CURRICULUM VITAE

Dong-Qing Wei

Updated 22nd September 2022

State Key Laboratory of Microbial Metabolism and College of Life Sciences and Biotechnology Shanghai Jiao Tong University 800 Dongchuan Road, Minhang District Shanghai, 200240, China Tel., 86-13918500529

E-mail:dqwei@sjtu.edu.cn

Website: https://scholar.google.com/citations?user=1rLPRzIAAAAJ&hl=zh-CN https://www.researchgate.net/profile/Dong-Qing_Wei

Education and Degrees

- 1987 1990 Post-doctoral Fellow, Department of Chemistry, University of British Columbia, Vancouver, B.C., Canada, with Gren Patey.
- 1985 1987 Ph.D., Chemical Physics Program at the Department of Physics, University of Puerto Rico, Rio Piedras, Puerto Rico, USA, thesis supervisor: Lesser Blum.
- 1982 1985 M.S., Department of Chemistry, Normal University of Henan, Xinxiang, Henan, People's Republic of China.
- 1978 1982 B.S., Department of Chemistry, Normal University of Henan, Xinxiang, Henan, People's Republic of China.

Skills & Activities

Computational Biology/Chemistry/Physics: Density Functional Theory, Bioinformatics Software/Tools/Techniques: Crystal, Ab Initio, Molecular Dynamics Simulation, Ferroelectrics, Lipase, Drug Design, Drug Discovery, Computational Chemistry, QM/MM, Hydrogen Bonding, Autodock, Molecular Mechanics, Molecular Dynamics, Intermolecular Interactions, Molecular Docking, Fluid, Electrolytes, Membrane Proteins, CPMD, Theoretical Chemistry, Cheminformatics and Computational Chemistry, Thermodynamics, Computer Simulations, Apoptosis Assays, Ferroelectric Materials, Free Energy, Phase Transitions, Computational Physics, Biostatistics, Statistical Mechanics, Virtual Screening, Conductivity, Pharmacophore, CSCW, Neuron, Electrostatics, Mutagenesis, Molecular Models, Algebra, Molecular Descriptors, Condensed Matter, Computational Systems Biology, Molecular Modeling, and Correlation.

Languages: English, French, Spanish (Castilian), Chinese

Academic and Research Positions

2006 – Present, Tenured Professor (长聘教授) at the State Key Laboratory of Microbial of Metabolism & Department of Bioinformatics and Biostatistics, College of Life Science and Biotechnology, Shanghai Jiao Tong University, Shanghai, China; an external member at Centre for Research in Molecular Modeling (CERMM), Concordia University in Montreal, Canada; and Adjunct Professor at Pengcheng National Lab.

2003-2006, The Haihe Distinguished Professor, appointed by the City of Tianjin at Tianjin Normal University, Directory General, Tianjin Institute of Bioinformatics and Drug Discoveries, Principal Scientist, Gordon Life Science Institute, San Diego, USA, and an external member at Centre for Research in Molecular Modeling (CERMM), Concordia University in Montreal, Canada.

1993 - 2003, Research Scientist, Centre de Recherche en Calcul Applique (CERCA, Research Center on Computation and its Application), Montreal, Quebec.

1990 - 2000, Professor of Chemistry, College of Chemistry and Molecular Engineering, Peeking University, Beijing, P.R. China.

1990 – 1992, Research Associate, Department of Chemistry, University of British Columbia, Vancouver, BC., Canada, with Gren Patey.

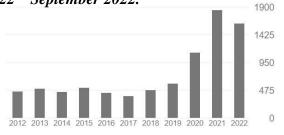
1985 – 1985, Assistant Professor, Department of Chemistry, Normal University of Henan, Xinxiang, Henan, People's Republic of China.

Scientific Publications

Publication Performance Metrics 22nd September 2022 by Google Scholar

Citation indices	All	Since 2017
Citations	12428	6067
h-index	61	36
i10-index	252	182

List of Publications (total SCI citations: 12428 times, original papers: 11,018, review and others: 1410) as of 22nd September 2022.



The 15 most significant contributions (cited 2050 times)

- 1. (Cited 445 times) Chou, K.-C., Wei D.Q., and W.-Z. Zhong, "Binding mechanism of coronavirus main proteinase with ligands and its implication to drug design against SARS". Biochemical and biophysical research communications, 2003. 308(1): p. 148-151.
- 2. (Cited 347 times) Wei, D. and G. Patey, "Orientational order in simple dipolar liquids: computer simulation of a ferroelectric nematic phase". Physical review letters, 1992. 68(13): p. 2043-2045.
- 3. (Cited 198 times) Wei, D. and G. Patey, "Ferroelectric liquid-crystal and solid phases formed by strongly interacting dipolar soft spheres". Physical Review A, 1992. 46(12): p. 7783-7792.
- 4. (Cited 182 times) Sirois, S., Wei D.Q., Q. Du, and K.-C. Chou, "Virtual screening for SARS-CoV protease based on KZ7088 pharmacophore points". Journal of chemical information and computer sciences, 2004. 44(3): p. 1111-1122.

- 5. (Cited 179 times) Wei, D. and D. Salahub, "Hydrated proton clusters and solvent effects on the proton transfer barrier: A density functional study". The Journal of chemical physics, 1994. 101(9): p. 7633-7642.
- 6. (Cited 141 times) Wang, J.-F., Wei D.Q., L. Li, S.-Y. Zheng, Y.-X. Li, and K.-C. Chou, "3D structure modeling of cytochrome P450 2C19 and its implication for personalized drug design". Biochemical and biophysical research communications, 2007. 355(2): p. 513-519.
- 7. (Cited 139 times) Wei, D. and D. Salahub, "Hydrated proton clusters: Ab initio molecular dynamics simulation and simulated annealing". The Journal of chemical physics, 1997. 106(14): p. 6086-6094.
- 8. (Cited 89 times) Chu, Y., A.C. Kaushik, X. Wang, W. Wang, Y. Zhang, X. Shan, D.R. Salahub, Y. Xiong, and Wei D.Q., "DTI-CDF: a cascade deep forest model towards the prediction of drug-target interactions based on hybrid features". Briefings in Bioinformatics, 2019, DOI: 10.1093/bib/bbz152.
- 9. (Cited 85 times) Chang, J., P. Lian, Wei D.Q., X.-R. Chen, Q.-M. Zhang, and Z.-Z. Gong, "Thermal decomposition of the solid phase of nitromethane: ab initio molecular dynamics simulations". Physical review letters, 2010. 105(18): p. 188302-188306.
- 10. (Cited 85 times) Wang, C., Wang, S., Li, D., Wei, D.Q.*, Zhao, J*. and Wang, J.*, "Human intestinal defensin 5 inhibits SARS-CoV-2 invasion by cloaking ACE2". Gastroenterology, 2020, DOI: 10.1053/j.gastro.2020.05.015.
- 11. (Cited 51 times) Tang, M., Z. Wang, Y. Zhou, W. Xu, S. Li, L. Wang, Wei D.Q., and Z. Qiao, "A novel drug candidate for Alzheimer's disease treatment: gx-50 derived from Zanthoxylum bungeanum". Journal of Alzheimer's Disease, 2013. 34(1): p. 203-213.
- 12. (Cited 40 times) Gu, R.-X., L.A. Liu, Wei D.Q., J.-G. Du, L. Liu, and H. Liu, "Free energy calculations on the two drug binding sites in the M2 proton channel". Journal of the American Chemical Society, 2011. 133(28): p. 10817-10825.
- 13. (Cited 34 times) Chu, Y., Shan, X., Chen, T., Jiang, M., Wang, Y., Wang, Q., Salahub, D.R., Xiong, Y. and Wei, D.Q., "DTI-MLCD: predicting drug-target interactions using multi-label learning with community detection method". Briefings in Bioinformatics, 2020, DOI: 10.1093/bib/bbaa205.
- 14. (Cited 30 times) Wang, Y., D. Hu, and Wei D.Q., "Transmembrane permeation mechanism of charged methyl guanidine". Journal of chemical theory and computation, 2014. 10(4): p. 1717-1726.
- 15. (Cited 8 times)Chu, Y., Zhang, Y., Wang, Q.K., Zhang, L.F., Wang, X.H., Wang, Y.J., , Salahub, D.R., Xu, Q., Wang, J.M., Jiang, X., Xiong, Y*, and Wei, D.Q.*, "A transformer-based model to predict peptide–HLA class I binding and optimize mutated peptides for vaccine design", Nature Machine Intelligence, 2022, 4(3):300-311, DOI: 10.1038/s42256-022-00459-7 (2021).

Other papers listed according to the number of citations (cited 8968 times)

16. (Cited 230 times) Khan, Abbas, et al. "Higher infectivity of the SARS-CoV-2 new variants is associated with K417N/T, E484K, and N501Y mutants: An insight from structural data." *Journal of cellular physiology* (2021), 236(10):7045-7057. doi: 10.1002/jcp.30367.

- 17. (Cited 141 times) Wei, D.-Q., J.-F. Wang, C. Chen, Y. Li, and K.-C. Chou, "Molecular modeling of two CYP2C19 SNPs and its implications for personalized drug design". Protein and peptide letters, 2008. 15(1): p. 27-32.
- 18. (Cited 135 times) Blum, L. and Wei D.Q., "Analytical solution of the mean spherical approximation for an arbitrary mixture of ions in a dipolar solvent". The Journal of chemical physics, 1987. 87(1): p. 555-565.
- 19. (Cited 130 times) Wei, D. and D. Salahub, "A combined density functional and molecular dynamics simulation of a quantum water molecule in aqueous solution". Chemical physics letters, 1994. 224(3-4): p. 291-296.
- 20. (Cited 126 times) Du, Q., S. Wang, Wei D.Q., S. Sirois, and K.-C. Chou, "Molecular modeling and chemical modification for finding peptide inhibitor against severe acute respiratory syndrome coronavirus main proteinase". Analytical Biochemistry, 2005. 337(2): p. 262-270.
- 21. (Cited 117 times) Dunbar, R.C., T.B. McMahon, D. Thoelmann, D.S. Tonner, D.R. Salahub, and Wei D.Q., "Zero-pressure thermal-radiation-induced dissociation of gas-phase cluster ions: comparison of theory and experiment for (H2O) 2Cl-and (H2O) 3Cl". Journal of the American Chemical Society, 1995. 117(51): p. 12819-12825.
- 22. (Cited 111 times) Wang, J.-F., K. Gong, Wei D.Q., Y.-X. Li, and K.-C. Chou, "Molecular dynamics studies on the interactions of PTP1B with inhibitors: from the first phosphate-binding site to the second one". Protein Engineering, Design & Selection, 2009. 22(6): p. 349-355.
- 23. (Cited 111 times) Khan, F.I., Wei D.Q., K.-R. Gu, M.I. Hassan, and S. Tabrez, "Current updates on computer aided protein modeling and designing". International journal of biological macromolecules, 2016. 85: p. 48-62.
- 24. (Cited 105 times) Iftimie, R., D. Salahub, Wei D.Q., and J. Schofield, "Using a classical potential as an efficient importance function for sampling from an ab initio potential". The Journal of Chemical Physics, 2000. 113(12): p. 4852-4862.
- 25. (Cited 102 times) Wang, S.-Q., Q.-S. Du, K. Zhao, A.-X. Li, Wei D.Q., and K.-C. Chou, "Virtual screening for finding natural inhibitor against cathepsin-L for SARS therapy". Amino Acids, 2007. 33(1): p. 129-135.
- 26. (Cited 101 times) Khan, M.T., Ali, A., Wang, Q., Irfan, M., Khan, A., Zeb, M.T., Zhang, Y.J., Chinnasamy, S. and Wei, D.Q., "Marine natural compounds as potents inhibitors against the main protease of SARS-CoV-2. A molecular dynamic study". Journal of Biomolecular Structure and Dynamics, 2020. pp.1-14.
- 27. (Cited 100 times) Du, Q.-S., S.-Q. Wang, Y. Zhu, Wei D.Q., H. Guo, S. Sirois, and K.-C. Chou, "Polyprotein cleavage mechanism of SARS CoV Mpro and chemical modification of the octapeptide". Peptides, 2004. 25(11): p. 1857-1864.
- 28. (Cited 94 times) Wei, D. and G. Patey, "Dynamics of molecular liquids: A comparison of different theories with application to wave vector dependent dielectric relaxation and ion solvation". The Journal of Chemical Physics, 93(2): p. 1399-1411.
- 29. (Cited 93 times) Liu, H., J. Zhao, Wei D.Q., and Z. Gong, "Structural and vibrational properties of solid nitromethane under high pressure by density functional theory". The Journal of chemical physics, 2006. 124(12): p. 124501-124510.
- 30. (Cited 92 times) Xiong, Y., Q. Wang, J. Yang, X. Zhu, and Wei D.Q., "PredT4SE-stack: prediction of bacterial type IV secreted effectors from protein sequences using a stacked ensemble method". Frontiers in Microbiology, 2018. 9: p. 2571-2580.

- 31. (Cited 89 times) Wang, Y., T. Zhao, Wei D.Q., E. Strandberg, A.S. Ulrich, and J.P. Ulmschneider, "How reliable are molecular dynamics simulations of membrane active antimicrobial peptides?". Biochimica et Biophysica Acta (BBA)-Biomembranes, 2014. 1838(9): p. 2280-2288.
- 32. (Cited 88 times) Wang, J.-F., Wei D.Q., Y. Lin, Y.-H. Wang, H.-L. Du, Y.-X. Li, and K.-C. Chou, "Insights from modeling the 3D structure of NAD (P) H-dependent D-xylose reductase of Pichia stipitis and its binding interactions with NAD and NADP". Biochemical and biophysical research communications, 2007. 359(2): p. 323-329.
- 33. (Cited 88 times) Wei, D., G. Patey, and A. Perera, "Orientational order in simple dipolar fluids: Density-functional theory and absolute-stability conditions". Physical Review E, 1993. 47(1): p. 506-512.
- 34. (Cited 87 times) Zhang, R., Wei D.Q., Q.-S. Du, and K.-C. Chou, "Molecular modeling studies of peptide drug candidates against SARS". Medicinal Chemistry, 2006. 2(3): p. 309-314.
- 35. (Cited 85 times) Sirois, S., G. Hatzakis, Wei D.Q., Q. Du, and K.-C. Chou, "Assessment of chemical libraries for their druggability". Computational Biology and Chemistry, 2005. 29(1): p. 55-67.
- 36. (Cited 80 times) Khan, M., S. Khan, A. Ali, H. Akbar, A.M. Sayaf, A. Khan, and Wei D.Q., "Immunoinformatics approaches to explore Helicobacter Pylori proteome (Virulence Factors) to design B and T cell multi-epitope subunit vaccine". Scientific reports, 2019. 9(1): p. 1-13.
- 37. (Cited 79 times) Xu, Q., Y. Xiong, H. Dai, K.M. Kumari, Q. Xu, H.-Y. Ou, and Wei D.Q., "PDC-SGB: Prediction of effective drug combinations using a stochastic gradient boosting algorithm". Journal of theoretical biology, 2017. 417: p. 1-7.
- 38. (Cited 78 times) Gao, W.-N., Wei D.Q., Y. Li, H. Gao, W.-R. Xu, A.-X. Li, and K.-C. Chou, "Agaritine and its derivatives are potential inhibitors against HIV proteases". Medicinal Chemistry, 2007. 3(3): p. 221-226.
- 39. (Cited 78 times) Ge, N.-N., Y.-K. Wei, G.-F. Ji, X.-R. Chen, F. Zhao, and Wei D.Q., "Initial decomposition of the condensed-phase β-HMX under shock waves: molecular dynamics simulations". The Journal of Physical Chemistry B, 2012. 116(46): p. 13696-13704.
- 40. (Cited 77 times) Wei, D.-Q., S. Sirois, Q.-S. Du, H.R. Arias, and K.-C. Chou, "Theoretical studies of Alzheimer's disease drug candidate 3-[(2, 4-dimethoxy) benzylidene]-anabaseine (GTS-21) and its derivatives". Biochemical and biophysical research communications, 2005. 338(2): p. 1059-1064.
- 41. (Cited 77 times) Khan, A., Khan, M., Saleem, S., Babar, Z., Ali, A., Khan, A.A., Sardar, Z., Hamayun, F., Ali, S.S. and Wei, D.Q., "Phylogenetic analysis and structural perspectives of RNA-dependent RNA-polymerase inhibition from SARs-CoV-2 with natural products". Interdisciplinary Sciences: Computational Life Sciences, 2020. 12(3), pp.335-348.
- 42. (Cited 76 times) Quimque, M.T.J., Notarte, K.I.R., Fernandez, R.A.T., Mendoza, M.A.O., Liman, R.A.D., Lim, J.A.K., Pilapil, L.A.E., Ong, J.K.H., Pastrana, A.M., Khan, A. and Wei, D.Q., "Virtual Screening-Driven Drug Discovery of SARS-CoV2 Enzyme Inhibitors Targeting Viral Attachment, Replication, Post-Translational Modification and Host Immunity Evasion Infection Mechanisms". Journal of Biomolecular Structure and Dynamics, 2020. pp.1-23.
- 43. (Cited 75 times) Khan, A., M. Junaid, A.C. Kaushik, A. Ali, S.S. Ali, A. Mehmood, and Wei D.Q., "Computational identification, characterization and validation of potential antigenic

- peptide vaccines from hrHPVs E6 proteins using immunoinformatics and computational systems biology approaches". PloS one, 2018. 13(5): p. e0196484.
- 44. (Cited 74 times) Wei, D. and L. Blum, "The mean spherical approximation for an arbitrary mixture of ions in a dipolar solvent: Approximate solution, pair correlation functions, and thermodynamics". The Journal of chemical physics, 1987. 87(5): p. 2999-3007.
- 45. (Cited 74 times) Chandra, A., Wei D.Q., and G. Patey, "The frequency dependent conductivity of electrolyte solutions". The Journal of chemical physics, 99(3): p. 2083-2094.
- 46. (Cited 73 times) Wang, J.-F., Wei D.Q., and K.-C. Chou, "Insights from investigating the interactions of adamantane-based drugs with the M2 proton channel from the H1N1 swine virus". Biochemical and biophysical research communications, 2009. 388(2): p. 413-417.
- 47. (Cited 72 times) Xiong, Y., J. Liu, and D.Q. Wei, "An accurate feature-based method for identifying DNA-binding residues on protein surfaces". Proteins: Structure, Function, and Bioinformatics, 2011. 79(2): p. 509-517.
- 48. (Cited 72 times) Attard, P., Wei D.Q., and G. Patey, "Critical comments on the nonlocal dielectric function employed in recent theories of the hydration force". Chemical Physics Letters, 1990. 172(1): p. 69-72.
- 49. (Cited 72 times) Khan, A., Ali, S.S., Khan, M.T., Saleem, S., Ali, A., Suleman, M., Babar, Z., Shafiq, A., Khan, M. and Wei, D.Q., "Combined drug repurposing and virtual screening strategies with molecular dynamics simulation identified potent inhibitors for SARS-CoV-2 main protease (3CLpro)". Journal of Biomolecular Structure and Dynamics, 2020. p.1-12.
- 50. (Cited 71 times) Gan, Y.-R., H. Huang, Y.-D. Huang, C.-M. Rao, Y. Zhao, J.-S. Liu, L. Wu, and Wei D.Q., "Synthesis and activity of an octapeptide inhibitor designed for SARS coronavirus main proteinase". Peptides, 2006. 27(4): p. 622-625.
- 51. (Cited 70 times) Li, Y., Wei D.Q., W.-N. Gao, H. Gao, B.-N. Liu, C.-J. Huang, W.-R. Xu, D.-K. Liu, H.-F. Chen, and K.-C. Chou, "Computational approach to drug design for oxazolidinones as antibacterial agents". Medicinal Chemistry, 2007. 3(6): p. 576-582.
- 52. (Cited 68 times) Wei, D.-Q., R. Zhang, Q.-S. Du, W.-N. Gao, Y. Li, H. Gao, S.-Q. Wang, X. Zhang, A.-X. Li, and S. Sirois, "Anti-SARS drug screening by molecular docking". Amino Acids, 2006. 31(1): p. 73-80.
- 53. (Cited 66 times) Wei, D. and G. Patey, "Rotational motion in molecular liquids". The Journal of chemical physics, 1989. 91(11): p. 7113-7129.
- 54. (Cited 66 times) Khan, A., Khan, M.T., Saleem, S., Junaid, M., Ali, A., Ali, S.S., Khan, M. and Wei, D.Q, "Structural Insights into the mechanism of RNA recognition by the N-terminal RNA-binding domain of the SARS-CoV-2 nucleocapsid phosphoprotein". Computational and Structural Biotechnology Journal, 2020.
- 55. (Cited 65 times) Chandra, A., Wei D.Q., and G. Patey, "Dielectric relaxation of electrolyte solutions: Is there really a kinetic dielectric decrement?". The Journal of chemical physics, 1993. 98(6): p. 4959-4966.
- 56. (Cited 64 times) Li, L., Wei D.Q., J.-F. Wang, and K.-C. Chou, "Computational studies of the binding mechanism of calmodulin with chrysin". Biochemical and biophysical research communications, 2007. 358(4): p. 1102-1107.
- 57. (Cited 63 times) Wei, D., J.-F. Truchon, S. Sirois, and D. Salahub, "Solvation of formic acid and proton transfer in hydrated clusters". The Journal of chemical physics, 2002. 116(14): p. 6028-6038.

- 58. (Cited 62 times) Lian, P., Wei D.Q., J.-F. Wang, and K.-C. Chou, "An allosteric mechanism inferred from molecular dynamics simulations on phospholamban pentamer in lipid membranes". PLoS One, 2011. 6(4): p. 18587-18593.
- 59. (Cited 60 times) Gu, R.-X., H. Gu, Z.-Y. Xie, J.-F. Wang, H.R. Arias, Wei D.Q., and K.-C. Chou, "Possible drug candidates for Alzheimer's disease deduced from studying their binding interactions with α7 nicotinic acetylcholine receptor". Medicinal Chemistry, 2009. 5(3): p. 250-262.
- 60. (Cited 60 times) Khan, S., A. Khan, A.U. Rehman, I. Ahmad, S. Ullah, A.A. Khan, S.S. Ali, S. Gul, and Wei D.Q., "Immunoinformatics and structural vaccinology driven prediction of multi-epitope vaccine against Mayaro virus and validation through in-silico expression". Infection, Genetics and Evolution, 2019.p.390-400.
- 61. (Cited 59 times) Zheng, H., Wei D.Q., R. Zhang, C. Wang, H. Wei, and K.-C. Chou, "Screening for new agonists against Alzheimer's disease". Medicinal Chemistry, 2007. 3(5): p. 488-493.
- 62. (Cited 59 times) Wang, Y., Wei D.Q., and J.-F. Wang, "Molecular dynamics studies on T1 lipase: insight into a double-flap mechanism". Journal of chemical information and modeling, 2010. 50(5): p. 875-878.
- 63. (Cited 59 times) Arias, H.R., R.-X. Gu, D. Feuerbach, B.-B. Guo, Y. Ye, and Wei D.Q., "Novel positive allosteric modulators of the human α7 nicotinic acetylcholine receptor". Biochemistry, 2011. 50(23): p. 5263-5278.
- 64. (Cited 59 times) Fan, H.-M., R.-X. Gu, Y.-J. Wang, Y.-L. Pi, Y.-H. Zhang, Q. Xu, and Wei D.Q., "Destabilization of Alzheimer's Aβ42 protofibrils with a novel drug candidate wgx-50 by molecular dynamics simulations". The Journal of Physical Chemistry B, 2015. 119(34): p. 11196-11202.
- 65. (Cited 58 times) Ali, A., A. Khan, A.C. Kaushik, Y. Wang, S.S. Ali, M. Junaid, S. Saleem, W.C. Cho, X. Mao, and Wei D.Q., "Immunoinformatic and systems biology approaches to predict and validate peptide vaccines against Epstein–Barr virus (EBV)". Scientific reports, 2019. 9(1): p. 720-732.
- 66. (Cited 57 times) Du, Q., S. Wang, Z. Jiang, W. Gao, Y. Li, Wei D.Q., and K.-C. Chou, "Application of bioinformatics in search for cleavable peptides of SARSCoV Mpro and chemical modification of octapeptides". Medicinal Chemistry, 2005. 1(3): p. 209-213.
- 67. (Cited 56 times) Wei, D., E. Proynov, A. Milet, and D. Salahub, "Solvation of the hydroxide anion: A combined DFT and molecular dynamics study". The Journal of Physical Chemistry A, 2000. 104(11): p. 2384-2395.
- 68. (Cited 56 times) Chandra, A., Wei D.Q., and G. Patey, "Microscopic theory of solvation dynamics in dipolar liquids". The Journal of chemical physics, 1993. 99(7): p. 4926-4931.
- 69. (Cited 56 times) Kaushik, A.C., S. Bharadwaj, S. Kumar, and Wei D.Q., "Nano-particle mediated inhibition of Parkinson's disease using computational biology approach". Scientific reports, 2018. 8(1): p. 9169-9177.
- 70. (Cited 56 times) Cui, H.-L., G.-F. Ji, X.-R. Chen, W.-H. Zhu, F. Zhao, Y. Wen, and Wei D.Q., "First-principles study of high-pressure behavior of solid β-HMX". The Journal of Physical Chemistry A, 2009. 114(2): p. 1082-1092.
- 71. (Cited 55 times) Gong, K., L. Li, J.-F. Wang, F. Cheng, Wei D.Q., and K.-C. Chou, "Binding mechanism of H5N1 influenza virus neuraminidase with ligands and its implication for drug design". Medicinal Chemistry, 2009. 5(3): p. 242-249.

- 72. (Cited 53 times) Liu, H., J. Zhao, G. Ji, Wei D.Q., and Z. Gong, "Vibrational properties of molecule and crystal of TATB: A comparative density functional study". Physics Letters A, 2006. 358(1): p. 63-69.
- 73. (Cited 52 times) Khan, F.I., M. Aamir, Wei D.Q., F. Ahmad, and M.I. Hassan, "Molecular mechanism of Ras-related protein Rab-5A and effect of mutations in the catalytically active phosphate-binding loop". Journal of Biomolecular Structure and Dynamics, 2017. 35(1): p. 105-118.
- 74. (Cited 49 times) Wei, H., R. Zhang, C. Wang, H. Zheng, A. Li, K.-C. Chou, and Wei D.Q., "Molecular insights of SAH enzyme catalysis and implication for inhibitor design". Journal of theoretical biology, 2007. 244(4): p. 692-702.
- 75. (Cited 49 times) Guo, X.-L., L. Li, Wei D.Q., Y.-S. Zhu, and K.-C. Chou, "Cleavage mechanism of the H5N1 hemagglutinin by trypsin and furin". Amino acids, 2008. 35(2): p. 375-382.
- 76. (Cited 49 times) Lian, P., H.-B. Guo, D. Riccardi, A. Dong, J.M. Parks, Q. Xu, E.F. Pai, S.M. Miller, Wei D.Q., and J.C. Smith, "X-ray structure of a Hg2+ complex of mercuric reductase (MerA) and quantum mechanical/molecular mechanical study of Hg2+ transfer between the C-terminal and buried catalytic site cysteine pairs". Biochemistry, 2014. 53(46): p. 7211-7222.
- 77. (Cited 49 times) Kaushik, A.C., X. Mao, C.-D. Li, Y. Li, Wei D.Q., and S. Sahi, "G-protein-coupled receptors function as logic gates for nanoparticle binding using systems and synthetic biology approach". Journal of Materials Research, 2019. 34(11): p. 1854-1867.
- 78. (Cited 49 times) Khan, Abbas, et al. "In silico and in vitro evaluation of kaempferol as a potential inhibitor of the SARS-CoV-2 main protease (3CLpro)." *Phytotherapy Research* (2021).
- 79. (Cited 48 times) Zeng, Q.-K., H.-L. Du, J.-F. Wang, Wei D.Q., X.-N. Wang, Y.-X. Li, and Y. Lin, "Reversal of coenzyme specificity and improvement of catalytic efficiency of Pichiastipitis xylose reductase by rational site-directed mutagenesis". Biotechnology letters, 2009. 31(7): p. 1025-1029.
- 80. (Cited 47 times) Liu, H., J. Zhao, J. Du, Z. Gong, G. Ji, and Wei D.Q., "High-pressure behavior of TATB crystal by density functional theory". Physics Letters A, 2007. 367(4-5): p. 383-388.
- 81. (Cited 47 times) Selvaraj, G., S. Kaliamurthi, A.C. Kaushik, A. Khan, Y.-K. Wei, W.C. Cho, K. Gu, and Wei D.Q., "Identification of target gene and prognostic evaluation for lung adenocarcinoma using gene expression meta-analysis, network analysis and neural network algorithms". Journal of biomedical informatics, 2018. 86: p. 120-134.
- 82. (Cited 45 times) Khan, M.T., A.C. Kaushik, S.I. Malik, S. Ali, and Wei D.Q., "Artificial neural networks for prediction of tuberculosis disease". Frontiers in microbiology, 2019. 10: p. 395-404.
- 83. (Cited 43 times) Wei, D., H. Guo, and D. Salahub, "Conformational dynamics of an alanine dipeptide analog: An ab initio molecular dynamics study". Physical Review E, 2001. 64(1): p. 011907-011911.
- 84. (Cited 42 times) Zhang, T., L.A. Liu, D.F. Lewis, and Wei D.Q., "Long-range effects of a peripheral mutation on the enzymatic activity of cytochrome P450 1A2". Journal of chemical information and modeling, 2011. 51(6): p. 1336-1346.

- 85. (Cited 42 times) Shan, X., X. Wang, C.-D. Li, Y. Chu, Y. Zhang, Y. Xiong, and Wei D.Q., "Prediction of CYP450 Enzyme-Substrate Selectivity Based on the Network-based Label Space Division Method". Journal of chemical information and modeling, 2019.
- 86. (Cited 41 times) Wei, D. and G. Patey, "Dielectric relaxation of electrolyte solutions". The Journal of chemical physics, 1991. 94(10): p. 6795-6806.
- 87. (Cited 41 times) Ge, N.-N., Y.-K. Wei, Z.-F. Song, X.-R. Chen, G.-F. Ji, F. Zhao, and Wei D.Q., "Anisotropic responses and initial decomposition of condensed-phase β-HMX under shock loadings via molecular dynamics simulations in conjunction with multiscale shock technique". The Journal of Physical Chemistry B, 2014. 118(29): p. 8691-8699.
- 88. (Cited 40 times) Khan, F.I., B. Nizami, R. Anwer, K.-R. Gu, K. Bisetty, M.I. Hassan, and Wei D.Q., "Structure prediction and functional analyses of a thermostable lipase obtained from Shewanella putrefaciens". Journal of Biomolecular Structure and Dynamics, 2017. 35(10): p. 2123-2135.
- 89. (Cited 40 times) Khan, M.T., A. Khan, A.U. Rehman, Y. Wang, K. Akhtar, S.I. Malik, and Wei D.Q., "Structural and free energy landscape of novel mutations in ribosomal protein S1 (rpsA) associated with pyrazinamide resistance". Scientific reports, 2019. 9(1): p. 7482-7494.
- 90. (Cited 39 times) Zi, J., D. Liu, P. Ma, H. Huang, J. Zhu, Wei D.Q., J. Yang, and C. Chen, "Effects of CYP2C9* 3 and CYP2C9* 13 on diclofenac metabolism and inhibition-based drugdrug interactions". Drug metabolism and pharmacokinetics, 2010. 25(4): p. 343-350.
- 91. (Cited 39 times) Zhang, Y.-F., X. Wang, A.C. Kaushik, Y. Chu, X. Shan, M.-Z. Zhao, Q. Xu, and Wei D.Q., "SPVec: A Word2vec-Inspired Feature Representation Method for Drug-Target Interaction Prediction". Frontiers in Chemistry, 2020. 7: p. 895.
- 92. (Cited 37 times) Sun, Y., Y. Xiong, Q. Xu, and Wei D.Q., "A hadoop-based method to predict potential effective drug combination". BioMed research international, 2014. 2014.
- 93. (Cited 37 times) Khan, Abbas, et al. "Immunogenomics guided design of immunomodulatory multi-epitope subunit vaccine against the SARS-CoV-2 new variants, and its validation through in silico cloning and immune simulation." *Computers in biology and medicine* 133 (2021): 104420.
- 94. (Cited 36 times) Li, J., Wei D.Q., J.-F. Wang, and Y.-X. Li, "A negative cooperativity mechanism of human CYP2E1 inferred from molecular dynamics simulations and free energy calculations". Journal of chemical information and modeling, 2011. 51(12): p. 3217-3225.
- 95. (Cited 35 times) Yuan, X.-L., Wei D.Q., X.-R. Chen, Q.-M. Zhang, and Z.-Z. Gong, "The first-principles calculations for the elastic properties of Zr2Al under compression". Journal of Alloys and Compounds, 2011. 509(3): p. 769-774.
- 96. (Cited 35 times) Lu, L.-Y., Wei D.Q., X.-R. Chen, D. Lian, G.-F. Ji, Q.-M. Zhang, and Z.-Z. Gong, "The first principle studies of the structural and vibrational properties of solid β-HMX under compression". Molecular Physics, 2008. 106(21-23): p. 2569-2580.
- 97. (Cited 35 times) Nangraj, A.S., Selvaraj, G., Kaliamurthi, S., Kaushik, A.C., Cho, W.C. and Wei, D.Q., "Integrated PPI-and WGCNA-Retrieval of Hub Gene Signatures Shared Between Barrett's Esophagus and Esophageal Adenocarcinoma". Frontiers in Pharmacology, 2020. 11.
- 98. (Cited 35 times)Khan, Abbas, et al. "SARS-CoV-2 new variants: Characteristic features and impact on the efficacy of different vaccines." *Biomedicine & Pharmacotherapy* (2021): 112176.
- 99. (Cited 35 times) Wang, Cheng, et al. "Human Cathelicidin Inhibits SARS-CoV-2 Infection: Killing Two Birds with One Stone." *ACS infectious diseases* (2021).

- 100. (Cited 34 times) Li, K., Y. Du, L. Li, and Wei D.Q., "Bioinformatics Approaches for Anticancer Drug Discovery". Current drug targets, 2020. 21(1): p. 3-17.
- 101. (Cited 34 times) Ayton, G., Wei D.Q., and G. Patey, "Liquid crystal phases of dipolar discotic particles". Physical Review E, 1997. 55(1): p. 447-454.
- 102. (Cited 33 times) Gu, R.-X., L.A. Liu, Y.-H. Wang, Q. Xu, and Wei D.Q., "Structural comparison of the wild-type and drug-resistant mutants of the influenza A M2 proton channel by molecular dynamics simulations". The Journal of Physical Chemistry B, 2013. 117(20): p. 6042-6051.
- 103. (Cited 33 times) Zhao, Y., J. Li, H. Gu, Wei D.Q., Y.-c. Xu, W. Fu, and Z. Yu, "Conformational preferences of π – π stacking between ligand and protein, analysis derived from crystal structure data geometric preference of π – π interaction". Interdisciplinary Sciences: Computational Life Sciences, 2015. 7(3): p. 211-220.
- 104. (Cited 33 times) Kaliamurthi, S., G. Selvaraj, A.C. Kaushik, K.-R. Gu, and Wei D.Q., "Designing of CD8+ and CD8+-overlapped CD4+ epitope vaccine by targeting late and early proteins of human papillomavirus". Biologics: targets & therapy, 2018. 12: p. 107-125.
- 105. (Cited 33 times) Cui, H.-L., G.-F. Ji, X.-R. Chen, Q.-M. Zhang, Wei D.Q., and F. Zhao, "Phase transitions and mechanical properties of octahydro-1, 3, 5, 7-tetranitro-1, 3, 5, 7-tetrazocine in different crystal phases by molecular dynamics simulation". Journal of Chemical & Engineering Data, 2010. 55(9): p. 3121-3129.
- 106. (Cited 33 times) Kotni, M.K., M. Zhao, and Wei D.Q., "Gene expression profiles and protein-protein interaction networks in amyotrophic lateral sclerosis patients with C9orf72 mutation". Orphanet journal of rare diseases, 2016. 11(1): p. 148-156.
- 107. (Cited 32 times) Chandra, A., Wei D.Q., and G. Patey, "Dielectric relaxation of dipolar liquids". The Journal of chemical physics, 1993. 99(3): p. 2068-2073.
- 108. (Cited 31 times) Li, J., Wei D.Q., J.-F. Wang, Z.-T. Yu, and K.-C. Chou, "Molecular dynamics simulations of CYP2E1". Medicinal Chemistry, 2012. 8(2): p. 208-221.
- 109. (Cited 31 times) Arias, H.R., R.-X. Gu, D. Feuerbach, and Wei D.Q., "Different interaction between the agonist JN403 and the competitive antagonist methyl lycaconitine with the human α7 nicotinic acetylcholine receptor". Biochemistry, 2010. 49(19): p. 4169-4180.
- 110. (Cited 31 times) Cui, S., Wei D.Q., H. Hu, W. Feng, and Z. Gong, "First-principles study of the structural and elastic properties of Cr2AlX (X= N, C) compounds". Journal of Solid State Chemistry, 2012. 191: p. 147-152.
- 111. (Cited 30 times) Lu, L.-Y., Wei D.Q., X.-R. Chen, G.-F. Ji, X.-J. Wang, J. Chang, Q.-M. Zhang, and Z.-Z. Gong, "The pressure-induced phase transition of the solid β–HMX". Molecular Physics, 2009. 107(22): p. 2373-2385.
- 112. (Cited 30 times) Kaliamurthi, S., G. Selvaraj, M. Junaid, A. Khan, K. Gu, and Wei D.Q., "Cancer Immunoinformatics: A Promising Era in the Development of Peptide Vaccines for Human Papillomavirus-induced Cervical Cancer". Current pharmaceutical design, 2018. 24(32): p. 3791-3817.
- 113. (Cited 30 times) Kaushik, A.C., S. Kumar, D.Q. Wei, and S. Sahi, "Structure based virtual screening studies to identify novel potential compounds for GPR142 and their relative dynamic analysis for study of type 2 diabetes". Frontiers in chemistry, 2018. 6: p. 23-37.

- 114. (Cited 30 times) Li, L., C.C. Koh, D. Reker, J. Brown, H. Wang, N.K. Lee, H.-h. Liow, H. Dai, H.-M. Fan, and L. Chen, "Predicting protein-ligand interactions based on bow-pharmacological space and Bayesian additive regression trees". Scientific reports, 2019. 9(1): p. 7703-7715.
- 115. (Cited 30 times) Khan, A., A.C. Kaushik, S.S. Ali, N. Ahmad, and Wei D.Q., "Deep-learning-based target screening and similarity search for the predicted inhibitors of the pathways in Parkinson's disease". RSC advances, 2019. 9(18): p. 10326-10339.
- 116. (Cited 30 times) Khan, A., J.M. Ashfaq-Ur-Rehman, C. Li, S. Saleem, F. Humayun, S. Shamas, S. Ali, Z. Babar, and Wei D.Q., "Dynamics insights into the gain of flexibility by Helix-12 in ESR1 as a mechanism of resistance to drugs in breast cancer cell lines". Front. Mol. Biosci. 6: 159. doi: 10.3389/fmolb, 2020.
- 117. (Cited 30 times) Attard, P., Wei D.Q., G. Patey, and G. Torrie, "The interaction between macroparticles in molecular fluids". The Journal of chemical physics, 1990. 93(10): p. 7360-7368.
- 118. (Cited 29 times) Wei, D., G. Torrie, and G. Patey, "Molecular solvent model for an electrical double layer: Effects of ionic polarizability". The Journal of chemical physics, 1993. 99(5): p. 3990-3997.
- 119. (Cited 29 times) Khan, Abbas, et al. "Preliminary Structural Data Revealed That the SARS-CoV-2 B. 1.617 Variant's RBD Binds to ACE2 Receptor Stronger Than the Wild Type to Enhance the Infectivity." *ChemBioChem* 22.16 (2021): 2641.
- 120. (Cited 28 times) Wei, D. and L. Blum, "Nonprimitive model of electrolytes: Analytical solution of the mean spherical approximation for an arbitrary mixture of sticky ions and dipoles". The Journal of chemical physics, 1988. 89(2): p. 1091-1100.
- 121. (Cited 28 times) Khan, Muhammad Tahir, et al. "SARS-CoV-2 nucleocapsid and Nsp3 binding: an in silico study." *Archives of Microbiology* 203.1 (2021): 59-66.
- 122. (Cited 28 times) Kaliamurthi, S., G. Selvaraj, S. Chinnasamy, Q. Wang, A.S. Nangraj, W. Cho, K. Gu, and Wei D.Q., "Exploring the Papillomaviral Proteome to Identify Potential Candidates for a Chimeric Vaccine against Cervix Papilloma Using Immunomics and Computational Structural Vaccinology". Viruses, 2019. 11(1): p. 63-88.
- 123. (Cited 28 times) Khan, Abbas, et al. "The SARS-CoV-2 B. 1.618 variant slightly alters the spike RBD–ACE2 binding affinity and is an antibody escaping variant: a computational structural perspective." *RSC Advances* 11.48 (2021): 30132-30147.
- 124. (Cited 28 times) Khan, M.T., Zeb, M.T., Ahsan, H., Ahmed, A., Ali, A., Akhtar, K., Malik, S.I., Cui, Z., Ali, S., Khan, A.S. and Ahmad, M, "SARS-CoV-2 nucleocapsid and Nsp3 binding: an in silico study". Archives of Microbiology, 2020. pp.1-8.
- 125. (Cited 27 times) Hussain, Irfan, et al. "Evolutionary and structural analysis of SARS-CoV-2 specific evasion of host immunity." *Genes & Immunity* 21.6 (2020): 409-419.
- 126. (Cited 27 times) Attard, P., Wei D.Q., and G. Patey, "On the existence of exact conditions in the theory of electrical double layers". The Journal of chemical physics, 1992. 96(5): p. 3767-3771.
- 127. (Cited 27 times) Yuan, X.-L., Wei D.Q., Y. Cheng, G.-F. Ji, Q.-M. Zhang, and Z.-Z. Gong, "Pressure effects on elastic and thermodynamic properties of Zr3Al intermetallic compound". Computational Materials Science, 2012. 58: p. 125-130.

- 128. (Cited 26 times) Dan, L., L. Lai-Yu, W. Dong-Qing, Z. Qing-Ming, G. Zi-Zheng, and G. Yong-Xin, "High-pressure behaviour of β-HMX crystal studied by DFT-LDA". Chinese Physics Letters, 2008. 25(3): p. 899-902.
- 129. (Cited 26 times) Wang, W., Q., Li, F., Xiong, Y, and Wei, D.Q., "MLCDForest: multi-label classification with deep forest in disease prediction for long non-coding RNAs". Briefings in Bioinformatics, 2020, DOI: 10.1093/bib/bbaa104.
- 130. (Cited 26 times) Liu, L., Y. Xiong, H. Gao, Wei D.Q., J.C. Mitchell, and X. Zhu, "dbAMEPNI: a database of alanine mutagenic effects for protein–nucleic acid interactions". Database, 2018. 2018. P. 1-7.
- 131. (Cited 26 times) Zhao, M. and Wei D.Q., "Rare diseases: Drug discovery and informatics resource". Interdisciplinary Sciences: Computational Life Sciences, 2018. 10(1): p. 195-204.
- 132. (Cited 25 times) Fazal, H., B.H. Abbasi, N. Ahmad, M. Ali, S. Shujait Ali, A. Khan, and Wei D.Q., "Sustainable production of biomass and industrially important secondary metabolites in cell cultures of selfheal (Prunella vulgaris L.) elicited by silver and gold nanoparticles". Artificial Cells, Nanomedicine, and Biotechnology, 2019. 47(1): p. 2553-2561.
- 133. (Cited 25 times) Chinnasamy, S., G. Selvaraj, C. Selvaraj, A.C. Kaushik, S. Kaliamurthi, A. Khan, S.K. Singh, and Wei D.Q., "Combining in silico and in vitro approaches to identification of potent inhibitor against phospholipase A2 (PLA2)". International Journal of Biological Macromolecules, 2020. 144: p. 53-66.
- 134. (Cited 24 times) Zhang, H.-Y., Q. Xu, Y.-K. Wang, T.-Z. Zhao, D. Hu, and Wei D.Q., "Passive transmembrane permeation mechanisms of monovalent ions explored by molecular dynamics simulations". Journal of chemical theory and computation, 2016. 12(10): p. 4959-4969.
- 135. (Cited 24 times) Xiong, Y., Y. Qiao, D. Kihara, H.-Y. Zhang, X. Zhu, and Wei D.Q., "Survey of machine learning techniques for prediction of the isoform specificity of cytochrome P450 substrates". Current drug metabolism, 2019. 20(3): p. 229-235.
- 136. (Cited 24 times) Hou, S., R.-X. Gu, and Wei D.Q., "Inhibition of β-amyloid channels with a drug candidate wgx-50 revealed by molecular dynamics simulations". Journal of chemical information and modeling, 2017. 57(11): p. 2811-2821.
- 137. (Cited 23 times) Wei, D. and L. Blum, "Internal energy in the mean spherical approximation as compared to Debye-Hueckel theory". Journal of Physical Chemistry, 1987. 91(16): p. 4342-4343.
- 138. (Cited 23 times) Wang, Y., A. Khan, A. Chandra Kaushik, M. Junaid, X. Zhang, and Wei D.Q., "The systematic modeling studies and free energy calculations of the phenazine compounds as anti-tuberculosis agents". Journal of Biomolecular Structure and Dynamics, 2019. 37(15): p. 4051-4069.
- 139. (Cited 23 times) Chang, J., X.-R. Chen, Wei D.Q., and X.-L. Yuan, "Elastic constants and anisotropy of β-BC2N under pressure". Physica B: Condensed Matter, 2010. 405(17): p. 3751-3755.
- 140. (Cited 23 times) Khan, M.T., A.U. Rehaman, M. Junaid, S.I. Malik, and Wei D.Q., "Insight into novel clinical mutants of RpsA-S324F, E325K, and G341R of Mycobacterium tuberculosis associated with pyrazinamide resistance". Computational and structural biotechnology journal, 2018. 16: p. 379-387.

- 141. (Cited 23 times) Bai, L.-Y., H. Dai, Q. Xu, M. Junaid, S.-L. Peng, X. Zhu, Y. Xiong, and Wei D.Q., "Prediction of effective drug combinations by an improved naive Bayesian algorithm". International journal of molecular sciences, 2018. 19(2): p. 467-479.
- 142. (Cited 23 times) Wang, X., Y. Wang, Z. Xu, Y. Xiong, and Wei D.Q., "ATC-NLSP: prediction of the classes of anatomical therapeutic chemicals using a network-based label space partition method". Frontiers in pharmacology, 2019. 10: p. 971-981.
- 143. (Cited 23 times)Khan, Abbas, et al. "An integrated systems biology and network-based approaches to identify novel biomarkers in breast cancer cell lines using gene expression data." *Interdisciplinary Sciences: Computational Life Sciences* (2020): 1-14.
- 144. (Cited 22 times) Zhang, X.-Q., J.-N. Yuan, G. Selvaraj, G.-F. Ji, X.-R. Chen, and Wei D.Q., "Towards the low-sensitive and high-energetic co-crystal explosive CL-20/TNT: from intermolecular interactions to structures and properties". Physical Chemistry Chemical Physics, 2018. 20(25): p. 17253-17261.
- 145. (Cited 22 times) Du, Q. and Wei D.Q., "Solvation and polarization of the N-methyl amine molecule in aqueous solution: A combined study of quantum mechanics and integral equation theory in three dimensions". The Journal of Physical Chemistry B, 2003. 107(48): p. 13463-13470.
- 146. (Cited 22 times) Wang, X.-D., J.-L. Huang, L. Yang, Wei D.Q., Y.-X. Qi, and Z.-L. Jiang, "Identification of human disease genes from interactome network using graphlet interaction". PloS one, 2014. 9(1): p. 86142-8612.
- 147. (Cited 21 times) Zhao, M., Q. Zhou, W. Ma, and Wei D.Q., "Exploring the ligand-protein networks in traditional Chinese medicine: current databases, methods, and applications". Evidence-Based Complementary and Alternative Medicine, 2013. 2013: p. 1-15.
- 148. (Cited 20 times) Sirois, S., C. Tsoukas, K.-C. Chou, Wei D.Q., C. Boucher, and G. Hatzakis, "Selection of molecular descriptors with artificial intelligence for the understanding of HIV-1 protease peptidomimetic inhibitors-activity". Medicinal Chemistry, 2005. 1(2): p. 173-184.
- 149. (Cited 20 times) Khan, A., S. Saleem, M. Idrees, S.S. Ali, M. Junaid, A.C. Kaushik, and Wei D.Q., "Allosteric ligands for the pharmacologically important Flavivirus target (NS5) from ZINC database based on pharmacophoric points, free energy calculations and dynamics correlation". Journal of Molecular Graphics and Modelling, 2018. 82: p. 37-47.
- 150. (Cited 20 times) Zhang, T., H. Dai, L.A. Liu, D.F. Lewis, and Wei D.Q., "Classification models for predicting cytochrome P450 enzyme-substrate selectivity". Molecular informatics, 2012. 31(1): p. 53-62.
- 151. (Cited 20 times) Junaid, M., M.T. Khan, S.I. Malik, and Wei D.Q., "Insights into the Mechanisms of the Pyrazinamide Resistance of Three Pyrazinamidase Mutants N11K, P69T, and D126N". Journal of chemical information and modeling, 2018. 59(1): p. 498-508.
- 152. (Cited 20 times) Chinnasamy, S., G. Selvaraj, A.C. Kaushik, S. Kaliamurthi, S. Chandrabose, S.K. Singh, R. Thirugnanasambandam, K. Gu, and Wei D.Q., "Molecular docking and molecular dynamics simulation studies to identify potent AURKA inhibitors: assessing the performance of density functional theory, MM-GBSA and mass action kinetics calculations". Journal of Biomolecular Structure and Dynamics, 2019: p. 1-11.
- 153. (Cited 19 times) Khan, Abbas, et al. "The Omicron (B. 1.1. 529) variant of SARS-CoV-2 binds to the hACE2 receptor more strongly and escapes the antibody response: Insights from structural and simulation data." *International Journal of Biological Macromolecules* (2022).

- 154. (Cited 19 time) Wang, Q., A. Mehmood, H. Wang, Q. Xu, Y. Xiong, and Wei D.Q., "Computational Screening and Analysis of Lung Cancer Related Non-Synonymous Single Nucleotide Polymorphisms on the Human Kirsten Rat Sarcoma Gene". Molecules, 2019. 24(10): p. 1951-1971.
- 155. (Cited 19 times) Mehmood, A., A.C. Kaushik, and D.Q. Wei, "Prediction and validation of potent peptides against herpes simplex virus type 1 via immunoinformatic and systems biology approach". Chemical biology & drug design, 2019.p. 1868-1883.
- 156. (Cited 19 times) Kaliamurthi, S., G. Selvaraj, L. Hou, Z. Li, Y. Wei, K. Gu, and Wei D.Q., "Synergism of essential oils with lipid based nanocarriers: Emerging trends in preservation of grains and related food products". Grain & Oil Science and Technology, 2019.p.21-26.
- 157. (Cited 19 times) Wei, D., G. Patey, and G. Torrie, "Double-layer structure at an ion-adsorbing surface". Journal of Physical Chemistry, 1990. 94(10): p. 4260-4268.
- 158. (Cited 19 times) Chu, Yanyi, et al. "MDA-GCNFTG: identifying miRNA-disease associations based on graph convolutional networks via graph sampling through the feature and topology graph." *Briefings in Bioinformatics* (2021).
- 159. (Cited 19 times) Wei, D. and G. Patey, "Dielectric relaxation of liquid mixtures". The Journal of chemical physics, 1991. 94(10): p. 6785-6794.
- 160. (Cited 18 times) Liu, H., J. Zhao, G. Ji, Z. Gong, and Wei D.Q., "Compressibility of liquid nitromethane in the high-pressure regime". Physica B: Condensed Matter, 2006. 382(1-2): p. 334-339.
- 161. (Cited 18 times) Khan, Muhammad Tahir, et al. "Structures of SARS-CoV-2 RNA-Binding Proteins and Therapeutic Targets." *Intervirology* 64.2 (2021): 55-68.
- 162. (Cited 18 times) Selvaraj, Gurudeeban, et al. "Are the Allergic Reactions of COVID-19 Vaccines Caused by mRNA Constructs or Nanocarriers? Immunological Insights." *Interdisciplinary Sciences: Computational Life Sciences* (2021): 1-4.
- 163. (Cited 18 times) Wang, Wei, et al. "LMI-DForest: A deep forest model towards the prediction of lncRNA-miRNA interactions." *Computational Biology and Chemistry* (2020): 107406.
- 164. (Cited 18 times) Wei, D. and G. Patey, "Dynamical properties of a ferroelectric nematic liquid crystal". Physical Review E, 1993. 47(4): p. 2954-2957.
- 165. (Cited 18 times) Khan, M.T., M. Junaid, X. Mao, Y. Wang, A. Hussain, S.I. Malik, and D.Q. Wei, "Pyrazinamide resistance and mutations L19R, R140H, and E144K in Pyrazinamidase of Mycobacterium tuberculosis". Journal of cellular biochemistry, 2019. 120(5): p. 7154-7166.
- 166. (Cited 18 times) Selvaraj, Gurudeeban, et al. "Are the Allergic Reactions of COVID-19 Vaccines Caused by mRNA Constructs or Nanocarriers? Immunological Insights." *Interdisciplinary Sciences: Computational Life Sciences* (2021): 1-4.
- 167. (Cited 18 times) Selvaraj, G., S. Kaliamurthi, S. Lin, K. Gu, and Wei D.Q., "Prognostic impact of tissue inhibitor of metalloproteinase-1 in non-small cell lung cancer: Systematic review and meta-analysis". Current medicinal chemistry, 2019.p.1-19.
- 168. (Cited 17 times) Wei, Y.-K., N.-N. Ge, G.-F. Ji, X.-R. Chen, L.-C. Cai, S.-Q. Zhou, and Wei D.Q., "Elastic, superconducting, and thermodynamic properties of the cubic metallic phase of AlH3 via first-principles calculations". Journal of Applied Physics, 2013. 114(11): p. 114905-114915.
- 169. (Cited 17 times) Wei, Y.-K., J.-N. Yuan, F.I. Khan, G.-F. Ji, Z.-W. Gu, and Wei D.Q., "Pressure induced superconductivity and electronic structure properties of scandium hydrides using first principles calculations". RSC Advances, 2016. 6(85): p. 81534-81541.

- 170. (Cited 17 times) Chen, Q., J.K. Buolamwini, J.C. Smith, A. Li, Q. Xu, X. Cheng, and Wei D.Q., "Impact of resistance mutations on inhibitor binding to HIV-1 integrase". Journal of chemical information and modeling, 2013. 53(12): p. 3297-3307.
- 171. (Cited 17 times) Kaushik, A.C., A. Kumar, Z. Peng, A. Khan, M. Junaid, A. Ali, S. Bharadwaj, and Wei D.Q., "Evaluation and validation of synergistic effects of amyloid-beta inhibitor—gold nanoparticles complex on Alzheimer's disease using deep neural network approach". Journal of Materials Research, 2019. 34(11): p. 1845-1853.
- 172. (Cited 16 times) Wang, J.-F., Wei D.Q., H.-L. Du, Y.-X. Li, and K.-C. Chou, "Molecular modeling studies on NADP-dependence of Candida tropicalis strain xylose reductase". The Open Bioinformatics Journal, 2008. 2(1): p. 72-79.
- 173. (Cited 16 times) Wei, D.-Q., G. Selvaraj, and A.C. Kaushik, "Computational Perspective on the Current State of the Methods and New Challenges in Cancer Drug Discovery". Current pharmaceutical design, 2018. 24(32): p. 3725-3726.
- 174. (Cited 16 times) Kaushik, A.C., A. Kumar, A.U. Rehman, M. Junaid, A. Khan, S. Bharadwaj, S. Sahi, and Wei D.Q., "Deciphering G-Protein-Coupled Receptor 119 Agonists as Promising Strategy against Type 2 Diabetes Using Systems Biology Approach". ACS Omega, 2018. 3(12): p. 18214-18226.
- 175. (Cited 15 times) Qi, Y.-Y., T. Zhang, Y. Cheng, X.-R. Chen, Wei D.Q., and L.-C. Cai, "Lattice dynamics and thermal conductivity of calcium fluoride via first-principles investigation". Journal of Applied Physics, 2016. 119(9): p. 095103-095110.
- 176. (Cited 15 times) Khan, A., A. Ali, M. Junaid, C. Liu, A.C. Kaushik, W.C. Cho, and Wei D.Q., "Identification of novel drug targets for diamond-blackfan anemia based on RPS19 gene mutation using protein-protein interaction network". BMC systems biology, 2018. 12(4): p. 39-51.
- 177. (Cited 15 times) Ding, S., Tian, Y., Cai, P., Zhang, D., Cheng, X., Sun, D., Yuan, L., Chen, J., Tu, W., Wei, D.Q. and Hu, Q.N., "novoPathFinder: a webserver of designing novel-pathway with integrating GEM-model". Nucleic Acids Research, 2020.
- 178. (Cited 15 times) Li, C.-D., Q. Xu, R.-X. Gu, J. Qu, and Wei D.Q., "The dynamic binding of cholesterol to the multiple sites of C99: as revealed by coarse-grained and all-atom simulations". Physical Chemistry Chemical Physics, 2017. 19(5): p. 3845-3856.
- 179. (Cited 15 times) Junaid, M., M. Shah, A. Khan, C.-D. Li, M.T. Khan, A.C. Kaushik, A. Ali, A. Mehmood, A.S. Nangraj, and S. Choi, "Structural-dynamic insights into the H. pylori cytotoxin-associated gene A (CagA) and its abrogation to interact with the tumor suppressor protein ASPP2 using decoy peptides". Journal of Biomolecular Structure and Dynamics, 2019. 37(15): p. 4035-4050.
- 180. (Cited 14 times) Lian, P., L.A. Liu, Y. Shi, Y. Bu, and Wei D.Q., "Tethered-hopping model for protein-DNA binding and unbinding based on Sox2-Oct1-Hoxb1 ternary complex simulations". Biophysical journal, 2010. 98(7): p. 1285-1293.
- 181. (Cited 14 times) Wang, J., C. Zhang, Wei D.Q., and Y. Li, "Docking and molecular dynamics studies on CYP2D6". Chinese Science Bulletin, 2010. 55(18): p. 1877-1880.
- 182. (Cited 14 times) Ursenbach, C., Wei D.Q., and G. Patey, "Activity coefficients of model aqueous electrolyte solutions: Sensitivity to the short range part of the interionic potential". The Journal of chemical physics, 1991. 94(10): p. 6782-6784.
- 183. (Cited 14 times) Zhang, X.-Q., X.-R. Chen, S. Kaliamurthi, G. Selvaraj, G.-F. Ji, and Wei D.Q., "Initial Decomposition of the Co-crystal of CL-20/TNT: Sensitivity Decrease under Shock Loading". The Journal of Physical Chemistry C, 2018. 122(42): p. 24270-24278.
- 184. (Cited 14 times) Mohammad, Anwar, et al. "Remdesivir MD Simulations Suggest a More Favourable Binding to SARS-CoV-2 RNA Dependent RNA Polymerase Mutant P323L Than Wild-Type." *Biomolecules* 11.7 (2021): 919.
- 185. (Cited 13 times)ul Qamar, Muhammad Tahir, et al. "Structural probing of HapR to identify potent phytochemicals to control Vibrio cholera through integrated computational approaches." *Computers in Biology and Medicine* (2021): 104929.
- 186. (Cited 13 times) Lian, P., J. Li, D.-Q. Wang, and Wei D.Q., "Car–Parrinello molecular dynamics/molecular mechanics (CPMD/MM) simulation study of coupling and uncoupling

- mechanisms of cytochrome P450cam". The Journal of Physical Chemistry B, 2013. 117(26): p. 7849-7856.
- 187. (Cited 13 times) Ye, Y., Y. Luo, C.F. Wang, D.Q. Wei, L.Z. Liu, and Y.F. Zhao, "Cycloaddition reaction of phosphonyl nitrile oxides to phosphaacetylene and alkene". Heteroatom Chemistry: An International Journal of Main Group Elements, 2009. 20(2): p. 95-100.
- 188. (Cited 13 times) Khan, S., Ali, S.S., Zaheer, I., Saleem, S., Ziaullah, Zaman, N., Iqbal, A., Suleman, M., Wadood, A., Rehman, A.U. and Khan, "Proteome-wide mapping and reverse vaccinology-based B and T cell multi-epitope subunit vaccine designing for immune response reinforcement against Porphyromonas gingivalis". Journal of Biomolecular Structure and Dynamics, 2020. p.1-15.
- 189. (Cited 13 times) Bai, Y., Z. Yu, R. Liu, N. Li, S. Yan, K. Yang, B. Liu, Wei D.Q., and L. Wang, "Pressure-Induced Crystallization and Phase Transformation of Para-xylene". Scientific reports, 2017. 7(1): p. 5321-5331.
- 190. (Cited 13 times) Wang, Y., X. Wang, Y. Xiong, C.-D. Li, Q. Xu, L. Shen, A. Chandra Kaushik, and Wei D.Q., "An integrated pan-cancer analysis and structure-based virtual screening of GPR15". International Journal of Molecular Sciences, 2019. 20(24): p. 6226.
- 191. (Cited 13 times) Kaliamurthi, S., A. Demir-Korkmaz, G. Selvaraj, E. Gokce-Polat, Y.-K. Wei, M.A. Almessiere, A. Baykal, K. Gu, and Wei D.Q., "Viewing the Emphasis on State-of-the-Art Magnetic Nanoparticles: Synthesis, Physical Properties, and Applications in Cancer Theranostics". Current pharmaceutical design, 2019.p.1505-15023.
- 192. (Cited 13 times) Li, K., C. Xu, Y. Du, M. Junaid, A.C. Kaushik, and D.Q. Wei, "Comprehensive epigenetic analyses reveal master regulators driving lung metastasis of breast cancer". Journal of Cellular and Molecular Medicine, 2019.p.5415-5431.
- 193. (Cited 12 times) Cui, H.-L., X.-R. Chen, G.-F. Ji, and Wei D.Q., "Structures and Phase Transition of GaAs under Pressure". Chinese Physics Letters, 2008. 25(6): p. 2169-2172.
- 194. (Cited 12 times) Wang, X., X. Zhu, M. Ye, Y. Wang, C. Li, Y. Xiong, and Wei D.Q., "STS-NLSP: a network-based label space partition method for predicting the specificity of membrane transporter substrates using a hybrid feature of structural and semantic similarity". Front. Bioeng. Biotechnol. 7: 306. DOI 10.3389/fbioe, 2019.p.306-319.
- 195. (Cited 12 times) Kaushik, A.C., Mehmood, A., Dai, X. and Wei, D.Q., "A comparative chemogenic analysis for predicting Drug-Target Pair via Machine Learning Approaches". Scientific Reports, 2020. 10(1), pp.1-11.
- 196. (Cited 12 times) Wang, Z.Y., Han, Q.Q., Deng, M.Y., Zhao, M.J., Apryani, E., Shoaib, R.M., Wei, D.Q. and Wang, Y.X., "Lemairamin, isolated from the Zanthoxylum plants, alleviates pain hypersensitivity via spinal α7 nicotinic acetylcholine receptors". Biochemical and Biophysical Research Communications, 2020.
- 197. (Cited 12 times) Du, Q., Wei D.Q., and K.-C. Chou, "Correlations of amino acids in proteins". Peptides, 2003. 24(12): p. 1863-1869.
- 198. (Cited 12 times) Cui, H.-L., G.-F. Ji, J.-J. Zhao, F. Zhao, X.-R. Chen, Q.-M. Zhang, and Wei D.Q., "Ab initio and molecular dynamics studies of solid β-HMX: effects of hydrostatic pressure and high temperature". Molecular Simulation, 2010. 36(9): p. 670-681.
- 199. (Cited 12 times) Cui, S., Wei D.Q., H. Hu, and Z. Gong, "Mechanical instability and ideal strengths of layered M2SC (M=Ti, Zr, and Hf) compounds". Journal of Applied Physics, 2013. 113(8): p. 083516-083523.
- 200. (Cited 12 times) Pan, C.T., Wang, S.Y., Yen, C.K., Kumar, A., Kuo, S.W., Zheng, J.L., Wen, Z.H., Singh, R., Singh, S.P., Khan, M.T. and Chaudhary, R.K., "Polyvinylidene Fluoride-Added Ceramic Powder Composite Near-Field Electrospinned Piezoelectric Fiber-Based Low-Frequency Dynamic Sensors". ACS omega, 2020. 5(28), p.17090-17101.
- 201. (Cited 12 times) Gul, H., Ali, S.S., Saleem, S., Khan, S., Khan, J., Wadood, A., Rehman, A.U., Ullah, Z., Ali, S., Khan, H. and Hussain, Z, "Subtractive proteomics and immunoinformatics approaches to explore Bartonella bacilliformis proteome (virulence factors) to design B and T cell multi-epitope subunit vaccine. Infection". Genetics and Evolution, 2020. p.104551.
- 202. (Cited 12 times) Khan, Abbas, et al. "Hantavirus: The Next Pandemic We Are Waiting For?." *Interdisciplinary Sciences: Computational Life Sciences* 13.1 (2021): 147-152.

- 203. (Cited 11 times) Stalin, Antony, et al. "An in-silico approach to identify the potential hot spots in SARS-CoV-2 spike RBD to block the interaction with ACE2 receptor." *Journal of Biomolecular Structure and Dynamics* (2021): 1-16.
- 204. (Cited 11 times) Zhang, Y., G. Ji, F. Zhao, Z. Gong, Wei D.Q., L. Chen, and W. Li, "Mesoscopic simulation of aggregate behaviour of fluoropolymers in the TATB-based PBX". Molecular Simulation, 2011. 37(03): p. 237-242.
- 205. (Cited 11 times) Huang, Siyuan, et al. "Circulating miR-1246 targeting UBE2C, TNNI3, TRAIP, UCHL1 genes and key pathways as a potential biomarker for lung adenocarcinoma: integrated biological network analysis." *Journal of Personalized Medicine* 10.4 (2020): 162.
- 206. (Cited 11 times) Wei, Y.K., Jia, L.Q., Fang, Y.Y., Wang, L.J., Qian, Z.X., Yuan, J.N., Selvaraj, G., Ji, G.F. and Wei, D.Q., "Formation and superconducting properties of predicted ternary hydride ScYH6 under pressures". International Journal of Quantum Chemistry, 2020. p.e26459.
- 207. (Cited 11 times)Suleman, Muhammad, et al. "Bioinformatics analysis of the differences in the binding profile of the wild-type and mutants of the SARS-CoV-2 spike protein variants with the ACE2 receptor." *Computers in Biology and Medicine* (2021): 104936.
- 208. (Cited 11 times) Du, Y., K. Li, X. Wang, K.C. Aman, M. Junaid, and Wei D.Q., "Identification of chlorprothixene as a potential drug that induces apoptosis and autophagic cell death in acute myeloid leukemia cells". The FEBS Journal, 2019.p.1-25.
- 209. (Cited 11 times) Suleman, Muhammad, et al. "Bioinformatics analysis of the differences in the binding profile of the wild-type and mutants of the SARS-CoV-2 spike protein variants with the ACE2 receptor." *Computers in Biology and Medicine* 138 (2021): 104936.
- 210. (Cited 11 times) Wei, Y.K., Jia, L.Q., Fang, Y.Y., Wang, L.J., Qian, Z.X., Yuan, J.N., Selvaraj, G., Ji, G.F. and Wei, D.Q., "Formation and superconducting properties of predicted ternary hydride ScYH6 under pressures". International Journal of Quantum Chemistry, 2020. p.e26459.
- 211. (Cited 11 times) Lin, Shenggeng, et al. "MDF-SA-DDI: predicting drug—drug interaction events based on multi-source drug fusion, multi-source feature fusion and transformer self-attention mechanism." *Briefings in Bioinformatics* (2021).
- 212. (Cited 11 times) Stalin, Antony, et al. "An in-silico approach to identify the potential hot spots in SARS-CoV-2 spike RBD to block the interaction with ACE2 receptor." *Journal of Biomolecular Structure and Dynamics* (2021): 1-16.
- 213. (Cited 10 times) Wei, D., "Field-induced phase transitions of simple dipolar fluids". Physical Review E, 1994. 49(3): p. 2454-2456.
- 214. (Cited 10 times) Wang, Y., X. Wang, Y. Xiong, A.C. Kaushik, J. Muhammad, A. Khan, H. Dai, and Wei D.Q., "New strategy for identifying potential natural HIV-1 non-nucleoside reverse transcriptase inhibitors against drug-resistance: an in silico study". Journal of Biomolecular Structure and Dynamics, 2019: p. 1-15.
- 215. (Cited 10 times) Wang, X., Liu, Y., Lu, F., Li, H., Gao, P. and Wei, D.Q., "Dipeptide Frequency of Word Frequency and Graph Convolutional Networks for DTA Prediction". Frontiers in Bioengineering and Biotechnology, 2020, 8, p.267.
- 216. (Cited 10 times) Ahmad, Irfan, et al. "Development of multi-epitope subunit vaccine for protection against the norovirus' infections based on computational vaccinology." *Journal of Biomolecular Structure and Dynamics* (2020): 1-12.
- 217. (Cited 10 times) Kaushik, A.C., A. Mehmood, A.K. Upadhyay, S. Paul, S. Srivastava, P. Mali, Y. Xiong, X. Dai, Wei D.Q., and S. Sahi, "CytoMegaloVirus Infection Database: A Public Omics Database for Systematic and Comparable Information of CMV". Interdisciplinary Sciences: Computational Life Sciences, 2019: p. 1-9.
- 218. (Cited 10 times) Ma, L.-N., Z.-Z. Du, P. Lian, and Wei D.Q., "A theoretical study on the mechanism of a superficial mutation inhibiting the enzymatic activity of CYP1A2". Interdisciplinary Sciences: Computational Life Sciences, 2014. 6(1): p. 25-31.
- 219. (Cited 10 times) Mehmood, A., M.T. Khan, A.C. Kaushik, A.S. Khan, M. Irfan, and Wei D.Q., "Structural dynamics behind clinical mutants of PncA-Asp12Ala, Pro54Leu, and His57Pro of Mycobacterium tuberculosis associated with pyrazinamide resistance". Frontiers in Bioengineering and Biotechnology, 2019. 7: p. 404.
- 220. (Cited 10 times) Kaushik, A.C., D. Gautam, A.S. Nangraj, Wei D.Q., and S. Sahi, "Protection of Primary Dopaminergic Midbrain Neurons Through Impact of Small Molecules Using Virtual

- Screening of GPR139 Supported by Molecular Dynamic Simulation and Systems Biology". Interdisciplinary Sciences: Computational Life Sciences, 2019: p. 1-11.
- 221. (Cited 10 times) Khan, M.T., Ali, S., Zeb, M.T., Kaushik, A.C., Malik, S.I. and Wei, D.Q., "Gibbs free energy calculation of mutation in PncA and RpsA associated with pyrazinamide resistance". Frontiers in Molecular Biosciences, 2020. 7.
- 222. (Cited 9 times) Kaushik, A.C., Wang, Y.J., Wang, X. and Wei, D.Q, "Irinotecan and vandetanib create synergies for treatment of pancreatic cancer patients with concomitant TP53 and KRAS mutations". Briefings in Bioinformatics, 2020.
- 223. (Cited 9 times) Zeb, Adnan, et al. "Genome-wide screening of vaccine targets prioritization and reverse vaccinology aided design of peptides vaccine to enforce humoral immune response against Campylobacter jejuni." *Computers in Biology and Medicine* 133 (2021): 104412.
- 224. (Cited 9 times) Huo, J., A. Murray, and Wei D.Q., "Adaptive visual and auditory map alignment in barn owl superior colliculus and its neuromorphic implementation". IEEE transactions on neural networks and learning systems, 2012. 23(9): p. 1486-1497.
- 225. (Cited 9 times) Xie, Z., T. Zhang, J.-F. Wang, K.-C. Chou, and Wei D.Q., "The computational model to predict accurately inhibitory activity for inhibitors towardsCYP3A4". Computers in biology and medicine, 2010. 40(11-12): p. 845-852.
- 226. (Cited 9 times) Dai, H., Q. Xu, Y. Xiong, W.-L. Liu, and Wei D.Q., "Improved prediction of michaelis constants in CYP450-mediated reactions by resilient back propagation algorithm". Current drug metabolism, 2016. 17(7): p. 673-680.
- 227. (Cited 9 times) Lian, P., H.-B. Guo, J.C. Smith, Wei D.Q., and H. Guo, "Catalytic mechanism and origin of high activity of cellulase TmCel12A at high temperature: a quantum mechanical/molecular mechanical study". Cellulose, 2014. 21(2): p. 937-949.
- 228. (Cited 9 times) Bai, Y., N. Li, C. Pei, Z. Yan, W. Li, and Wei D.Q., "High-pressure transformations of ortho-xylene probed by combined infrared and Raman spectroscopies". Solid State Communications, 2018. 269: p. 96-101.
- 229. (Cited 9 times) Li, L., Y. Xiong, Z.-Y. Zhang, Q. Guo, Q. Xu, H.-H. Liow, Y.-H. Zhang, and Wei D.Q., "Improved feature-based prediction of SNPs in human cytochrome P450 enzymes". Interdisciplinary Sciences: Computational Life Sciences, 2015. 7(1): p. 65-77.
- 230. (Cited 9 times) Quimque, Mark Tristan, et al. "Potential Cancer-and Alzheimer's Disease-Targeting Phosphodiesterase Inhibitors from Uvaria alba: Insights from In Vitro and Consensus Virtual Screening." *ACS omega* 6.12 (2021): 8403-8417.
- 231. (Cited 9 times) Li, K., Y. Du, Wei D.Q., and F. Zhang, "CEBPE expression is an independent prognostic factor for acute myeloid leukemia". Journal of translational medicine, 2019. 17(1): p. 188-199.
- 232. (Cited 9 times) Kaushik, A.C., A. Mehmood, S. Peng, Y.-J. Zhang, X. Dai, and Wei D.Q., "A-CaMP: a tool for anti-cancer and antimicrobial peptide generation". Journal of Biomolecular Structure and Dynamics, 2020: p. 1-9.
- 233. (Cited 9 times) Li, D., Y. Gao, X. Pan, Wei D.Q., B. Guo, C. Yang, and B. Liu, "MD and DSC study of bioactive structural stability of insulin in various imidazolium ionic liquids". Journal of Molecular Liquids, 2019. 277: p. 971-976.
- 234. (Cited 9 times) Mou, Kejie, et al. "Emerging mutations in envelope protein of SARS-CoV-2 and their effect on thermodynamic properties." *Informatics in medicine unlocked* 25 (2021): 100675.
- 235. (Cited 9 times) Selvaraj, Gurudeeban, et al. "Identifying potential drug targets and candidate drugs for COVID-19: biological networks and structural modeling approaches." *F1000Research* 10 (2021).
- 236. (Cited 9 times)Dai, Qiuying, et al. "MDA-CF: Predicting MiRNA-Disease associations based on a cascade forest model by fusing multi-source information." *Computers in Biology and Medicine* 136 (2021): 104706.
- 237. (Cited 9 times)Mou, Kejie, et al. "Emerging Mutations in Nsp1 of SARS-CoV-2 and Their Effect on the Structural Stability." *Pathogens* 10.10 (2021): 1285.
- 238. (Cited 8 times) Zhang, Xueting, et al. "Semiconductor-to-metal transition in HfSe2 under high pressure." *Journal of Alloys and Compounds* 867 (2021): 158923.

- 239. (Cited 8 times) Inayat, Nagina, et al. "iEnhancer-DHF: Identification of Enhancers and Their Strengths Using Optimize Deep Neural Network With Multiple Features Extraction Methods." *IEEE Access* 9 (2021): 40783-40796.
- 240. (Cited 8 times) Kaliamurthi, Satyavani, et al. "Structure-based virtual screening reveals Ibrutinib and Zanubrutinib as potential repurposed drugs against COVID-19." *International Journal of Molecular Sciences* 22.13 (2021): 7071.
- 241. (Cited 8 times) Khan, A., Umbreen, S., Hameed, A., Fatima, R., Zahoor, U., Babar, Z., Waseem, M., Hussain, Z., Rizwan, M., Zaman, N. and Ali, S., "In Silico Mutagenesis Based Remodeling of SARs-CoV-1 Peptide (ATLQAIAS) to Inhibit SARs-CoV-2: Structural-dynamics and Free Energy Calculations", 2020.
- 242. (Cited 8 times) Humayun, Fahad, et al. "Abrogation of SARS-CoV-2 interaction with host (NRP1) Neuropilin-1 receptor through high-affinity marine natural compounds to curtail the infectivity: A structural-dynamics data." *Computers in Biology and Medicine* (2021): 104714.
- 243. (Cited 8 times) Khan, Abbas, et al. "In Silico Mutagenesis-Based Remodelling of SARS-CoV-1 Peptide (ATLQAIAS) to Inhibit SARS-CoV-2: Structural-Dynamics and Free Energy Calculations." *Interdisciplinary Sciences: Computational Life Sciences* 13.3 (2021): 521-534.
- 244. (Cited 8 times) Dong, E., Liu, B., Dong, Q., Shi, X., Ma, X., Liu, R., Zhu, X., Luo, X., Li, X., Li, Y. and Li, Q., "Effects of pressure on the structure and properties of layered ferromagnetic Cr2Ge2Te6". Physica B: Condensed Matter, 2020. 595, p.412344.
- 245. (Cited 8 times) Khan, Muhammad Tahir, et al. "SARS-CoV-2 Genome from the Khyber Pakhtunkhwa Province of Pakistan." *ACS omega* 6.10 (2021): 6588-6599.
- 246. (Cited 8 times) Kaushik, A.C., Mehmood, A., Wei, D.Q. and Dai, X., "Systems biology integration and Screening of reliable prognostic markers to create synergies in the control of lung cancer patients". Frontiers in Molecular Biosciences, 2020, 7.
- 247. (Cited 8 times) Li-Wei, D., Z. Ji-Jun, J. Guang-Fu, G. Zi-Zheng, and W. Dong-Qing, "First-principles study of orthorhombic perovskites MgSiO3 up to 120 GPa and its geophysical implications". Chinese Physics Letters, 2006. 23(8): p. 2334-2337.
- 248. (Cited 8 times) Chen, Q., X. Cheng, Wei D.Q., and Q. Xu, "Molecular dynamics simulation studies of the wild type and E92Q/N155H mutant of Elvitegravir-resistance HIV-1 integrase". Interdisciplinary Sciences: Computational Life Sciences, 2015. 7(1): p. 36-42.
- 249. (Cited 8 times) Zhang, P., F. Meng, Z. Gong, G. Ji, S. Cui, and Wei D.Q., "First-principles study of structure and properties of ω-Ti2Zr". Computational Materials Science, 2013. 74: p. 129-137.
- 250. (Cited 8 times) Yuan, J., G. Ji, X. Chen, Wei D.Q., F. Zhao, and Q. Wu, "Phase transition, thermodynamics properties and IR spectrum of α-and γ-RDX: First principles and MD studies". Chemical Physics Letters, 2016. 644: p. 250-254.
- 251. (Cited 8 times) Li, W., B. Yang, Y. Wang, Wei D.Q., C. Whiteley, and X. Wang, "Molecular modeling of substrate selectivity of Candida antarctica lipase B and Candida rugosa lipase towards c9, t11-and t10, c12-conjugated linoleic acid". Journal of Molecular Catalysis B: Enzymatic, 2009. 57(1-4): p. 299-303.
- 252. (Cited 8 times) Guan, X., Wei D.Q., and D. Hu, "Free Energy Calculation of Transmembrane Ion Permeation: Sample with a Single Reaction Coordinate and Analysis along Transition Path". Journal of chemical theory and computation, 2019. 15(2): p. 1216-1225.
- 253. (Cited 8 time) Li, C.-D., M. Junaid, H. Chen, A. Ali, and Wei D.Q., "Helix-Switch Enables C99 Dimer Transition between the Multiple Conformations". Journal of chemical information and modeling, 2018. 59(1): p. 339-350.
- 254. (Cited 7 times) Junaid, M., Li, C.D., Li, J., Khan, A., Ali, S.S., Jamal, S.B., Saud, S., Ali, A. and Wei, D.Q., "Structural insights of catalytic mechanism in mutant pyrazinamidase of Mycobacterium tuberculosis". Journal of Biomolecular Structure and Dynamics, 2020. p.1-14.
- 255. (Cited 7 times) Lodhi, Madeeha Shahzad, et al. "A novel formulation of theranostic nanomedicine for targeting drug delivery to gastrointestinal tract cancer." *Cancer Nanotechnology* 12.1 (2021): 1-27.
- 256. (Cited 7 times) Mehmood, Aamir, et al. "Bringing Structural Implications and Deep Learning-Based Drug Identification for KRAS Mutants." *Journal of Chemical Information and Modeling* 61.2 (2021): 571-586.

- 257. (Cited 7 times) Wei, Y.-K., X.-M. Zhao, M.-M. Li, J.-X. Yu, S. Gurudeeban, Y.-F. Hu, G.-F. Ji, and Wei D.Q., "Detoxification of aflatoxins on prospective approach: effect on structural, mechanical, and optical properties under pressures". Interdisciplinary Sciences: Computational Life Sciences, 2018. 10(2): p. 311-319.
- 258. (Cited 7 times) Shi, Zhiwen, et al. "Prediction of blood-brain barrier permeability of compounds by fusing resampling strategies and eXtreme gradient boosting." *IEEE Access* 9 (2020): 9557-9566.
- 259. (Cited 7 times) Zhang, Y., G. Ji, Z. Gong, and Wei D.Q., "New coupling mechanism of the silane coupling agents in the TATB-based PBX". Molecular Simulation, 2013. 39(5): p. 423-427
- 260. (Cited 7 times) Gao, J., L. Li, X. Wu, and Wei D.Q., "BioNetSim: a Petri net-based modeling tool for simulations of biochemical processes". Protein & cell, 2012. 3(3): p. 225-229.
- 261. (Cited 7 times) Yu, Y., X. Liu, J. He, M. Zhang, H. Li, Wei D.Q., and Y. Song, "Appendant structure plays an important role in amyloidogenic cystatin dimerization prior to domain swapping". Journal of Biomolecular Structure and Dynamics, 2012. 30(1): p. 102-112.
- 262. (Cited 7 times) Xu, K., Wei D.Q., X.-R. Chen, and G.-F. Ji, "Thermal decomposition of solid phase nitromethane under various heating rates and target temperatures based on ab initio molecular dynamics simulations". Journal of molecular modeling, 2014. 20(10): p. 2438-2448.
- 263. (Cited 7 times) Yuan, J.-N., Y.-K. Wei, X.-Q. Zhang, X.-R. Chen, G.-F. Ji, M.K. Kotni, and Wei D.Q., "Shock response of 1, 3, 5-trinitroperhydro-1, 3, 5-triazine (RDX): The CN bond scission studied by molecular dynamics simulations". Journal of Applied Physics, 2017. 122(13): p. 135901-135910.
- 264. (Cited 7 times) Lv, H.-M., X.-L. Guo, R.-X. Gu, and Wei D.Q., "Free energy calculations and binding analysis of two potential anti-influenza drugs with Polymerase basic protein-2 (PB2)". Protein and peptide letters, 2011. 18(10): p. 1002-1009.
- 265. (Cited 7 times) Junaid, M., C.D. Li, M. Shah, A. Khan, H. Guo, and D.Q. Wei, "Extraction of molecular features for the drug discovery targeting protein-protein interaction of Helicobacter pylori CagA and tumor suppressor protein ASSP2". Proteins: Structure, Function, and Bioinformatics, 2019.p. 837-849.
- 266. (Cited 6 times) Wang, J.-F., Wei D.Q., C.-F. Wang, Y. Ye, Y.-X. Li, Y. Luo, W.-W. Wang, L.-Z. Liu, and Y.-F. Zhao, "A theoretical study on the mechanism of 2: 11, 3 dipolar cycloaddition reactions". Journal of Theoretical and Computational Chemistry, 2007. 6(04): p. 861-867.
- 267. (Cited 6 times) Yao, Y., T. Zhang, Y. Xiong, L. Li, J. Huo, and D.Q. Wei, "Mutation probability of cytochrome P450 based on a genetic algorithm and support vector machine". Biotechnology journal, 2011. 6(11): p. 1367-1376.
- 268. (Cited 6 times) Zhu, J., Y. Cheng, T.-C. Bai, Y. Lu, Z. Chang, Wei D.Q., and G. Stell, "Solvent dynamics effect in condensed-phase electron-transfer reactions". The Journal of Physical Chemistry B, 2008. 112(12): p. 3735-3745.
- 269. (Cited 6 times) Selvaraj, G., C. Selvaraj, and Wei D.Q., "Computational Advances in Chronic Diseases Diagnostics and Therapy-II". Current Drug Targets, 2020. 21(2): p. 103-104.
- 270. (Cited 6 times) Ali, A., M. Junaid, A. Khan, A. Kaushik, and A. Mehmood, "Identification of Novel Therapeutic Targets in Myelodysplastic Syndrome Using Protein-Protein Interaction Approach and Neural Networks". J Comput Sci Syst Biol, 2018. 11: p. 184-189.
- 271. (Cited 6 time) Khan, F.I., K. Bisetty, K.-R. Gu, S. Singh, K. Permaul, M.I. Hassan, and Wei D.Q., "Molecular dynamics simulation of chitinase I from Thermomyces lanuginosus SSBP to ensure optimal activity". Molecular Simulation, 2017. 43(7): p. 480-490.
- 272. (Cited 6 times) Kaushik, A.C., Y.-J. Wang, X. Wang, A. Kumar, S.P. Singh, C.-T. Pan, Y.-L. Shiue, and Wei D.Q., "Evaluation of anti-EGFR-iRGD recombinant protein with GOLD nanoparticles: synergistic effect on antitumor efficiency using optimized deep neural networks". RSC Advances, 2019. 9(34): p. 19261-19270.
- 273. (Cited 6 times) Dai, H., W. Wang, Q. Xu, Y. Xiong, and Wei D.-Q., "Estimation of Probability Distribution and Its Application in Bayesian Classification and Maximum Likelihood Regression". Interdisciplinary Sciences: Computational Life Sciences, 2019. 11(3): p. 559-574.
- 274. (Cited 6 times)Suleman, Muhammad, et al. "Immunoinformatics and Immunogenetics-Based Design of Immunogenic Peptides Vaccine against the Emerging Tick-Borne Encephalitis Virus

- (TBEV) and Its Validation through In Silico Cloning and Immune Simulation." *Vaccines* 9.11 (2021): 1210.
- 275. (Cited 6 times) Khan, Taimoor, et al. "CytomegaloVirusDb: multi-Omics knowledge database for Cytomegaloviruses." *Computers in Biology and Medicine* (2021): 104563.
- 276. (Cited 6 times) Jiang, Mingming, et al. "NeuroPpred-Fuse: an interpretable stacking model for prediction of neuropeptides by fusing sequence information and feature selection methods." *Briefings in Bioinformatics* (2021).
- 277. (Cited 6 times) Li, Zonglun, et al. "Pressure-Tailored Band Engineering for Significant Enhancements in the Photoelectric Performance of CsI3 in the Optical Communication Waveband." *Advanced Functional Materials* (2021): 2108636.
- 278. (Cited 6 times) Shah, Abdullah, et al. "Comparative mutational analysis of SARS-CoV-2 isolates from Pakistan and structural-functional implications using computational modelling and simulation approaches." *Computers in biology and medicine* 141 (2022): 105170.
- 279. (Cited 5 times) Chen, H.F., M.Y. Wu, Z. Wang, and D.Q. Wei, "Insight into the metabolism rate of quinone analogues from molecular dynamics simulation and 3D-QSMR methods". Chemical biology & drug design, 2007. 70(4): p. 290-301.
- 280. (Cited 5 times) Li, L., Wei D.Q., J.-F. Wang, and K.-C. Chou, "SCYPPred: a web-based predictor of SNPs for human cytochrome P450". Protein and peptide letters, 2012. 19(1): p. 57-61.
- 281. (Cited 5 times) Selvaraj, C., G. Selvaraj, S. Kaliamurthi, W.C. Cho, Wei D.Q., and S.K. Singh, "Ion Channels as Therapeutic Targets for Type 1 Diabetes Mellitus". Current drug targets, 2019. 20: p. 132-147.
- 282. (Cited 5 times) Li, F., C. Chen, S.-Y. Qu, M.-Z. Zhao, X. Xie, X. Wu, L. Li, X. Wang, Q. Ding, and Q. Xu, "The Disulfide Bond between Cys22 and Cys27 in the Protease Domain Modulate Clotting Activity of Coagulation Factor X". Thrombosis and haemostasis, 2019. 119(06): p. 871-881.
- 283. (Cited 5 times) Selvaraj, Chandrabose, et al. "Interrogation of Bacillus anthracis SrtA active site loop forming open/close lid conformations through extensive MD simulations for understanding binding selectivity of SrtA inhibitors." *Saudi Journal of Biological Sciences* (2021).
- 284. (Cited 5 times) Celik, Ismail, et al. "Computational prediction of the effect of mutations in the receptor-binding domain on the interaction between SARS-CoV-2 and human ACE2." Molecular Diversity (2022).
- 285. (Cited 5 times) Ren, H.-C., J.-N. Yuan, W.-S. Xu, T.-N. Chen, G.-F. Ji, and Wei D.Q., "Two-Dimensional Infrared Spectra of Cationic Dopamine under Different Electric Fields: Theoretical Studies from the Density Function Theory Anharmonic Potential". The Journal of Physical Chemistry C, 2018. 122(31): p. 17994-18004.
- 286. (Cited 5 times)Kaushik, Aman Chandra, et al. "CoronaPep: An Anti-coronavirus Peptide Generation Tool." *IEEE/ACM Transactions on Computational Biology and Bioinformatics* (2021).
- 287. (Cited 5 times) Ali, Sharafat, et al. "Proteome wide vaccine targets prioritization and designing of antigenic vaccine candidate to trigger the host immune response against the Mycoplasma genitalium infection." *Microbial Pathogenesis* 152 (2021): 104771.
- 288. (Cited 5 times) Khan, M.T., Kaushik, A.C., ul ain Rana, Q., Malik, S.I., Khan, A.S., Wei, D.Q., Sajjad, W., Ahmad, S., Ali, S. and Irfan, M., "Characterization and synthetic biology of lipase from Bacillus amyloliquefaciens strain". Archives of Microbiology, 2020. pp.1-10.
- 289. (Cited 5 times) Khan, M., Iqbal, N., Khan, S., Muhammad Khan, D., Khan, A. and Wei, D.Q, "Prediction of Recombination Spots Using Novel Hybrid Feature Extraction Method via Deep Learning Approach". Frontiers in Genetics, 2020. 11, p.1052.
- 290. (Cited 5 times) Khan, Muhammad Tahir, et al. "Insight into the drug resistance whole genome of Mycobacterium tuberculosis isolates from Khyber Pakhtunkhwa, Pakistan." *Infection, Genetics and Evolution* 92 (2021): 104861.

- 291. (Cited 5 times) Ali, A., Khan, M.T., Khan, A., Ali, S., Chinnasamy, S., Akhtar, K., Shafiq, A. and Wei, D.Q., "Pyrazinamide resistance of novel mutations in pncA and their dynamic behavior". RSC Advances, 2020. 10(58), pp.35565-35573.
- 292. (Cited 5 times)El Khatabi, Khalil, et al. "Identification of novel acetylcholinesterase inhibitors through 3D-QSAR, molecular docking, and molecular dynamics simulation targeting Alzheimer's disease." *Journal of Molecular Modeling* 27.10 (2021): 1-13.
- 293. (Cited 5 times) Zhao, M., H.-T. Chang, Q. Zhou, T. Zeng, C.-S. Shih, Z.-P. Liu, L. Chen, and Wei D.Q., "Predicting protein-ligand interactions based on chemical preference features with its application to new D-amino acid oxidase inhibitor discovery". Current pharmaceutical design, 2014. 20(32): p. 5202-5211.
- 294. (Cited 4 times) Bai, Y., D. Wang, Z.-X. Yu, Y. Jia, F.-Y. Zhu, Wei D.Q., W.-Z. Zhong, and K.-C. Chou, "Ecdysterone determination of Niuxi by near-infrared diffuse reflection spectroscopy". Spectroscopy, 2006. 21(5).
- 295. (Cited 4 times) Wei, D. and L. Blum, "Solvation thermodynamic functions in the mean spherical approximation: Behavior near the solvent critical region". The Journal of chemical physics, 1995. 102(10): p. 4217-4226.
- 296. (Cited 4 times) Wei, D.-Q., Y.-J. Wang, L. Wang, J.-H. Hu, Z.-Z. Gong, Y.-X. Guo, and Y.-S. Zhu, "Molecular dynamics simulations of dipolar fluids in orientationally ordered phases". Physical Review E, 2007. 75(6): p. 061702-061707.
- 297. (Cited 4 times) Zhang, L., G.-F. Ji, F. Zhao, C.-M. Meng, and Wei D.Q., "The first-principle studies of the crystal phase transitions: Fd3m-MgAl2O4→ F4-3m-MgAl2O4". Physica B: Condensed Matter, 2011. 406(3): p. 335-338.
- 298. (Cited 4 times) Wei, D.-Q., L. Gao, J. Zhang, L.-W. Yan, J.-H. Hu, L. Chen, Z.-Z. Gong, Y.-X. Guo, and Y. Han, "Role of dipole elongation in orientationally ordered liquids". Physical Review E, 2011. 83(6): p. 061703-0161708.
- 299. (Cited 4 times) Cui, S., Wei D.Q., Q. Zhang, Z. Gong, and H. Hu, "Newly found phase transition and mechanical stability of AuAl2: A first-principles study". Journal of Alloys and Compounds, 2013. 574: p. 486-489.
- 300. (Cited 4 times) Yang, H.-F., X.-N. Zhang, Y. Li, Y.-H. Zhang, Q. Xu, and Wei D.Q., "Theoretical studies of intracellular concentration of micro-organisms' metabolites". Scientific reports, 2017. 7(1): p. 9048-9059.
- 301. (Cited 4 times) Wei, D., "Orientational Order in Simple Polar Liquids: Effect of the Quadrupole Moment". Molecular Crystals and Liquid Crystals Science and Technology. Section A. Molecular Crystals and Liquid Crystals, 1995. 269(1): p. 89-98.
- 302. (Cited 4 times) Khan, Sohail, et al. "Targeting the N-terminal domain of the RNA-binding protein of the SARS-CoV-2 with high affinity natural compounds to abrogate the protein-RNA interaction: a molecular dynamics study." *Journal of Biomolecular Structure and Dynamics* (2021): 1-9.
- 303. (Cited 4 times) Song, K.-F., G.-F. Ji, K.M. Kumari, and Wei D.Q., "Blending effect between n-decane and toluene in oxidation: a ReaxFF study". Molecular Simulation, 2018. 44(1): p. 21-33.
- 304. (Cited 4 times) Wang, Y.-K., Wei D.Q., R.-X. Gu, H.-M. Fan, and J. Ulmschneider, "Applications of rare event dynamics on the free energy calculations for membrane protein systems". Canadian Journal of Chemistry, 2013. 91(9): p. 769-774.
- 305. (Cited 4 times) Khan, Taimoor, et al. "A computational perspective on the dynamic behaviour of recurrent drug resistance mutations in the pncA gene from Mycobacterium tuberculosis." *RSC Advances* 11.4 (2021): 2476-2486.
- 306. (Cited 4 times) Cai, X., F. Li, H. Lei, S. Qu, C. Qian, D. Xiang, D.Q. Wei, W. Wu, Q. Xu, and X. Wang, "p. R180C mutation of glycosyltransferase B leads to B subgroup, an in vitro and in silico study". Vox sanguinis, 2018. 113(5): p. 476-484.
- 307. (Cited 4 times) Khan, M.T., Chinnasamy, S., Cui, Z., Irfan, M. and Wei, D.Q, "Mechanistic analysis of A46V, H57Y, and D129N in pyrazinamidase associated with pyrazinamide resistance". Saudi Journal of Biological Sciences, 2020.

- 308. (Cited 4 times) Kaliamurthi, S., G. Selvaraj, S. Chinnasamy, Q. Wang, A.S. Nangraj, W.C. Cho, K. Gu, and Wei D.Q., "Immunomics Datasets and Tools: To Identify Potential Epitope Segments for Designing Chimeric Vaccine Candidate to Cervix Papilloma". Data, 2019. 4(1): p. 31-48.
- 309. (Cited 4 times) Guan, X., Wei D.Q., and D. Hu, "Free Energy Calculations on the Water-Chain-Assisted and the Dehydration Mechanisms of Transmembrane Ion Permeation". Journal of chemical theory and computation, 2019.
- 310. (Cited 4 times) Li, D., Lian, X., Xu, Y., Dong, Q., Liu, B., Wei, D. and Guo, B., "The Inhibitory Mechanism of L-Lysine Hydrochloride on the Ice Crystals Growth by Molecular Dynamics". Chemical Physics, 2020. p. 110889.
- 311. (Cited 4 times) Magpantay, Hilbert D., et al. "Antibacterial and COX-2 Inhibitory Tetrahydrobisbenzylisoquinoline Alkaloids from the Philippine Medicinal Plant Phaeanthus ophthalmicus." *Plants* 10.3 (2021): 462.
- 312. (Cited 4 times) Zhou, Deshan, et al. "LUNAR Drug Screening for Novel Coronavirus Based on Representation Learning Graph Convolutional Network." *IEEE/ACM Transactions on Computational Biology and Bioinformatics* (2021).
- 313. (Cited 4 times) Khan, Abbas, et al. "Computational modelling of potentially emerging SARS-CoV-2 spike protein RBDs mutations with higher binding affinity towards ACE2: A structural modelling study." *Computers in biology and medicine* 141 (2022): 105163.
- 314. (Cited 4 times) Khan, Taimoor, Abbas Khan, and Dong-Qing Wei. "MMV-db: vaccinomics and RNA-based therapeutics database for infectious hemorrhagic fever-causing mammarenaviruses." *Database* 2021 (2021).
- 315. (Cited 4 time) Magpantay, Hilbert D., et al. "Antibacterial and COX-2 Inhibitory Tetrahydrobisbenzylisoquinoline Alkaloids from the Philippine Medicinal Plant Phaeanthus ophthalmicus." *Plants* 10.3 (2021): 462.
- 316. (Cited 4 times) Khan, Abbas, et al. "Structural-Dynamics and Binding Analysis of RBD from SARS-CoV-2 Variants of Concern (VOCs) and GRP78 Receptor Revealed Basis for Higher Infectivity." *Microorganisms* 9.11 (2021): 2331.
- 317. (Cited 4 times) Li, D., J. Cai, B. Guo, Wei D.Q., D.Q. Liu, and B. Liu, "Controllable crystallization of urea crystal face (0 0 1) by molecular simulation". Journal of Crystal Growth, 2019.125139-125147.
- 318. (Cited 4 times) Li, Juanying, et al. "Size and morphology effects on the high pressure behaviors of Mn 3 O 4 nanorods." *Nanoscale Advances* 2.12 (2020): 5841-5847.
- 319. (Cited 3 times) Khan, F.I., K. Bisetty, Wei D.Q., and M.I. Hassan, "A pH based molecular dynamics simulations of chitinase II isolated from Thermomyces lanuginosus SSBP". Cogent Biology, 2016. 2(1): p. 1168336-1168346.
- 320. (Cited 3 times)Zheng, Bowen, et al. "Towards an Ensemble Vaccine against the Pegivirus Using Computational Modelling Approaches and Its Validation through In Silico Cloning and Immune Simulation." *Vaccines* 9.8 (2021): 818.
- 321. (Cited 3 times)Khan, Anwar Sheed, et al. "Characterization of rifampicin-resistant Mycobacterium tuberculosis in Khyber Pakhtunkhwa, Pakistan." *Scientific reports* 11.1 (2021): 1-10.
- 322. (Cited 3 time) Fan, P., Q.-H. Lin, Y. Guo, L.-L. Zhao, H. Ning, M.-Y. Liu, and Wei D.Q., "The PPI network analysis of mRNA expression profile of uterus from primary dysmenorrheal rats". Scientific reports, 2018. 8(1): p. 351-361.
- 323. (Cited 3 times) Kaushik, A.C., A. Kumar, C.-Y. Yu, S.-W. Kuo, S.-S. Liang, S.P. Singh, X. Wang, Y.-J. Wang, C.-K. Yen, and X. Dai, "PCL-DOX Macro drops: Evaluation of enhanced intracellular delivery of doxorubicin in metastatic cancer cells by in-silico and in-vitro approach". New Journal of Chemistry, 2019.p. 12241-12256.
- 324. (Cited 3 times) Kaushik, Aman Chandra, et al. "Exosomal ncRNAs profiling of mycobacterial infection identified miRNA-185-5p as a novel biomarker for tuberculosis." *Briefings in Bioinformatics* (2021).
- 325. (Cited 3 times) Lai-Yu, L., W. Dong-Qing, C. Xiang-Rong, and J. Guang-Fu, "First-principles calculations of structures and electronic properties of solid pentaerythritol under pressure". Chinese Physics Letters, 2008. 25(9): p. 3368-3371.

- 326. (Cited 3 times) He, J., Wei D.Q., J.-F. Wang, and K.-C. Chou, "Predicting protein-ligand binding sites based on an improved geometric algorithm". Protein and peptide letters, 2011. 18(10): p. 997-1001.
- 327. (Cited 3 time) Cui, H.L., X.R. Chen, G.F. Ji, and D.Q. Wei. The Mesoscopic Structure of β-HMX-Based PBXs by Dissipative Particle Dynamics Simulation. Advanced Materials Research Vols 403-408 (2012) pp 4430-4434.
- 328. (Cited 3 times) Wei, Y.K., N.N. Ge, G.F. Ji, X.R. Chen, L.C. Cai, and D.Q. Wei, "Pressure induced metallization of SiH4 (H2) 2 via first-principles calculations". Computational Materials Science, 2014. 88: p. 116-123.
- 329. (Cited 3 times) Peng, S., X. Zhang, Y. Lu, X. Liao, K. Lu, C. Yang, J. Liu, W. Zhu, and Wei D.Q., "High-scalable Collaborated Parallel Framework for Large-scale Molecular Dynamic Simulation on Tianhe-2 Supercomputer". IEEE/ACM transactions on computational biology and bioinformatics, 2018.
- 330. (Cited 3 times) Yang, L., D. Li, B. Guo, and Wei D.Q., "Theoretical Study on the Inclusion Interaction of β-Cyclodextrin with Gabapentin and Its Stability". Journal of Structural Chemistry, 2019. 60(4): p. 564-574.(Cited 3 time) Zhang, H.-Y., Q. Xu, Y. Xiong, S.-L. Peng, K.M. Kumari, and Wei D.Q., "Membrane defect and water leakage caused by passive calcium permeation". Journal of Molecular Liquids, 2018. 270: p. 227-233.
- 331. (Cited 3 times) Qu, S.Y., Q. Xu, W. Wu, F. Li, C.D. Li, R. Huang, Q. Ding, and D.Q. Wei, "An Unexpected Dynamic Binding Mode between Coagulation Factor X and Rivaroxaban Reveals Importance of Flexibility in Drug Binding". Chemical biology & drug design, 2019.p. 1664-1671.
- 332. (Cited 3 times) Pan, Cheng-Tang, et al. "Characterization of Piezoelectric Properties of Ag-NPs Doped PVDF Nanocomposite Fibres Membrane Prepared by Near Field Electrospinning." *Combinatorial Chemistry & High Throughput Screening* 23 (2020): 0-0.
- 333. (Cited 2 times) Humayun, Fahad, et al. "Structure-guided design of multi-epitopes vaccine against variants of concern (VOCs) of SARS-CoV-2 and validation through In silico cloning and immune simulations." *Computers in biology and medicine* 140 (2022): 105122.
- 334. (Cited 2 times) Idrees, Muhammad, et al. "Core-Proteomics-Based Annotation of Antigenic Targets and Reverse-Vaccinology-Assisted Design of Ensemble Immunogen against the Emerging Nosocomial Infection-Causing Bacterium Elizabethkingia Meningoseptica." *International Journal of Environmental Research and Public Health*, vol. 19, no. 1, Dec. 2021, p. 194.
- 335. (Cited 2 times) Khan, Abbas, et al. "Computational Evaluation of Abrogation of HBx-Bcl-xL Complex with High-Affinity Carbon Nanotubes (Fullerene) to Halt the Hepatitis B Virus Replication." *Molecules* 26.21 (2021): 6433.
- 336. (Cited 2 times) Mukhtar, Farwa, et al. "Emerging mutations in spike and other structural proteins of SARS-CoV-2." *Authorea Preprints* (2021).
- 337. (Cited 2 times) Ren, Hai Chao, et al. "Quasi-Static Two-Dimensional Infrared Spectra of the Carboxyhemoglobin Subsystem under Electric Fields: A Theoretical Study." *The Journal of Physical Chemistry B* 124.43 (2020): 9570-9578.
- 338. (Cited 2 times) Wei, D., J. Lu, J. Zhu, and X. Zhou, "THE STUDIES ON SALT EFFECT OF THE ISOTHERMAL VAPOUR-LIQUID EQUILIBRIUM——ORGANIC SOLVENT-WATER-INORGANIC SALT SYSTEM". Acta Physico-Chimica Sinica, 1987. 3(05): p. 520-524.
- 339. (Cited 2 times) Du, Q., Wei D.Q., and A. Li, "Quantum chemical description for molecular lipophilicity and hydrophilicity (I)-Molecular lipophilic and hydrophilic surface". CHEMICAL JOURNAL OF CHINESE UNIVERSITIES-CHINESE, 2004. 25(9): p. 1706-1710.
- 340. (Cited 2 times) Zhang, T., Q. Zhou, Y. Pang, Y. Wang, C. Jin, J. Huo, L.A. Liu, and Wei D.Q., "CYP-nsSNP: A specialized database focused on effect of non-synonymous SNPs on function of CYPs". Interdisciplinary Sciences: Computational Life Sciences, 2012. 4(2): p. 83-89.
- 341. (Cited 2 times) Zhou, C., J. Liang, S. Cheng, T. Shi, K. Houk, Wei D.Q., and Y.-L. Zhao, "Ab initio molecular metadynamics simulation for S-nitrosylation by nitric oxide: S-nitroxide as the key intermediate". Molecular Simulation, 2017. 43(13-16): p. 1134-1141.

- 342. (Cited 2 times) Kaushik, A.C., Mehmood, A., Wei, D.Q. and Dai, X., "Robust Biomarker Screening Using Spares Learning Approach for Liver Cancer Prognosis". Frontiers in Bioengineering and Biotechnology, 2020. 8, p.241.
- 343. (Cited 2 times) Kaushik, A.C., Mehmood, A., Wei, D.Q. and Dai, X., "Robust Biomarker Screening Using Spares Learning Approach for Liver Cancer Prognosis". Frontiers in Bioengineering and Biotechnology, 2020. 8, p.241.
- 344. (Cited 2 times) Kaushik, A.C., Mehmood, A., Dai, X. and Wei, D.Q, "WeiBI (web-based platform): Enriching integrated interaction network with increased coverage and functional proteins from genome-wide experimental OMICS data". Scientific reports, 2020 10(1), p.17.
- 345. (Cited 2 times) Selvaraj, G., S. Kaliamurthi, and Wei D.Q., "Emerging Trends on Nanoparticles and Nano-materials in Biomedical Applications-II". Current pharmaceutical design, 2019. 25(24): p. 2607-2608.
- 346. (Cited 2 times) Kaushik, A.C., Li, M., Mehmood, A., Dai, X. and Wei, D.Q., ACPS: An accurate bioinformatics tool for precision-based anti-cancer peptide generation via omics data. Chemical Biology & Drug Design, 2020.
- 347. (Cited 2 times) Li, Z., Chinnasamy, S., Zhang, Y. and Wei, D.Q., "Molecular dynamics simulation and binding free energy calculations of microcin J25 binding to the FhuA receptor". Journal of Biomolecular Structure and Dynamics, 2020. p.1-10.
- 348. (Cited 2 times) Kaushik, A.C., Mehmood, A., Dai, X. and Wei, D.Q, "WeiBI (web-based platform): Enriching integrated interaction network with increased coverage and functional proteins from genome-wide experimental OMICS data". Scientific reports, 2020 10(1), p.17.
- 349. (Cited 2 times) Kaushik, Aman Chandra, et al. "Pan-cancer Analysis and Drug Formulation for GPR139 and GPR142." *Frontiers in pharmacology* 11 (2020): 2154.
- 350. (Cited 2 time) Khan, Abbas, et al. "BC-TFdb: a database of transcription factor drivers in breast cancer." *Database* 2021 (2021).
- 351. (Cited 2 times) Kaushik, Aman Chandra, et al. "Globally ncRNAs expression profiling of TNBC and screening of functional lncRNA." *Frontiers in bioengineering and biotechnology* 8 (2020).
- 352. (Cited 2 times) Guo, X., Y. Zhu, Y. Li, P. Shi, H. Zhou, J. Yao, Z. Huang, and Wei D.Q., "Genetic insight of the H5N1 hemagglutinin cleavage site". Chinese Science Bulletin, 2007. 52(17): p. 2374-2379.
- 353. (Cited 2 times) Yang, F., S. Hu, B. Li, V.M. Dwyer, H. Hassan, Wei D.Q., and P. Shi, "A study of the dynamic relation between physiological changes and spontaneous expressions". Scientific reports, 2017. 7(1): p. 7081-7092.
- 354. (Cited 2 times) Kaushik, A.C., Mehmood, A., Dai, X. and Wei, D.Q, "WeiBI (web-based platform): Enriching integrated interaction network with increased coverage and functional proteins from genome-wide experimental OMICS data". Scientific reports, 2020 10(1), p.1-7.
- 355. (Cited 2 times)Lin, Tao, et al. "Retainable Superconductivity and Structural Transition in 1T-TaSe2 Under High Pressure." *Inorganic Chemistry* 60.15 (2021): 11385-11393.
- 356. (Cited 2 times) Nangraj, Asma Sindhoo, et al. "Insights into mutations induced conformational changes and rearrangement of Fe2+ ion in pncA gene of Mycobacterium Tuberculosis to decipher the mechanism of resistance to pyrazinamide." *Frontiers in molecular biosciences* 8 (2021).
- 357. (Cited 2 times) Khan, Abbas, et al. "BC-TFdb: a database of transcription factor drivers in breast cancer." *Database* 2021 (2021).
- 358. (Cited 1 time) Tan, P., Z. Fu, L. Petridis, S. Qian, D. You, Wei D.Q., J. Li, and L. Hong, "A Two-Fold Structural Classification Method for Determining the Accurate Ensemble of Protein Structures". Communications in Computational Physics, 2019. 25(4): p. 1010-1023.
- 359. (Cited 1 times) Du, W.S.Q.L.H. and Q.S.W.D. Qing, "Prediction of Protein Structural Classes Based on Correlations of Amino Acid Residues [J]". Acta Physico-chimica Sinica, 2004. 5 p. 498-502.
- 360. (Cited 1 time) Tianjun, W., F. Guangying, and W. Dongqing, "The reversed-phase HPLC of dispersing agent NNO and determination of molecular weight distributions". JOURNAL OF DISPERSION SCIENCE AND TECHNOLOGY, 1988. 9(4): p. 363-669.

- 361. (Cited 1 time) Wang, Y., Q. Zhou, H. Dai, T. Zhang, and Wei D.Q., "Prediction of the functional consequences of single amino acid substitution in human cytochrome P450". Molecular Simulation, 2012. 38(14-15): p. 1297-1307.
- 362. (Cited 1 time) Khan, Ayyaz, et al. "Core amino acid substitutions in HCV-3a isolates from Pakistan and opportunities for multi-epitopic vaccines." *Journal of Biomolecular Structure and Dynamics* (2020): 1-16.
- 363. (Cited 1 time) Tu, W., S. Ding, L. Wu, Z. Deng, H. Zhu, X. Xu, C. Lin, C. Ye, M. Han, and M. Zhao, "SynBioEcoli: a comprehensive metabolism network of engineered E. coli in three dimensional visualization". Quantitative Biology, 2017. 5(1): p. 99-104.
- 364. (Cited 1 time) Lodhi, Madeeha Shahzad, et al. "A Novel Method of Affinity Purification and Characterization of Polygalacturonase of Aspergillus flavus by Galacturonic Acid engineered Magnetic Nanoparticle." *Food Chemistry* (2021): 131317.
- 365. (Cited 1 time) Ali, Mahboob, et al. "Chalcones: As Potent α-amylase Enzyme Inhibitors; Synthesis, In Vitro, and In Silico Studies." *Medicinal Chemistry* 17.8 (2021): 903-912.
- 366. (Cited 1 time)Cui, H.l., G.f. Ji, X.r. Chen, and D.q. Wei, "Mesoscopic Simulation of Aggregate Behaviour of Polymers in β-HMX-based PBXs". Chinese Journal of Chemical Physics, 2013. 26(4): p. 462-466.
- 367. (Cited 1 time) Humayun, Fahad, et al. "Computational Method for Classification of Avian Influenza A Virus Using DNA Sequence Information and Physicochemical Properties." *Frontiers in Genetics* 12 (2021): 10.
- 368. (Cited 1 time) Khan, Abbas, et al. "HantavirusesDB: Vaccinomics and RNA-based therapeutics database for the potentially emerging human respiratory pandemic agents." *Microbial Pathogenesis* 160 (2021): 105161.
- 369. (Cited 1 time) Tang, Xingyu, et al. "Deep6mAPred: A CNN and Bi-LSTM-based deep learning method for predicting DNA N6-methyladenosine sites across plant species." *Methods* (2022).
- 370. (Cited 1 time) Shafiq, Athar, et al. "Investigation of the binding and dynamic features of A. 30 variant revealed higher binding of RBD for hACE2 and escapes the neutralizing antibody: A molecular simulation approach." *Computers in Biology and Medicine* 146 (2022): 105574.
- 371. (Cited 1 time) Khan, Taimoor, et al. "Potential Immunogenic Activity of Computationally Designed mRNA-and Peptide-Based Prophylactic Vaccines against MERS, SARS-CoV, and SARS-CoV-2: A Reverse Vaccinology Approach." *Molecules* 27.7 (2022): 2375.
- 372. (Cited 1 time) Khan, Abbas, et al. "A protein coupling and molecular simulation analysis of the clinical mutants of androgen receptor revealed a higher binding for Leupaxin, to increase the prostate cancer invasion and motility." *Computers in Biology and Medicine* 146 (2022): 105537.
- 373. (Cited 1 time) Ren, Hai-Chao, et al. "Revealing the Relationship between Electric Fields and the Conformation of Oxytocin Using Quasi-Static Amide-I Two-Dimensional Infrared Spectra." *ACS Omega* (2022).
- 374. (Cited 1 time) Shah, Ismail, et al. "Evaluation and identification of essential therapeutic proteins and vaccinomics approach towards multi-epitopes vaccine designing against Legionella pneumophila for immune response instigation." *Computers in Biology and Medicine* (2022): 105291.
- 375. (Cited 0 times) Wei, D., J. Lu, and T. Wei, "The mean association number: Ultraviolet spectroscopic study of PbCl 2– H 2 O– C 2 H 5 OH". Journal of solution chemistry, 1986. 15(9): p. 743-748.
- 376. (Cited 0 times) Huang, Juan, et al. "Controlling the Substrate Specificity of an Enzyme through Structural Flexibility by Varying the Salt-Bridge Density." *Molecules* 26.18 (2021): 5693.
- 377. (Cited 0 times) DU, Q.S.W. and D. Qing, "Application of bridge functions in statistical mechanical integral equation theory 3d-RISM-HNC and improvement in the calculation of solvation free energy". Acta Chimica Sinica, 2004. 2.p.120-126.
- 378. (Cited 0 times) Wang, C.-F., H.-Q. Zheng, H.-C. Wei, R. Zhang, H.-F. Chen, and Wei D.Q., "Structure and vibrational frequencies of Ph3PCl2 with discrete solvent molecules and in gas phase". Journal of Theoretical and Computational Chemistry, 2007. 6(03): p. 511-521.
- 379. (Cited 0 times) Kai, X., C. Xiang-Rong, W. Dong-Qing, and G. Qing-Quan, "Surface Corrugation in Rotational and Diffractive Scattering of O2 from LiF (001)". Communications in Theoretical Physics, 2010. 54(6): p. 743-748.

- 380. (Cited 0 times) Xu, K., Wei D.Q., Z.-Y. Wang, P. Lian, L. Huang, X.-R. Chen, L. Chen, Q. Zhang, and G.-F. Ji, "Extraordinary mechanical properties of monatomic C3N2 chain". Molecular Simulation, 2015. 41(4): p. 256-261.
- 381. (Cited 0 times) Huang, R., Xu, Q. and Wei, D.Q., "A significant equilibrium distinguishing DNA/RNA-like phases in 2T SAW model on Husimi tetrahedron lattice". Results in Physics, 2020. p.103110.
- 382. (Cited 0 times) Feng, Xiaoyi, et al. "A drug information embedding method based on graph convolution neural network." 2020 IEEE International Conference on E-health Networking, Application & Services (HEALTHCOM). IEEE, 2021.
- 383. (Cited 0 times) Ahmad, Namra, et al. "Computational Modeling of Immune Response Triggering Immunogenic Peptide Vaccine Against the Human Papillomaviruses to Induce Immunity Against Cervical Cancer." *Viral Immunology* (2021).
- 384. (Cited 0 times) Khan, M. Tahir, et al. "CYP1A2, 2A13, and 3A4 network and interaction with aflatoxin B1." *World Mycotoxin Journal* 14.2 (2021): 179-189.
- 385. (Cited 0 times) Ahmad, Namra, et al. "Computational Modeling of Immune Response Triggering Immunogenic Peptide Vaccine Against the Human Papillomaviruses to Induce Immunity Against Cervical Cancer." *Viral Immunology* (2021).
- 386. (Cited 0 times) Wang, Xianfang, et al. "Prediction of Protein Solubility Based on Sequence Feature Fusion and DDcCNN." *Interdisciplinary Sciences: Computational Life Sciences* (2021): 1-14.
- 387. (Cited 0 times) Azim, Nazia, et al. "A Static Analysis of Wnt/β-Catenin and Wnt/Ca 2+ Biological Regulatory Networks for ARVC Using Automata Network Model." *IEEE Access* 9 (2021): 107611-107624.
- 388. (Cited 0 times) Mehmood, Aamir, et al. "Discovering potent inhibitors against the Mpro of the SARS-CoV-2. A medicinal chemistry approach." *Computers in Biology and Medicine* (2022): 105235.
- 389. (Cited 0 times) Jiang, Xue, et al. "A Self-attention Graph Convolutional Network for Precision Multi-tumour Early Diagnostics with DNA Methylation Data." *bioRxiv* (2022).
- 390. (Cited 0 times) Yang, Yaning, et al. "SVPath: an accurate pipeline for predicting the pathogenicity of human exon structural variants." *Briefings in Bioinformatics* 23.2 (2022): bbac014.
- 391. (Cited 0 times) Li, Cheng-Dong, et al. "Effect of Cholesterol on C99 Dimerization: Revealed by MD Simulations." *Frontiers in Molecular Biosciences*: 346.
- 392. (Cited 0 times) Ali, Imtiaz, et al. "Crystal structure of Acetyl-CoA carboxylase (AccB) from Streptomyces antibioticus and insights into the substrate-binding through in silico mutagenesis and biophysical investigations." *Computers in Biology and Medicine* (2022): 105439.
- 393. (Cited 0 times) Ren, Hai-Chao, et al. "Intermolecular Vibration Energy Transfer Process in Two CL-20-Based Cocrystals Theoretically Revealed by Two-Dimensional Infrared Spectra." *Molecules* 27.7 (2022): 2153.
- 394. (Cited 2 times) Khan, Taimoor, et al. "Subtractive proteomics assisted therapeutic targets mining and designing ensemble vaccine against Candida auris for immune response induction." *Computers in Biology and Medicine* (2022): 105462.
- 395. (Cited 0 times) Khan, M. T., et al. "Inhibitory effect of thymoquinone from Nigella sativa against SARS-CoV-2 main protease. An in-silico study." *Brazilian Journal of Biology* 84 (2022).
- 396. (Cited 0 times) Ahmad, Hussain, et al. "Structural and dynamic investigation of non-synonymous variations in Renin–AGT complex revealed altered binding via hydrogen-bonding network reprogramming to accelerate the hypertension pathway." *Chemical Biology & Drug Design* (2022).
- 397. (Cited 0 times) Zhang, Hui-Yuan, Qin Xu, and Dong-Qing Wei. "Ion permeation across the membrane: A comprehensive comparison analysis on passive permeations of differently charged ions." *Journal of Molecular Liquids* 359 (2022): 119339.
- 398. (Cited 0 times) Li, Cheng-Dong, et al. "Effect of Cholesterol on C99 Dimerization: Revealed by Molecular Dynamics Simulations." *Frontiers in molecular biosciences* 9 (2022): 872385.

- 399. (Cited 0 times) Khan, Abbas, et al. "Structural and molecular insights into the mechanism of resistance to enzalutamide by the clinical mutants in androgen receptor (AR) in castration-resistant prostate cancer (CRPC) patients." *International Journal of Biological Macromolecules* 218 (2022): 856-865.
- 400. (Cited 0 times) Huang, Guohua, et al. "Enhancer-LSTMAtt: A Bi-LSTM and Attention-Based Deep Learning Method for Enhancer Recognition." *Biomolecules* 12.7 (2022): 995.
- 401. (Cited 0 times) Suleman, Muhammad, et al. "Screening of immune epitope in the proteome of the Dabie bandavirus, SFTS, to design a protein-specific and proteome-wide vaccine for immune response instigation using an immunoinformatics approaches." *Computers in Biology and Medicine* 148 (2022): 105893.
- 402. (Cited 0 times) Khan, Muhammad Tahir, et al. "Diversity and novel mutations in membrane transporters of Mycobacterium tuberculosis." *Briefings in Functional Genomics* (2022).
- 403. (Cited 0 times) Suleman, Muhammad, et al. "Sequence-Structure functional Implications and molecular simulation of high deleterious Non-Synonymous substitutions revealed the mechanism of drug resistance in glioma." *Frontiers in Pharmacology*: 3250.
- 404. (Cited 0 times) Li, Daixi, et al. "Investigating the stabilization of IFN-α2a by replica exchange molecular dynamics simulation." *Journal of Molecular Modeling* 28.8 (2022): 1-12.
- 405. (Cited 0 times) Lin, Shenggeng, et al. "DeepPSE: Prediction of polypharmacy side effects by fusing deep representation of drug pairs and attention mechanism." *Computers in Biology and Medicine* (2022): 105984.
- 406. (Cited 0 times) Pan, Deng, et al. "Virulence and Biofilm Inhibition of 3-Methoxycinnamic Acid against Agrobacterium tumefaciens." *Journal of Applied Microbiology* (2022).
- 407. (Cited 0 times) Cao, Yi, et al. "Identifying the kind behind SMILES—anatomical therapeutic chemical classification using structure-only representations." *Briefings in Bioinformatics* (2022).
- 408. (Cited 0 times) Wang, Jiangang, et al., "Structural communication fingerprinting and dynamic investigation of RBD-hACE2 complex from BA.1 x AY.4 recombinant variant (Deltacron) of SARS-CoV-2 to decipher the structural basis for enhanced transmission". Journal of biomolecular Structure & Dynamics (2022) [Just accepted]
- 409. (Cited 0 times) Khan, Taimoor, et al.; "Towards Specie-Specific Ensemble Vaccine Candidates against Mammarenaviruses Using Optimized Structural Vaccinology Pipeline and Molecular Modelling Approaches" Microbial Pathogenesis (2022) [Just accepted]

Editorials (cited 22 times)

(Cited 20 times) Wang, J.-F. and Wei D.Q., "Role of structural bioinformatics and traditional Chinese medicine databases in pharmacogenomics". Pharmacogenomics, 2009. 10(8): p. 1213-1215.

- 413. (Cited 2 times) Wei, D.Q., "Editorial (Hot Topic: New Drug Design Based on Multi-Targets and System Biology Approach in Light of Real Time DNA Sequencing Technologies)". Current topics in medicinal chemistry, 2012. 12(12): p. 1309-1309.
- 414. (Cited 0 times) Khan, A, Ali, L, Wei D.Q., "Breast Cancer Resistance, Biomarkers and Therapeutics Development in the Era of Artificial Intelligence" Frontiers in Molecular Biosciences, 2022 [Just accepted]

415. Special Issues Edited (cited 4 times)

- 416. (Cited 2 times) Current Topics in Medicinal Chemistry, Volume 12, Number 12 [Hot topic: New Drug Design Based on Multi-targets and System Biology Approach in Light of Real Time DNA Sequencing Technologies (Guest Editor: Dong-Qing Wei).
- 417. (Cited 1 time) Current Drug Metabolism, Volume 13, Number 7, August 2012 [Hot Topic: SNPS of Drug Metabolic Enzymes and Personalized Medicine-Part II (Guest Editor: Dong-Qing Wei)].

- 418. (Cited 1 time) Wei, D.-Q. and Q. Chen, "SNPs of Drug Metabolic Enzymes and Personalized Medicine Part II". Current drug metabolism, 2012. 13(7): p. 951-951.
- 419. (Cited 0 times) Current Drug Metabolism, Volume 12, Number 5, June 2011 [Hot Topic: SNPS of Drug Metabolic Enzymes and Personalized Medicine-Part I (Guest Editor: Dong-Qing Wei)].
- 420. (Cited 0 times) Miaomiao Zhao, Yunpeng Cai, Yanjie Wei, FengfengZhou, Jiankui He, GuanyuWang, Dong-Qing Wei and Caiwan Zhang, "Personalized Clinical Data Screening: Special Issue on Health Informatics", Computers in Biology and Medicine. 61, 161-162(2015).
- 421. (Cited 0 times) Muhammad Rizwan Javed, Dong-Qing Wei, Aman Chandra Kaushik, "CADD and Molecular Dynamics Simulations: Potential Impacts to Conventional Medicines", Combinatorial Chemistry & High Throughput Screening, 2020.
- 422. (Cited 0 times) Xiaofeng Dai, Dong-Qing Wei and Jianying Zhang, "Molecular Biomarkers for Cancer Control", Frontier in Genetics, 2020.
- 423. (Cited 0 times)Gurudeeban SelvarajAman Chandra Kaushik, Dong-Qing Wei and Yi Pan, "Computational Genomics and Molecular Medicine for Emerging COVID-19", IEEE/ACM transactions on computational biology and bioinformatics, 2021.
- 424. (cited 0 times) Gurudeeban Selvaraj, Satyavani Kaliamurthi and Dong-Qing Wei, "Advances in Drug Design and Development for Human Therapeutics Using Artificial Intelligence II", Biomolecules, 2022.
- 425. (Cited 0 times) Liaqat Ali, Abbas Khan and Dong-Qing Wei, "Breast Cancer Resistance, Biomarkers and Therapeutics Development in the Era of Artificial Intelligence", Frontiers in Molecular Biosciences, 2022.
- 426. (Cited 0 times) Muhammad Tahir Ul Qamar, Sajjad Ahmad, Abbas Khan and Dong-Qing Wei, "Immunotherapeutics Development against Hantaviruses", Frontiers in Immunology, 2022.
- 427. (Cited 0 times) Mohadeseh Zarei Ghobadi, Majid Teymoori-Rad, Gurudeeban Selvaraj, Dong-Qing Wei, "Computational Systems Immunovirology", Frontier in Immunology, 2022.
- 428. (Cited 0 times) Abbas Khna, Dong-Qing Wei and Syed Shujait Ali, "Vaccine Development for SARS-CoV-2 and Zoonotic Diseases", Vaccines, 2022.

Invited reviews (cited 852 times)

- 429. (Cited 152 times) Chou, K.-C., Wei D.Q., Q.-S. Du, S. Sirois, and W.-Z. Zhong, "Progress in computational approach to drug development against SARS". Current Medicinal Chemistry, 2006. 13(27): p. 3263-3270.
- 430. (Cited 129 times) Wang, J.-F., Wei D.Q., and K.-C. Chou, "Drug candidates from traditional Chinese medicines". Current Topics in Medicinal Chemistry, 2008. 8(18): p. 1656-1665.
- 431. (Cited 86 times) Wang, J.-F., C.-C. Zhang, K.-C. Chou, and Wei D.Q., "Structure of cytochrome p450s and personalized drug". Current medicinal chemistry, 2009. 16(2): p. 232-244.
- 432. (Cited 79 times) Wang, J.-F., Wei D.Q., and K.-C. Chou, "Pharmacogenomics and personalized use of drugs". Current topics in medicinal chemistry, 2008. 8(18): p. 1573-1579.
- 433. (Cited 79 times) Kem, W., F. Soti, K. Wildeboer, S. LeFrancois, K. MacDougall, Wei D.Q., K.-C. Chou, and H.R. Arias, "The nemertine toxin anabaseine and its derivative DMXBA (GTS-21): Chemical and pharmacological properties". Marine Drugs, 2006. 4(3): p. 255-273.

- 434. (Cited 67 times) Chen, Q., T. Zhang, J.-F. Wang, and Wei D.Q., "Advances in human cytochrome p450 and personalized medicine". Current drug metabolism, 2011. 12(5): p. 436-444.
- 435. (Cited 41 times) Gu, R.-X., L.A. Liu, and Wei D.Q., "Structural and energetic analysis of drug inhibition of the influenza A M2 proton channel". Trends in pharmacological sciences, 2013. 34(10): p. 571-580.
- 436. (Cited 39 times) Muhammad, J., A. Khan, A. Ali, L. Fang, W. Yanjing, Q. Xu, and Wei D.Q., "Network pharmacology: exploring the resources and methodologies". Current topics in medicinal chemistry, 2018. 18(12): p. 949-964.
- 437. (Cited 26 times) Zhao, M. and Wei D.Q., "Rare diseases: Drug discovery and informatics resource". Interdisciplinary Sciences: Computational Life Sciences, 2018. 10(1): p. 195-204.
- 438. (Cited 23 times) Zhang, T., Q. Chen, L. Li, L. Angela Liu, and Wei D.Q., "In silico prediction of cytochrome P450-mediated drug metabolism". Combinatorial chemistry & high throughput screening, 2011. 14(5): p. 388-395.
- 439. (Cited 18 times) Selvaraj, G., S. Kaliamurthi, S. Lin, K. Gu, and Wei D.Q., "Prognostic impact of tissue inhibitor of metalloproteinase-1 in non-small cell lung cancer: Systematic review and meta-analysis". Current medicinal chemistry, 2019.
- 440. (Cited 17 times) Xu, J., Q. Gu, H. Liu, J. Zhou, X. Bu, Z. Huang, G. Lu, D. Li, Wei D.Q., and L. Wang, "Chemomics and drug innovation". Science China Chemistry, 2013. 56(1): p. 71-85.
- 441. (Cited 15 times) Wang, Y.-J., F.I. Khan, Q. Xu, and Wei D.Q., "Recent studies of mitochondrial slc25: Integration of experimental and computational approaches". Current Protein and Peptide Science, 2018. 19(5): p. 507-522.
- 442. (Cited 14 times) Sirois, S., R. Zhang, W. Gao, H. Gao, Y. Li, H. Zheng, and Wei D.Q., "Discovery of potent anti-SARS-CoV Mpro inhibitors". Current Computer-Aided Drug Design, 2007. 3(3): p. 191-200.
- 443. (Cited 14 times) Chen, Q., Z. Wang, and Wei D.Q., "Progress in the applications of flux analysis of metabolic networks". Chinese Science Bulletin, 2010. 55(22): p. 2315-2322.
- 444. (Cited 13 times) Zhang, T., M. Zhao, Y. Pang, W. Zhang, L. Angela Liu, and Wei D.Q., "Recent progress on bioinformatics, functional genomics, and metabolomics research of cytochrome P450 and its impact on drug discovery". Current topics in medicinal chemistry, 2012. 12(12): p. 1346-1355.
- 445. (Cited 13 times) Li, L. and Wei D.Q., Bioinformatics tools for discovery and functional analysis of single nucleotide polymorphisms, in Advance in structural bioinformatics. 2015, Springer. p. 287-310.
- 446. (Cited 9 times) Gu, R.-X., Y.-Q. Zhong, and Wei D.Q., "Structural basis of agonist selectivity for different nAChR subtypes: insights from crystal structures, mutation experiments and molecular simulations". Current pharmaceutical design, 2011. 17(17): p. 1652-1662.
- 447. (Cited 9 times) Khan, M.T., A.C. Kaushik, A.I. Bhatti, Y.-J. Zhang, S. Zhang, A.J. Wei, S.I. Malik, and Wei D.Q., "Marine Natural Products and Drug Resistance in Latent Tuberculosis". Marine drugs, 2019. 17(10): p. 549.
- 448. (Cited 8 times) Guo, X., J.-F. Wang, Y. Zhu, and Wei D.Q., "Recent progress on computer-aided inhibitor design of H5N1 influenza A virus". Current computer-aided drug design, 2010. 6(2): p. 139-146.

449. (Cited 1 time) Zhang, H.-Y., Q. Xu, F. Li, P.-C. Tian, Y.-H. Wang, Y. Xiong, Y.-H. Zhang, and Wei D.Q., "Recent progresses of simulations on passive membrane permeations in China". Molecular Simulation, 2016. 42(10): p. 799-808.

Monographs (cited 35 times)

- 450. (Cited 10 times) Wei, D., Q. Xu, T. Zhao, and H. Dai, Advance in structural bioinformatics. 2015: Springer, ISBN: 978-94-017-9245-5.
- 451. (Cited 9 times) Wei, D.-Q., Y. Ma, W.C. Cho, Q. Xu, and F. Zhou, Translational Bioinformatics and Its Application. 2017: Springer, ISBN: 978-94-024-1045-7.
- 452. (Cited 5 times) Multiscale Dynamics Simulations, : Nano and Nano-bio Systems in Complex Environments by D.R. Salahub and Dong-Qing Wei, RSC, October, 2021 (DOI: 10.1039/9781839164668), ISBN: 978-1-83916-178-0.
- 453. (Cited 4 times) Liu, L.A., Wei D.Q., and Y. Li, Interdisciplinary Research and Applications in Bioinformatics, Computational Biology, and Environmental Sciences. 2011: IGI Global, ISBN:1609600649.
- 454. (Cited 4 times) Liu, L.A., Wei D.Q., Y. Li, and H. Lei, Handbook of Research on Computational and Systems Biology: Interdisciplinary Applications. 2011: Igi Global. ISBN: 9781609604912.
- 455. (Cited 3 times) Dong-Qing Wei, Xijun Wang(editors), "Theory and Application of Computational Chemistry", AIP Conference Proceedings Volume 1102, American Institute of Physics Press, 2009, ISBN: 9780735406377.
- 456. (Cited 0 times) Zizheng Gong (Editor), Shaokai Luo, Dong-Qing, Wei, Yongxin Guo and Junyi Guo (Associate Editors), "The progress of Interdisciplinary Research For Mathematics, Mechanics, Physics and High New Technology", Vol. 13, Scientific Publishing, Beijing, 2010, ISBN 9787030282620.
- 457. (Cited 0 times) Rong-Xiu Li, Dong-Qing Wei and Yan Feng, "The Protein Structure Simulation and Design", Chinese Chemical Industrial, 2011, ISBN: 9787122101839.
- 458. (Cited 0 times) Jianguo Du, Dong-Qing Wei, "Experimental and Theoretical Studies of Minerals and Rocks at High Temperature and Pressure", Earth Quake Press, Beijing, 2011, ISBN: 9787502833149.
- 459. (Cited 0 times) Dong-Qing Wei of the editors lead by Xia Li, Bioinformatics, People's Health Press, 2011, ISBN: 9787117129381.
- 460. (Cited 0 times) Dong-Qing Wei of the editors lead by Ming Chen, Bioinformatics, Scientific Press, 2012-01-01 (ISBN: 9787030332059).
- 461. (Cited 0 times) Dong-Qing Wei, Ruo-Xu Gu, Peng Lian, Tao Zhang and Ying Wang, "Molecular Simulation and Computer Aided Drug Design", Shanghai Jiao Tong University Press, 2012, ISBN: 978-7-313-07980-0, Publishing Date: 2012-03026.
- 462. (Cited 0 times) Dong-Qing Wei, Hao Dai, Gui-Hua Jia, Yong-Hong Zhang and Qin Xu, "Computer Aided Drug Design", Shanghai Jiao Tong University Press, 2012, ISBN:978-7-313-16652-4/R, published: 2017-03

Book and proceeding chapters (cited 168 times)

- 463. (Cited 28 times) Fan, H., R. Gu, and Wei D.Q., The α7 nAChR selective agonists as drug candidates for Alzheimer's disease, in Advance in Structural Bioinformatics. 2015, Springer. p. 353-365.
- 464. (Cited 27 times) S. Gao and Dong-Qing Wei, "Data analysis in single-cell transcriptome sequencing, in Computational Systems Biology". 2018, Springer. p. 311-326.
- 465. (Cited 21 times) Zhao, M., Q. Zhou, W. Ma, and Wei D.Q., "Exploring the ligand-protein networks in traditional Chinese medicine: current databases, methods, and applications". Evidence-Based Complementary and Alternative Medicine, 2013. 2013.
- 466. (Cited 17 times) Q. Chen and Wei D.Q., Human cytochrome P450 and personalized medicine, in Advance in Structural Bioinformatics. 2015, Springer. p. 341-351.
- 467. (Cited 13 times) Li, L. and Wei D.Q., Bioinformatics tools for discovery and functional analysis of single nucleotide polymorphisms, in Advance in structural bioinformatics. 2015, Springer. p. 287-310.
- 468. (Cited 12 times) Y. Xiong, X.L. Zhu, H. Dai and Dong-Qing Wei, "Survey of Computational Approaches for Prediction of DNA-Binding Residues on Protein Surfaces", in Methods in molecular biology, March 2018, DOI: 10.1007/978-1-4939-7717-8_13.
- 469. (Cited 10 times) Wei, D., Q. Xu, T. Zhao, and H. Dai, Introduction, in Advance in structural bioinformatics. 2015: Springer.
- 470. (Cited 9 times) Wei, D.-Q., Y. Ma, W.C. Cho, Q. Xu, and F. Zhou, Translational Bioinformatics and Its Application. 2017: Springer.
- 471. (Cited 7 times) Gu, R., L.A. Liu, and Wei D.Q., Drug inhibition and proton conduction mechanisms of the influenza A M2 proton channel, in Advance in Structural Bioinformatics. 2015, Springer. p. 205-226.
- 472. (Cited 7 times) Chou, K.-C., Wei D.Q., Q.-S. Du, S. Sirois, H.-B. Shen, and W.-Z. Zhong, Study of inhibitors against SARS coronavirus by computational approaches, in Viral Proteases and Antiviral Protease Inhibitor Therapy. 2009, Springer. p. 1-23.
- 473. (Cited 4 times) Wang, J.-F., C.-C. Zhang, J.Y. Yan, K.C. Chou, and D.Q. Wei, "Molecular modeling of CYP proteins and its implication for personal drug design". Automation in Proteomics and Genomics, 2009.
- 474. (Cited 4 times) Wei, D., F. Zhang, and T.K. Woo. Ab initio molecular dynamics simulations of molecular collisions of nitromethane. in AIP Conference Proceedings. 2002. American Institute of Physics.
- 475. (Cited 4 times) Dong-Qing Wei, F. Zhang and T. K. Woo, "First-Principle Simulations of Energetic Molecular Liquids", AIP Conference Proceedings, 620, 407-410(2002).
- 476. (Cited 4 times) Zhang, T. and Wei D.Q., Recent progress on structural bioinformatics research of cytochrome P450 and its impact on drug discovery, in Advance in Structural Bioinformatics. 2015, Springer. p. 327-339.
- 477. (Cited 4 times) Wang, Y.-K., Wei D.Q., R.-X. Gu, H.-M. Fan, and J. Ulmschneider, "Applications of rare event dynamics on the free energy calculations for membrane protein systems". Canadian Journal of Chemistry, 2013. 91(9): p. 769-774.
- 478. (Cited 3 times) Kaushik, A.C., S. Bharadwaj, A. Kumar, A. Dhar, and Wei D.Q., New Trends in Artificial Intelligence: Applications of Particle Swarm Optimization in Biomedical Problems, in Intelligent System. 2018, BoD–Books on Demand. p. 193-207.

- 479. (Cited 3 times) Lian, P. and Wei D.Q., An application of QM/MM simulation: the second protonation of cytochrome P450, in Advance in Structural Bioinformatics. 2015, Springer. p. 311-324.
- 480. (Cited 3 times) A. C.Kaushik, S. Bharadwaj, A. Kumar, A. Dhar and Dong-Qing Wei, "New Trends in Artificial Intelligence: Applications of Particle Swarm Optimization in Biomedical Problems", In Intelligent System, August 2018, DOI: 10.5772.
- 481. (Cited 1 time) Bratko, D., L. Blum, and Wei D.Q., A model of ion hydration, in Interactions of Water in Ionic and Nonionic Hydrates. 1987, Springer. p. 27-31.
- 482. (Cited 1 time) Salahub, D.R., A. Martinez, and Wei D.Q., Structure, reactivity and dynamics of atomic and molecular clusters using density functional theory (DFT) and other tools, in Theory of Atomic and Molecular Clusters. 1999, Springer. p. 157-180.
- 483. (Cited 1 time) J.J. Zhu, T.C. Bai and Dong-Qing Wei, "The Electron-Transfer Rate Processes in Biological Systems", in Theory and Application of Computational Chemistry, AIP Conference Proceedings Volume 1102, American Institute of Physics Press, 2009.
- 484. (Cited 0 times) A. Khan and Dong-Qing Wei, "Machine Learning Algorithms for the Analysis of Molecular Dynamics Trajectories", in Multiscale Dynamics Simulations: Nano and Nanobio Systems in Complex Environments by D.R. Salahub and Dong-Qing Wei, RSC, October 2021 (DOI: 10.1039/9781839164668), ISBN: 978-1-83916-178-0.
- 485. (Cited 0 times) Satyavani Kaliamurthi, Gurudeeban Selvaraj, William C. Cho, Dong-Qing Wei, and Gilles H. Peslherbe, "The Implication of miRNA Signature in the Characteristic Features and Diagnosis of Lung Cancer", in Handbook of Cancer and Immunollogy, Springer Nature, March 2023, https://doi.org/10.1007/16833_2023_130.
- 486. (Cited 0 times) Aman Chandra Kaushik, Shakti Sahi and Dong-Qing Wei, "Computational Methods for Structure-Based Drug Design Through System Biology", in Computational Methods for Estimating the Kinetic Parameters of Biological Systems, January 2022, DOI: 10.1007/978-1-0716-1767-0_9.
- 487. (Cited 0 times) X.Q. Guan and Dong-Qing Wei, "Free Energy Analysis Algorithms along Transition Paths and Transmembrane Ion Permeation", in Multiscale Dynamics Simulations: Nano and Nano-bio Systems in Complex Environments by D.R. Salahub and Dong-Qing Wei, RSC October 2021(DOI: 10.1039/9781839164668), ISBN: 978-1-83916-1780-0.
- 488. (Cited 0 times) Berard, D., Wei D.Q., and D. Salahub. Towards a density functional treatment of chemical reactions in complex media. in Pacific Symposium on Biocomputing. Pacific Symposium on Biocomputing. 1997.
- 489. (Cited 0 times) Dong-Qing Wei, "Theory and Simulation of Strongly Interacting Dipolar Fluids: Ferroelectric Liquid Crystals, Ferro-fluids and Electrorheological Fluids", Proceeding of the First International Conference on Frontiers of Physics: Looking to the 21st Century, 438-441, World Scientific, Singapore, 1997.
- 490. (Cited 0 times) Dong-Qing Wei and D.R. Salahub, "DFT Ab Initio Molecular Dynamics and Combined DFT and Molecular Dynamics Simulations", in Combined Quantum Mechanical and Molecular Mechanical Methods, Eds, J. Gao and Mark Thompson, John Wiley, 1998.
- 491. (Cited 0 times) Dong-Qing Wei, "Recent Advances on the Molecular Theories of Electrolyte Solution: Equilibrium Structure, Thermodynamics and Dynamics", note from the plenary lectures, The Problems of Solvation and Complex Formation in Solutions, Ivanovo, Russia, 1998.

- 492. (Cited 0 times) Wang, X.J., C.Y. Dai, Y.L. Zhang, and D.Q. Wei. Application of QM/MM Methods in Theoretical Studies of Enzyme Catalysis. in AIP Conference Proceedings. 2009. American Institute of Physics.
- 493. (Cited 0 times) Jing-Fang Wang and Dong-Qing Wei, "Protein Structure", in Bioinformatics, People's Health Press, 2010 (ISBN: 9787117129381).
- 494. (Cited 0 times) Liang Zhang, Guang-fu Ji, Jian-jin Tan and Dong-Qing Wei, "Progress on The Experimental and Theoretical Studies of MgAl2O4 at High Temperature and Pressure", Experimental and Theoretical Studies of Minerals and Rocks at High Temperature and Pressure, eds, Jianguo Du et al, Earth Quake Press, Beijing, 2011.
- 495. (Cited 0 times) Dong-Qing Wei, Hao Dai and Yong-Hong Zhang, "Bioinformatics and Drug Discovery", Inspiration of Frontier of Life Sciences, edited by Zhong-Dong Qiao and Lin He, The Scientific Press, 2016, ISBN 978-7-03-049733-8.

Papers in journals not cited by SCI

- 496. Jianping Lv, Dong-Qing Wei*, Yonghua Wang and Qin Xu, "Thermal Stability of Lipase 5 Based on MD Simulations", J. Atomic Mol. Phys. 32, 128-134(2016).
- 497. Junhui Gao, Zeren Guan, Zhao Liu, Yingfu Zhong and Dong-Qing Wei, "The Implement of AutoDockVina on the Super-Computer Taihu Light", Harold of Scientific Innovation. 23, 89(2016).
- 498. Lin Huang, Xiao-Li Yuan, Shao-Xin Cui and Dong-Qing Wei*, "The Compression Behaviors of Zirconium From The First Principle Calculations", J. Atomic Mol. Phys. 29(6), 1069 1076(2012).
- 499. Xiao-Li Yuan, Dong-Qing Wei, Yan Cheng, Qing-Ming Zhang, Zi-Zheng Gong, "Thermodynamic properties of Zr2Al under high pressure from first-principles calculations", J. At. Mol. Sci. 3, 160(2012).
- 500. Shuo Liu, Jiang Chang, Dong-Qing Wei, Xiang-Rong Chen, Qing-Ming Zhang, Zi-Zheng Gong, Guang-Fu Ji, and Yong-Xin Guo, "Properties and phase transitions of the solid β-HMX: different force fields", J.At. Mol. Sci. 3, 41-48 (2012).
- 501. Huiqin Zheng and Dong-Qing Wei, "Acetylcholinestemse Inhibitor Screening and Rational Design", Journal of Henan Normal University. 39, 127-130(2011).
- 502. Liang Zhang, Guangfu Ji, Feng Zhao, Dong-Qing Wei, "The First Principle Studies of Pressure Induced Structure Change For SrZrO3 And ZnCr2O4", Journal of Sichuan University (Natural Science Edition). 48, 142 -146(2011).
- 503. Huiqin Zheng, Dong-Qing Wei and Yun Li "Three-Dimensional Quantitative Structure-Activity Relationship of Cyclohexene Neuraminidase Inhibitors", Journal of Xinyang Normal University. 24, 331 -333, (2011).
- 504. H. Du, J. Wang, Y. Zeng, F. Ling, Dong-Qing Wei, Y. Lin and X. Wang, "Progress on the drug metabolism of CYP2C19", Journal of Guangdong College of Pharmacy. 24, 532-535(2008).
- 505. H. Du, J. Wang, Y. Zeng, F. Ling, Dong-Qing Wei, Y. Lin and X. Wang, "Bioinformatic Analysis of Xylose Reductase for Site-Directed Mutagenesis", Journal of South China University of Technology. 36, 122-127(2008).
- 506. H. Gao, Chunfang Wang, Gao Weina, Li Yun, Wei Dong-Qing*, "Quantitatives Structure-Activity Relationship and Modification of HIV Protease Inhibitors", J. Atomic Mol. Phys. 24, 675(2007).

- 507. Wei-Na Gao, Bing-Ni Liu, Xiao-Wen Ren, Wei-Ren Xu and Dong-Qing Wei*, "Theoretical modifications of agaritine based on HIV protease structures", Chinese Herb Medicine. 38, 5(2007).
- 508. Yingjie Wang, Dong-Qing Wei*, Lu Wang, Zizheng Gong, Yongxin Guo, "Theoretical Studies of Dipolar Fluids", J. Atomic Mol. Phys. 24, 451(2007).
- 509. Lu Wang, Jijun Zhao Zhao*, Yingjie Wang, Zizheng Gong, Yongxin Guo, Dong-Qing Wei, "Structure, Electronic Properties, and Oxygen Adsorption of Al13 Cluster by First-Principles Calculations", J. Atom. Mol. Phys. 24, 559(2007).
- 510. Chun-fang Wang, Jing-fang Wang, Lin Li and Dong-Qing Wei*, "Examples of Applying Molecular Modelling on Chemistry and Biology Problems", J. Atomic Mol. Phys. 26, No. 2, 316-320(2007).
- 511. Hong Liu, Jijun Zhao, Zi-Zheng Gong Zheng Wei, Jianguo Du, Guangu Ji and Dong-Qing Wei, "Crystal Structure of TATB Under High Pressure", J. Atomic Mol. Phys. 26, No. 2, 291 297(2007)
- 512. Huiqin Zheng, Yun Li, Zhiqin Jiang, Qishi Du, Dong-Qing Wei, "2D-QSAR Study of Carbocyclic Derivates Neuraminidase Inhibitors", J. Tianjin Normal University. 26, 13 16(2006).
- 513. Hui Gao, Weina Gao, Yun Li, Weiren Xu, Qishi Du, and Dong-Qing Wei, "Modeling Studies of Peptide Inhibitors For The HIV Protease", J. Atomic Mol. Phys.23, 6-10(2006).
- 514. Weina Gao, Yun Li, Rui Zhang, Hui Gao, Weiren Xu, Dengke Liu, Aixiu Li, Qishi Du, Xin Zhang, and Dong-Qing Wei, "HIV Inhibitor Screening Based on The Traditional Chinese Medicines Database", Acta Pharmaceutica Sinica. 41(3), 241 246 (2006) (Figure was placed on the cover of the issue, and selected as the excellent paper of 2006).
- 515. Li Yun, Hang Chang-jiang, Gao Hui, Gao Wei-Na, Liu Deng-Ke, Xu Wei-Ren, Du Qi-shi, Wei Dong-Qing "Synthesis of (R)-N-[3-[3-Fluoro-4-morpholinyl]phenyl]-2-oxo-5- oxazolidinyl] methanol", Journal of Tianjin Normal University. 25, 13-15(2005).
- 516. Mei Dong, Xin Zhang, Aixiu Li and Dong-Qing Wei, "The Structure and Functions of the PEVK Poly-peptide Domain of Muscle Titin Protein", Chemistry of Life. 25, 113(2004).
- 517. Du Qishi, Wang Shuqing, Wei Dong-Qing, Li Aixiu, "Study of Amino Acids Correlation in Proteins and Application in Structural Analysis", Chemical Journal on Internet (CJI), 6, 40, 2004.
- 518. Jijun Zhao, Hong Liu, Zi-Zheng Gong ad Dong-Qing Wei, "Ab Initio Studies of Organic Molecular Crystals", Energetic Materials, 12, Additional Issue, 497(2004).
- 519. Zi-Zheng Gong, Zhang Xu-Dong, Han Gao, He Bi, Hong Liu, Dong-Qing Wei and Guo Yongxin, "New Progress in Metallic Hydrogen Research", Energetic Materials. 12, Additional Issue, 669 (2004).
- 520. Dong-Qing Wei and J.S. Lu, "Nonprimitive Statistical Mechanical Theory of Electrolyte in Mixed Solvents", Fenzi Kexue Xuebao. 4, 63(1986).
- 521. Dong-Qing Wei and J.S. Lu, "The Application of Vapor-Liquid Equilibrium on The Salt Effect in Mixed Solvents", J. Normal University Henan (Natural Science Etition). 2, 35 -38 (1986).
- 522. Dong-Qing Wei and J.S. Lu, "The Thermodynamics Behaviors of The System of Salts in Mixed Solvent-The Critical Phenomena of System Acetone-Water-LiCl and System Acetone-Water-KCl", J. Normal University Henan (Natural Science Edition). 2, 46-59 (1985).
- 523. Dong-Qing Wei, "Markov Process of Formation of Cloud", Henan Meteorology. 1, 1985.

Conference Proceedings

- 516. Gurudeeban Selvaraj, Satyavani Kaliamurthi, Yang-Kai Wei, Keren Gu, Dong-Qing Wei: Abstract: Comprehensive gene expression meta-analysis and integrated bioinformatic studies of shared signatures between AD and SQ in NSCLC. International Conference on "From Computational Biophysics to Systems Biology 2018 (CBSB2018)", Shenzhen, China, on May 20-23, 2018, Shenzhen, China; 05/2018.
- 517. Satyavani Kaliamurthi, Gurudeeban Selvaraj, Yang-Kai Wei, Keren Gu, Dong-Qing Wei: Abstract: Designing of CD8+ and CD4+ T- cell epitope-based vaccine by targeting whole genome sequence of Asian subtype Human papilloma virus 5. International Conference on "From Computational Biophysics to Systems Biology 2018 (CBSB2018)", on May 20-23, 2018, Shenzhen, China; 05/2018
- 518. Shaoliang Peng, Xiaoyu Zhang, Yutong Lu, Xiangke Liao, Kai Lu, Canqun Yang, Jie Liu, Weiliang Zhu, Dong-Qing Wei: mAMBER: A CPU/MIC collaborated parallel framework for AMBER on Tianhe-2 supercomputer. 2016 IEEE International Conference on Bioinformatics and Biomedicine (BIBM); 12/2016, DOI:10.1109/BIBM.2016.7822595
- 519. Fenglei Yang, Sijung Hu, Xiaoyun Ma, Harnani Hassan, Dong-Qing Wei: A new engineering approach to reveal correlation of physiological change and spontaneous expression from video images. BiOS SPIE 2015, San Francisco; 02/2015, DOI:10.1117/12.2077120
- 520. Peng Lian, Hao-Bo Guo, Jeremy C. Smith, Hong Guo, Dong-Qing Wei: The Catalytic Mechanism and Hyperthermophilic Nature of Cellulase TmCel12A Reveal a Possible Pathway to Improve the Efficiency of Cellulosic Biofuel Production. International Conference on Computational and Systems Biology; 01/2012
- 521. Peng Lian, Jue Li, Dongqi Wang, Dong-Qing Wei: ab initio Dynamics Simulations Reveal the Mechanism of Compound I's Formation during Catalytic Circle of Cytochrome P450cam. 3rd International Conference on Computational and Systems Biology; 10/2011
- 522. Peng Lian, Jing-Fang Wang, Dong-Qing Wei: Molecular Dynamics Studies on phospholamban in membranes. International Conference on Computational and Systems Biology; 10/2009
- 523. Peng Lian, Yongxiang Shi, Yuxiang Bu, Dong-Qing Wei: Study on Interaction between DNA and POU Cooperating with HMG by Molecular Dynamics Simulation. Theory and Applications of Computational Chemistry 2008, Shanghai, China; 09/2008
- 524. Jing-Yi Yan, Jing-Fang Wang, Dong-Qing Wei: Interactions of CYP2C9 with Different Substrates and its Implications for Metabolic Mechanism. Bioinformatics and Biomedical Engineering, 2008. ICBBE 2008. The 2nd International Conference on; 06/2008, DOI:10.1109/ICBBE.2008.45
- 525. Jing-Fang Wang, Lin-Li, Dong-Qing Wei: Discovery of Anti-Hiv Drugs Using Computer Aided Drug Designing Tools. Bioinformatics and Biomedical Engineering, 2007. ICBBE 2007. The 1st International Conference on; 08/2007, DOI:10.1109/ICBBE.2007.87

Abstracts

526. Tangzhen Zhao, Yukun Wang, Qin Xu and Dong-Qing Wei, "Molecular Dynamics Simulations of Calcium Binding Sites in the RCK Domain of the Mthk Gating Ring", 59th Annual Meeting of the Biophysical-Society, Baltimore, MD, 2015.

- 527. Dong-Qing Wei, "Structural Bioinformatics and Chinese Traditional Medicine Database For Drug Design and Personalized Medicine", J Antivir Antiretrovir 3 (4), 169 (ISSN: 1948 -5964).
- 528. D.R. Salahub, H. Guo, E. Proynov, S. Sirois, J.F. Truchon, and Dong-Qing Wei, "Biomolecular modeling with density functional theory and other tools: Aspects of enzymic mechanisms", Book of Abstracts, 215th ACS National Meeting, Dallas, March 29-April 2 (1998), PHYS-264. Publisher: American Chemical Society, Washington, D. C.
- 529. Dong-Qing Wei and D.R. Salahub, "DFT ab initio molecular dynamics (MD) and combined DFT and MD simulation", Book of Abstracts, 214th ACS National Meeting, Las Vegas, NV, September 7-11(1997), COMP-170. Publisher: American Chemical Society, Washington, D. C.
- 530. Dong-Qing Wei and Dennis R Salahub, "Density Functional Theory Ab Initio Molecular Dynamics and Combined Density Functional Theory and Molecular Dynamics Simulations", ACS Symposium Series. 712, 159-171(1998). Publisher: American Chemical Society, Washington, D. C.
- 531. DR Salahub, F. Bohr, ME Casida, JG Guan, C. Jamorski and Dong-Qing Wei," Electronic Response Properties From Density Functional Theory", Book of Abstracts. 208, COMP-33 (1994). Publisher: American Chemical Society, Washington, D. C.

Presentations made at conferences (163)

- 1. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging, Cancer Immunology and COVID-19", 8th National Conference of Bioinformatics and Computational Biology, July 22-25, 2022, Guangzhou, China.
- 2. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", China Big Data Technology Conference, Jan 23-24, 2021. A nationwide conference by the Chinese Computer Society at Changsha, Hunan.
- 3. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", Symposium on New Technologies in Bioinformatics and AI, March 19-21, 2021, A nationwide conference by Xiangtan University at Xiangtan, Hunan.
- 4. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", China Biomedical Technology Innovation and Development Summit Forum and the Second Xiangjiang Academician Forum Activity, May 08, 2021, A nationwide conference by Changsha Science and Technology Association, and Changsha High-tech Industrial Development Zone Management Committee at Xiangtan, Hunan.
- 5. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", High-Performance Computing and Artificial Intelligence Technology Symposium, June 17, 2021, A nationwide conference organized by ACM SIGHPC CHINA, School of Information Science and Engineering, Hunan University, National Supercomputing Changsha Center at Changsha, Hunan.
- 6. Dong-Qing Wei, "AIDD and drug candidates by super-computing", "Biorefining and Synthetic Biology" China-Thailand Cooperation and Exchange Forum, September 27, 2021, A bilateral conference organized by Shanghai Jiao Tong University, Thailand National Research Center for Genetic Engineering and Biotechnology (BIOTEC) at Shanghai.
- 7. Dong-Qing Wei, "AIDD and drug candidates by super-computing", The 16th International Congress of Genomics, October 25-31, 2021, A global conference organized by BGI, The China National GeneBank (CNGB), GigaScience at Qingdao.

- 8. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", The 10th National Conference on Bioinformatics and Systems Biology, A global conference organized by the Chinese Society of Bioinformatics at Chengdu.
- 9. Dong-Qing Wei, "Artificial intelligence supercomputing precision drug discovery: anti-aging and new crown", The 14th Annual Conference of the Chinese Society of Biological Engineering, October 16-17, 2021, A nationwide conference organized by the Biological Resources Committee of Chinese Society of Biological Engineering, Shanxi Biological Engineering Society at Shanxi Jinzhong.
- 10. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", 2021 3rd China Active Health Conference, October 22-24, 2021, A nationwide conference organized by China Active Health Innovation Alliance, Beijing Precision Medicine Society at Shanghai.
- 11. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", ISBRA 2021 & ICBDS 2021, November 26-28, 2021, A nationwide conference by Shenzhen Institute of Advanced Technology, Chinese Academy of Sciences at Shenzhen.
- 12. Dong-Qing Wei, "Precision medicine and health research of lyrebellin", Academician expert Jinan trip, November 02, 2021, organized by China Association for Science and Technology, Chinese Society of Biological Engineering at Jinan, Shandong.
- 13. Dong-Qing Wei, "AIDD and drug candidates by super-computing", the fourth conference on Computational and Mathematical Bioinformatics and Biophysics, December 13-17, 2021, A bilateral conference organized by TSIMF Sanya, Hainan.
- 14. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", The 11th Jiangsu Provincial Conference on Bioinformatics, December 04-05, 2021, A nationwide conference by Bioinformatics Professional Committee of Jiangsu Biomedical Engineering Society at Nanking.
- 15. Dong-Qing Wei, "Artificial Intelligence Supercomputing Precision Drug Discovery: Anti-Aging and COVID-19", Wu Wenjun Artificial Intelligence Science and Technology Award 10th Anniversary Award Ceremony and 2020 Chinese Intelligent Industry Annual Conference, April 10-12 2021, A nationwide conference by the Chinese Society of Intelligent Engineering at Suzhou.
- 16. Dong-Qing Wei, "2020 China Big Data Technology Conference", Attended Invited breakout sessions at the national conference organized by the Chinese Computer Society from January 23-24th, 2021 at Changsha, Hunan.
- 17. Dong-Qing Wei, "A nationwide Symposium on New Technologies in Bioinformatics and AI", Attended the Report of the General Assembly organized by Xiangtan University from March 19-21, 2021 at Xiangtan, Hunan.
- 18. Dong-Qing Wei, "The 18th nationwide Yangtze River Delta Science and Technology Forum", Invited reports organized by the Shanghai Association of Science and Technology, Jiangsu Science and Technology Association, Zhejiang Science and Technology Association, Anhui Science and Technology Association, Songjiang District People's Government on May 20th, 2021 at Shanghai.
- 19. Dong-Qing Wei, "A nationwide China Biomedical Technology Innovation and Development Summit Forum and the Second Xiangjiang Academician Forum Activity", Reports of the

- general assembly organized by Changsha Science and Technology Association, Changsha High-tech Industrial Development Zone Management Committee, Changsha Biomedicine (Gene Technology) Industry Chain Promotion Office, Hunan Pharmaceutical Industry Association on May 08, 2021, at Changsha, Hunan.
- 20. Dong-Qing Wei, "A nationwide 2021 High-Performance Computing and Artificial Intelligence Technology Symposium" organized by ACM SIGHPC CHINA, School of Information Science and Engineering, Hunan University, National Supercomputing Changsha Center on June 17, 2021, at Changsha, Hunan.
- 21. Dong-Qing Wei, "A nationwide 8th "From Atoms to Earth" Symposium on High-Pressure Science and Earth Science", organized by Key Laboratory of High Temperature and Pressure of Earth's Internal Matter, Chinese Academy of Sciences, Institute of Geochemistry, Chinese Academy of Sciences, Experimental Mineral, Petrological, and Geochemistry Professional Committee of Chinese Society of Mineralogy, Petrology, and Geochemistry, Fluid Geoscience Committee of Chinese Geophysical Society, High-Pressure Physics Committee of Chinese Physical Society, Chinese Interdisciplinary Science Society from July 2-5, 2021 at Guizhou, China.
- 22. Dong-Qing Wei, "A nationwide invited Symposium on Protein Structure Prediction Based on Next-Generation Supercomputing Platform", organized by National Supercomputing Shenzhen Center from Wu Wenjun Artificial Intelligence Science and Technology Award 10thAnniversAugust 28-29, 2021 at Shenzhen.
- 23. Dong-Qing Wei, "A bilateral Biorefining and Synthetic Biology" China-Thailand Cooperation and Exchange Forum", organized by Shanghai Jiao Tong University, Thailand National Research Center for Genetic Engineering and Biotechnology (BIOTEC) on September 27th, 2021 at Shanghai, China.
- 24. Dong-Qing Wei, "A nationwide Medical Mathematics Satellite Academic Conference", participation organized by the Medical Mathematics Committee of the Chinese Mathematical Society on October 24th, 2021 at Kunming, Yunnan.
- 25. Dong-Qing Wei, "The nationwide 10th National Conference on Bioinformatics and Systems Biology", Invited breakout sessions organized by the Chinese Society of Bioinformatics from October 25-28, 2021 at Chengdu.
- 26. Dong-Qing Wei, "The 14th nationwide Annual Conference of the Chinese Society of Biological Engineering" organized by Biological Resources Committee of Chinese Society of Biological Engineering, Shanxi Biological Engineering Society from October 16-17, 2021 at Shanshi, Jinzhing.
- 27. Dong-Qing Wei, "The nationwide 2021 3rd China Active Health Conference", organized by China Active Health Innovation Alliance, Beijing Precision Medicine Society from October 22-24, 2021 at Shanghai.
- 28. Dong-Qing Wei, "The 9th nationwide China Computer Society Big Data Academic Conference", participation organized by the Chinese Computer Society from November 19-21, 2021 at Guangzhou, Guangdong.
- 29. Dong-Qing Wei, "ISBRA 2021 & ICBDS 2021", A nationwide Report of the General Assembly organized by Shenzhen Institute of Advanced Technology, Chinese Academy of Sciences from November 26-28, 2021 at Shenzhen, China.

- 30. Dong-Qing Wei, "Academician expert Jinan trip", A nationwide trip for a report of the general assembly organized by China Association for Science and Technology, Chinese Society of Biological Engineering on November 2nd, 2021 at Jinan, Shandong.
- 31. Dong-Qing Wei, "The fourth nationwide conference on Computational and Mathematical Bioinformatics and Biophysics", organized by TSIMF from December 13-17, 2021 at Sanya, Hainan.
- 32. Dong-Qing Wei, "The 11th nationwide Jiangsu Provincial Conference on Bioinformatics", organized by Bioinformatics Professional Committee of Jiangsu Biomedical Engineering Society from December 04-05, 2021 at Nanking.
- 33. Dong-Qing Wei, "A nationwide Wu Wenjun Artificial Intelligence Science and Technology Award 10th Anniversary Award Ceremony and 2020 Chinese Intelligent Industry Annual Conference", attended invited breakout sessions organized by the Chinese Society of Intelligent Engineering from April 10-12th 2021 at Suzhou.
- 34. Dong-Qing Wei, "Artificial Intelligence Supercomputing and Accurate Drug Discovery", Supercomputing and AI-assisted COVID-19 Prevention and Cure, Dec. 6-8, 2020, The 3rd Conference of Computational Biology and Bioinformation Processing, Yantai, China, plenary talk.
- 35. Dong-Qing Wei, "Artificial Intelligence Supercomputing and Accurate Drug Discovery", Supercomputing and AI-assisted COVID-19 Prevention and Cure, Aug. 28, 2020, Online Conference, invited talk.
- 36. Dong-Qing Wei, "Artificial Intelligence Supercomputing and Accurate Drug Discovery", The 3rd Worldwide Chinese Conference of Computational Biology, Aug. 3-6, 2020, Online Conference, invited talk.
- 37. Dong-Qing Wei, "Artificial Intelligence Supercomputing and Accurate Drug Discovery", Big Data and AI, On-line Conference, March 13-16, 2020, invited talk.
- 38. Dong-Qing Wei, "Thermal Decomposition of The Solid Phase Nitromethane: Ab Initio Molecular Dynamics Simulations", Workshop of Shock Wave Physics, Aug. 19-23, Shanghai, China, plenary talk.
- 39. Dong-Qing Wei, G.N. Patey, "Rotational Motion in Molecular Liquids", "The Canadian Society for Chemistry Conference", June 1989, Victoria, B.C.
- 40. Dong-Qing Wei, G.M. Torrie and G.N. Patey, "Molecular Solvent Model For an Electrical Double Layer: Effects of Ionic Polarizability", "76th Canadian Society for Chemistry Conference and Exhibition", June 1993, Sherbrooke, Quebec, Canada.
- 41. Dong-Qing Wei and Abbas Khan, "Identification of Novel Drug Targets for Diamond Blackfan Anemia (DBA) Based on RPS19 Gene Mutation, Using Protein-Protein Interaction Network", Invited Talk, 2017 International Conference of System Biology, Shenzhen, China, Aug. 18-21, 2017.
- 42. Jing-Yi Yan, Dong-Qing Wei, Jing-Fang Wang, "Interactions of CYP2C9 with Different Substrates and its Implications for Metabolic Mechanism", The 2nd IEEE International Conference on Bioinformatics and Biomedical Engineering, Wuhan, May 2008 (Co-chairman and invited talk).
- 43. Jing-fang Wang, Lin Li, Dong-Qing Wei and Kuo-Chen Chou, "Discovery of Anti-HIV Drugs Using Computer Aided Drug Design Tool", The 1st IEEE International Conference on

- Bioinformatics and Biomedical Engineering, Wuhan, July 2007(Division chairman and invited talk).
- 44. Dong-Qing Wei, "Ultra-fast Chemical Reactions and Energetic Materials Under Shock Wave", Symposium on High Pressure Science and Earth Science, Xiangyang, China, Aug. 2-4, 2019.
- 45. Dong-Qing Wei, "Personalized Medicine in Era of Bio Data-Discovery of WGX-50 and Antiaging Function", Huaxiang International Forum, Changsha, China, June 16-18, 2019.
- 46. Dong-Qing, Wei, "Rare Event Dynamics Involving Membrane Systems and CADD", 2019 Symposium of Jiangsu and Zhejiang Bioinformatics, Wuxi, China, April 19, 2019.
- 47. Dong-Qing Wei, "Precision Drug Discovery Based on Plant Extractions and Chinese Traditional Medicine Database", 6th National Conference on Computational Biology and Bioinformatics, Chengdu, China, March 29-31, 2019.
- 48. Dong-Qing Wei, "Deep Learning For Precision Medicine of Cancer Treatments", 8th National Conference on System Biology and Bioinformatics, Macau, Oct. 22-24, 2018.
- 49. Qin Xu and Dong-Qing Wei, "Multiple interconverting conformations of C99 dimer characterized by MD simulations", International Workshop on Molecular Simulation 2018, Shanghai, China, August 25 27, 2018.
- 50. Dong-Qing Wei, "Discovery of Wgx-50 and its Anti-Aging Function", Keynote Talk, 17th, Chinese National Conference of Interdisciplinary Sciences, Hulunbeier, China, Aug. 01-05, 2018.
- 51. Ru-Gu and Dong-Qing Wei, "Rare Event Dynamics Involving Membrane Systems and CADD", Plenary Talk, 2nd Chinese Conference on Computational Biophysics and Molecular Simulations, Zhuhai, China, June 07-10, 2018.
- 52. Dong-Qing Wei, "Wgx-50 and its Role Anti-Aging and Radiation", Plenary Talk, 1st High Level Forum of Radiation Biology, Ningbo, China, June 06-08, 2018.
- 53. Dong-Qing Wei, "Discovery of Wgx-50 and its Anti-Aging Function", Keynote Talk, 11th Chinese National Neurology Doctors, Changsha, China, June 01-04, 2018.
- 54. Dong-Qing Wei, "Rare Event Dynamics Involving Membrane Systems and CADD", Invited Talk, Computational Biophysics and Systems Biology" (CBSB2018), Shenzhen, China, May 20-23, 2018.
- 55. Qin Xu and Dong-Qing Wei, "Rare Event Dynamics Involving Membrane Systems and CADD", Invited Talk, 5th National Bioinformatics Conference organized by Chinese Bio-Engineering Society, Tangshan, China, April 21-22, 2018.
- 56. Dong-Qing Wei, "WGX-50 and its Role As a Drug Candidate of AD and Anti-Aging", Plenary Talk, Symposium On Natural Medicine, Hangzhou, China, Aug. 9-12, 2017.
- 57. Dong-Qing Wei, "Personalized Drug, Precision Medicine, WGX-50 and its Role As a Drug Candidate of AD and Anti-Aging", Plenary Talk, Conference on Functional Genomics and System Biology, Harbin, China, August 9-11, 2017.
- 58. Dong-Qing Wei, "SNPs of CYP450 and Personalized Drug, Precision Medicine", Plenary Talk, 3rd Conference on Digital Medicine and Big Data Analysis, Changchun, China, August 6-8, 2017.
- 59. Dong-Qing Wei, "Chemistry and Phase Transition of Deep Carbon", Plenary Talk, 7th Conference From Atom to Earth, High Pressure Earth Science, Changchun, China, August 6-8, 2017.

- 60. Dong-Qing Wei and Guang-Fu Ji, "Phase Transition and The Ultra-fast Chemical Reactions of Energetic Materials Explored by Ab Initio MD Simulations", Plenary Talk at the 10th Conference of Computational Nano-Science and New Materials, Jishou, China July 16-19, 2017.
- 61. Dong-Qing Wei and Guang-Fu Ji, "Ultra-fast Chemical Reactions of Energetic Materials Explored by Ab Initio MD Simulations", Invited Talk at the 13th National Conference of Quantum Chemistry, Dalian, China June 8-11, 2017.
- 62. Dong-Qing Wei, "Initial Chemical Reactions of Explosives Under Shockwave Impact", the deMon Workshop, Calgary, Canada, May 11-17, 2017.
- 63. Dong-Qing Wei, "SNPs of CYP450, Personalized Drug, Precision Medicine, WGX-50 and its Role As a Drug Candidate of AD and Anti-Aging", Plenary Talk at the 4thNational Conference of Bioinformatics, organized by the Bioinformatics Division, Chinese Biotechnology Association, Changsha, China, April 22-24, 2017.
- 64. Dong-Qing Wei, "Application and Assessment of Health Medicine Big Data", Plenary Talk at the First National Conference on the Assessment and Protection of Health & Medicine Big Data Application, organized by the Division of the Assessment and Protection of Health & Medicine Big Data Application, Chinese Association of Health Information, Beijing, China, April 7-8, 2017.
- 65. Dong-Qing Wei, "SNPs of CYP450, Personalized Drug, Precision Medicine, WGX-50 and its Role As a Drug Candidate of AD and Anti-Aging", Plenary Talk at the 2016 Big Dada Technology Conference, Beijing, China, Dec. 8-10, 2016.
- 66. Dong-Qing Wei, "Rare Event Dynamics and Free Energy of Membrane Systems", Invited Talk at the 4th International Conference on Molecular Simulation (ICMS-2016), Shanghai, China, October 23-26,2016.
- 67. Dong-Qing Wei, "Rare Event Dynamics, Ion Permeation and Free Energy of Membrane Systems", Invited Talk at the 7th National Conference of Bioinformatics and System Biology, Chengdu, China, October 7-9,2016.
- 68. Dong-Qing Wei, "Simulations of Protein Dynamics on Super-Computer and CADD", Symposium of High-Performance Computation, Beijing, September 22-24, 2016.
- 69. Dong-Qing Wei, "Rare Event Dynamics of Ion Permeation Across Membranes and Chains", Theory and Application of Computational Chemistry, Seattle, USA, August 13-September 5, 2016.
- 70. Dong-Qing Wei, "An Improved Feature-Based Approach to Predict Effective Drug Combinations", Invited Talk at the 10th International Conference on Systems Biology, Weihai, China, August 19-22, 2016.
- 71. Dong-Qing Wei, "Precision Medicine Informatics", Plenary Talk at the 16th National Conference of Interdisciplinary Sciences, Dandong, China, August 5-8, 2016.
- 72. Dong-Qing Wei, "SNPs of CYP450 and Molecular Metabolism", Plenary Talk at the 7th National Conference of Microbial Genetics, Hohhot, China, August 5-8, 2016.
- 73. Dong-Qing Wei, "SNPs of CYP450and Molecular Metabolism and Toxicity", Plenary Talk at the6th Agricultural Microbiology, 15th Insecticidal Microorganism, 11th entomogenous fungi, Shanghai, China, July 17-19, 2016.
- 74. Dong-Qing Wei, "Free Energy Calculations of Membrane Systems and CADD", 2016 Shanghai Workshop on Frontiers in Molecular Biophysics, Shanghai, China, July 23-26, 2016.

- 75. Dong-Qing, Wei, "QM/MM Studies of Enzyme Catalysis", Plenary Talk at The 21st International Workshop on Quantum Systems in Chemistry, Physics, and Biology (QSCP-XXI) Vancouver, Canada, July 2-9, 2016.
- 76. Dong-Qing Wei, "SNPs of CYP450and Molecular Metabolism and Toxicity", Plenary Talk at the 2016 Cross-Strait Predictive Toxicology Conference:Environment, Food and Health, Taichung, Taiwan, May 5-7, 2016.
- 77. Dong-Qing Wei, "Ab Initio MD and Fast Reactions of Energetic Materials", Plenary Talk at the International Workshop Molecular Simulation, 2016, Xian, China, April 15-17, 016.
- 78. Dong-Qing Wei, "HPC, Personalized Medicine and WGX-50", Plenary Talk at the Conference of Big Data and Precision Medicine, Shanghai, China, March 25-27, 2016.
- 79. Dong-Qing Wei, "Dynamics of Passive Membrane Permeations", Plenary Talk at the High-Performance Computing Symposium, Changshu, Jiangsu, China, Nov. 27-28,2015.
- 80. Dong-Qing Wei, "High Pressure Physics and Chemistry of Some Carbon Containing System in The Geophysical Environment", Plenary Talk at the 5th National Conference on High Pressure Geophysics-From Atoms to Earth, Beijing, China, Nov. 28-30, 2015.
- 81. Dong-Qing Wei, "Drug Screening Technology Based on The Traditional Chinese Medicine Database and Anti-Aging Function of Wgx-50, a Molecule Extracted From Sichuan Pepper" Plenary Talk at the Annual National Conference on Active Components From Herbs, Changsha, China, Nov. 11-18, 2015.
- 82. Yukun Wang, Ruoxu Gu and Dong-Qing Wei, "Free Energy Calculation For Membrane Systems and CADD", Invited Talk at the 9th International Conference on Systems Biology (ISB 2015), Aug. 21-24, 2015, Luoyang, China.
- 83. Yukun Wang, Ruoxu Gu and Dong-Qing Wei, "Free Energy Calculation For Membrane Systems and CADD", Invited Talk at the 13th National Conference on Computational Chemistry, Nov. 19-22, 2015, Guangzhou, China.
- 84. Dong-Qing Wei, "Drug Screening Technology Based on The Traditional Chinese Medicine Database and Anti-Aging Function of Wgx-50, a Molecule Extracted From Sichuan Pepper", Plenary Talk at the 2nd National Conference on New and Green Technology of Pharmacology, Lasha, China, July 25-27, 2015.
- 85. Yukun Wang, Ruoxu Gu and Dong-Qing Wei, "Free Energy Calculation For Membrane Systems and CADD", Plenary Talk at the Annual National Conference on System Biology and Bioinformatics, Zhuhai, China, May 5-7, 2015.
- 86. Dong-Qing Wei, Yukun Wang, Ruoxu Gu, Huameng Fan, Dan Hu and Jacob Ulmschneide, "Rare Event Dynamics: Ion Transportation Through Protein channels and Across Membranes and CADD", Plenary Talk at the Young Scholar Frontier Symposium on Quantitative Biology Development, Beijing, China, May 8-10, 2015.
- 87. Dong-Qing Wei, "Physics and Chemistry of Deep Carbon Circulation", Plenary Talk at the 3rd National Conferences on Geo-biochemistry, Wuhan, China, March 17-18, 2015.
- 88. Dong-Qing Wei, "A Drug Candidate from Traditional Chinese Medicine and Its Potential Role Against AD And in Anti-aging", Symposium of Traditional Medicine, Cha University and Cha Hospitals, Seoul, Korea, Jan. 19-22, 2015.
- 89. Dong-Qing Wei, Yukun Wang, Ruoxu Gu, Huameng Fan, Dan Hu and Jacob Ulmschneide, "Rare Event Dynamics: Ion Transportation Through Protein channels and Across Membranes

- and CADD", Plenary Talk at the Workshop on Frontiers of Molecular Simulations, Beijing, China, Jan. 4-5, 2015.
- 90. Yukun Wang, Ruoxu Gu and Dong-Qing Wei, "Free Energy Calculation For Membrane Systems and CADD", Invited Talk at 6th National Conference on Bioinformatics and System Biology, Nanjing, China, Oct. 6-9, 2014.
- 91. Dong-Qing Wei, Kai Xu, Yanzhi Bai, Shouxin Cui and Guangfu Ji, "Preliminary Studies of Carbon Circulation- Chemical Reactions of Systems Consists of C, H, N, O in The High-Pressure Regime, And Solid Phases of Carbon Dioxide", Plenary talk at The Oriental Forum, Shanghai, China, Sept. 28-30, 2014.
- 92. Yukun Wang, Ruoxu Gu and Dong-Qing Wei, "Rare Event Dynamics: Ion Transportation Through Protein channels and Across Membranes and CADD", Plenary Talk at 14thNational Conference of Interdisciplinary Sciences, Zhengzhou, China, Sept. Aug. 7-10, 2014.
- 93. Nina Ge, Guangfu Ji and Dong-Qing Wei, "Quantum Chemical Simulation of Chemical Reactions in the decomposition of explosives", Plenary Talk at the 10th National Explosion Mechanics, Guiyang, July 26-30, 2014.
- 94. Yukun Wang, Ruoxu Gu and Dong-Qing Wei, "Rare Event Dynamics: Ion Transportation Through Protein channels and Across Membranes and CADD", Plenary Talk at 26th Canadian Symposium on Theoretical and Computational Chemistry (CSTCC), Montreal, Quebec, Canada, July 1-5, 2014.
- 95. Peng Lian, Dong-Qing Wei, Hong Guo, Jeremy Smith "QM/MM Studies of Enzyme Catalysis", Plenary Talk at deMon workshop, Los Cabos, Mexico, April 28-May 1st, 2014.
- 96. Dong-Qing Wei and Yan-Zhi Bai, "Compression and Chemical Reactions For Systems Consists of C, H, N, O, And High Pressure Phases of Carbon Dioxide and Para-xylene (p-xylene)", 2nd Conference on Geochemistry and Geo-biochemistry", Wuhan, March 16-18, 2014.
- 97. Dong-Qing Wei, "Bioinformatics Studies of CYP450 and Personalized Drug Metabolism", Plenary Talk at the 4th International Conferences on Computational and System Biology (ICCSB), Shenzhen, China, Nov. 14-16, 2013.
- 98. Yukun Wang, Ruoxu Gu and Dong-Qing Wei, "Rare Event Dynamics: Ion Transportation Through Protein channels and Across Membranes", Invited talk at 12th National Computational Chemistry Conference, Suzhou, China, Oct. 21-24, 2013.
- 99. Ruoxu Gu, Yukun Wang and Dong-Qing Wei, "Rare Event Dynamics: Ion Transportation Through Protein channels and Across Membranes", Invited talk at 6th Asian and Pacific Conference on Theoretical and Computational Chemistry (APCTCC6), Gyeongju, Korea, July 10-13, 2013.
- 100. Li Li, Hai Dai and Dong-Qing Wei, "Bioinformatics Studies of CYP450 SNPs and Personalized Drug Metabolism", Invited talk at the 94th AAAS Pacific Division Annual Meeting, Las Vegas, USA, June 16-19, 2013.
- 101. Kai Xu, Juqiang Jiang and Dong-Qing Wei, "Introduction to deMon-GUI and QM/MM Studies of Enzyme Catalytic Reactions", Plenary Talk at deMon workshop, Toulouse, France, June 20-24, 2013.
- 102. Dong-Qing Wei, Kai Xu, Yanzhi Bai, Shouxin Cui and Guangfu Ji, "Preliminary Studies of Carbon Circulation- Chemical Reactions of Systems Consists of C, H, N, O in The High Pressure Regime, And Solid Phases of Carbon Dioxide", Plenary talk at Symposium of High

- Pressure Science and Technology in Memory of Prof. Fuqian Jing, Wuhan University of Science and Technology, Wuhan, China, Sept. 24-27, 2012.
- 103. Dong-Qing Wei, Yukun Wang, Ruoxu Gu, Huameng Fan, Dan Hu and Jacob Ulmschneider, "Rare Event Dynamics and Its Applications on the Free Energy Calculations for Membrane Protein Systems", Invited talk at the Theory and Application of Computational Chemistry (TACC 2012), Pavia, Italy, Sept. 1-8, 2012.
- 104. Dong-Qing Wei, "Simulations of Chemical and Biological Systems From Explosives to Membrane Proteins", Plenary talk at the Worldwide Chinese Computational Biology and Molecular Simulation Conference, Dalian, Aug. 9-12, 2012.
- 105. Dong-Qing Wei, Ruo-Xu Gu, Peng Lian and Huai-meng Fan, "Structural Bioinformatics and Chinese Traditional Medicine Database For Drug Design", Invited talk at the 5th National Conference of Bioinformatics, Harbin, Aug. 7-10, 2012.
- 106. Dong-Qing Wei, "Molecular Simulations of Solid Explosives", Plenary talk at the 9thNational Conference on Explosive Mechanics, Xining, China, July26-31 2012.
- 107. Dong-Qing Wei, "Computer Simulations A Must-having Tools for Biological and Material Sciences", Plenary talk at the 15thNational Conference of Chinese Interdisciplinary Sciences", Yinchuan, China, Aug. 1-4, 2012.
- 108. Dong-Qing Wei, Ruo-Xu Gu, Peng Lian and Huai-meng Fan, "Simulations of Chemical and Biological Systems From Explosives to Membrane Proteins", invited talk at the Professor Nick Quirke 60th birthday symposium at Imperial College of London, London, England, July 6, 2012.
- 109. Dong-Qing Wei, "Structural Bioinformatics and Chinese Traditional Medicine Database For Drug Design", invited talk at the International Symposium on Molecular Cognition and Translational Research of Neuropsychiatric Disorders, in Shanghai, China, April 28-30, 2012.
- 110. Dong-Qing Wei, "Structural Bioinformatics and Chinese Traditional Medicine Database For Drug Design", Invited talk at The 28th Congress of Chinese Chemical Society, Chengdu, China, April 13-16, 2011.
- 111. Dong-Qing Wei, "Structural Bioinformatics and Chinese Traditional Medicine Database For Drug Design and Personalized Medicine", invited talk at the International Conference and Exhibition on Virology, Boston, USA, 2011, 5-7 September 2011.
- 112. Dong-Qing Wei, "Reaction Mechanism of Solid Explosive", Invited talk at The National Conference on Theoretical and Quantum Chemistry, Hefei, China, July 2011.
- 113. Dong-Qing Wei, "Reaction Mechanism of Solid Explosive", Plenary talk at Conference From Atom to Earth, Dalian, China, July 2011.
- 114. Dong-Qing Wei, "Reaction Mechanism of Solid Explosive", Plenary talk at The National Conference on Dynamics Response, Tianyuan, China, July 2011.
- 115. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalised Drug Design", Asian Biotechnology Congress, May 11-15, Shanghai, China, Invited talk.
- 116. Dong-QingWei, "Computational Chemistry for Some Real Chemical and Biological Problems", Plenary talk at deMon Workshop, Bremen, Germany, June 2011.
- 117. Dong-Qing Wei, "Computational Chemistry For Some Real Chem/Bio Problems: Reaction Mechanism of Explosives, Personalized Drug Design, Membrane Proteins and Protein-DNA Interactions", Indian Theoretical Chemistry Symposium, Dec. 8-12, Kanpur, India, plenary talk.

- Dong-Qing Wei, "Structural Bioinformatics For Real Biological Problems: Personalized Drug Design, Membrane Proteins And The Protein-DNA Interactions", Sino-German Workshop on Computational systems biology approaches for cancer research and biomarker discovery, January 11-15, 2010 Zhejiang University, Hangzhou, China, invited talk.
- 119. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalized Drug Design", 2008 Analysis of Effective Components of Chinese Traditional Medicines, Haikou, Nov. 11-13, 2009, plenary talk.
- 120. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalized Drug Design",2009 International Workshop on Computational and Integrative Biology, a satellite meeting of the International Conference of Integrative Biology, September 18th to 20th, 2009, Hangzhou, China, invited talk.
- 121. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalized Drug Design", Recent Progress in Computer Simulations in Molecular Sciences, Seoul, Korea, June 14-16, 2009, plenary talk.
- 122. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalized Drug Design", Westlake International Conference on Personalized Medicine, Hangzhou, China, May 29-30, 2009, invited talk.
- 123. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalized Drug Design", 2008 World Gene Congress, Fushan, Oct. 5-7 2008, invited talk.
- 124. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalized Drug Design", 2008 Analysis of Effective Components of Chinese Traditional Medicines, Shenzhen, Oct. 6-8, 2008, invited talk.
- 125. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalized Drug Design", Theory and Application of Computational Chemistry (TACC 2008), Shanghai, Sept. 23-27 2008, chairman and plenary talk.
- 126. Dong-Qing Wei, "Structural Bioinformatics, Traditional Chinese Medicine Database (TCMD) and Personalized Drug Design", 3Rd National Conference on Bioinformatics and System Biology, Wuhan, Oct.7-9 2008, invited talk.
- 127. Dong-Qing Wei, "Structural Bioinformatics and Personalized Drug Design", 5th MMPH Congress, Emeishan, Aug. 2008. Plenary talk and 5th Jiao Shanqing MMPH award.
- 128. Lin Li and Dong-Qing Wei, "Discovery of Drugs Using Computer Aided Drug Design Tool", 10th International Congress on Amino Acids and Proteins, Chalkidiki, Greece, August 2007 (invited talk).
- 129. Jing-fang Wang and Dong-Qing Wei, "Molecular Modeling of CYP450 and Personalized Drug Design", 9th Computational Conference of Chinese Chemical Society, Chengdu, Aug. 2007.
- 130. Dong-Qing Wei, "Computer Aided Drug Design Against HIV Based on Traditional Chinese Medicine Database", World AIDS Day, Tianjin, Nov. 2006 (invited talk).

131.

- 132. Dong-Qing Wei, "Inhibitor Design Against Viruses", International Symposium for Chinese Medicinal Chemists, ISCMC, Nanjing, October 2006(invited talk).
- 133. Dong-Qing Wei, "Bioinformatics and Inhibitor Design Against Viruses", 25th Chinese Chemical Societies, Changchun, China, July 2006(invited talk).

- 134. Liu Hong, Zhao Jijun, WEI Dong-Qing, GONG Zizheng, First-Principles Study of Solid Nitromethane under High Pressure, International Autumn Seminar on Propellants, Explosives and Pyrotechnics, Beijing, Nov. 2005 (invited talk).
- 135. Weina Gao and Dong-Qing Wei, "Bioinformatics and Its Application on Inhibitor Design Against SARS", 3rd International Symposium Computational Methods in Toxicology and Pharmacology Integrating Internet Resources (CMTPI-2005), Shanghai, Oct. 2005.
- 136. Dong-Qing Wei and Qishi Du, "A Combined Study of Ab Initio Quantum Mechanics and Integral Equation Theory in Three-Dimensions", 9th Quantum Chemical Conference, Guilin, Oct. 2005(invited talk and session chair).
- 137. Qishi Du and Dong-Qing Wei et al invited talks of 8 papers on "Studies of the SARS drug design" and other research, Chinese Chemical Societies, Changsha, P.R. China, April 2004 (invited talk).
- 138. Dong-Qing Wei, invited talk, "Bioinformatics And Its Application to Drug Design Against SARS", International Conference on Theory and Application of Computational Chemistry (TACC), Geonjiu, Korea, Feb. 2004(invited talk).
- 139. Dong-Qing Wei, plenary talk, "ab initio Molecular Dynamics Simulations of Molecular Liquids", 12th High Pressure Physics Conference, Huangshan, P.R. China, Sept. 2004 (plenary talk).
- 140. Jijun Zhao, Hong Liu, Zi-Zheng Gong ad Dong-Qing Wei, invited talk, "Ab Initio Studies of Organic Molecular Crystals", Xiamen, P.R. China, Nov. 2004(special invited talk).
- 141. Zi-Zheng Gong, Zhang Xu-Dong, Han Gao, He Bi, Hong Liu, Dong-Qing Wei and Guo Yongxin, invited talk "New Progress in Metallic Hydrogen Research", Xiamen, P.R. China, Nov. 2004.
- 142. Rui Zhang, Yu Ji, Aixiu Li, Xin Zhang, Qishi Du, Kuo-Chen Chou, and Dong-Qing Wei, "Scoring and Docking Studies of the SARS-CoV Mpro binding with a few inhibitors", The Second Annual World Congress, The Human Proteome Organization, Montreal, 2003 (invited talk).
- 143. S.S. Decker and T.K. Woo, Dong-Qing Wei and F. Zhang, "Combined QM/MM and Ab Initio Molecular Dynamics of Nitromethane at High Pressure", in "12th International Detonation Symposium", San Diego, 2002.
- 144. Dong-Qing Wei, F. Zhang and T. K. Woo, "First-Principle Simulations of Energetic Molecular Liquids", in "Shock Wave Physics", Atlanta, 2001.
- 145. Dong-Qing Wei, "Application of Quantum/Classical Molecular Dynamics to Biologically Interesting Systems", in "New Perspectives for Computer-aided Drug Design", Montreal, April 1999.
- 146. Dennis R. Salahub, Hong Guo, Emil Proynov, Suzanne Sirois, Jean-Francois Truchon and Dong-Qing Wei, "Biomolecular Modeling with Density Functional Theory and Other Tools: Aspects of Enzymatic Mechanisms", in "ACS Meeting", Dallas, 1998 (invited talk).
- 147. Dennis R. Salahub, Steeve Chrétien, Anne Milet, Emil Proynov, Suzanne Sirois, Dong-Qing Wei, "Activation Energies and Dynamics from DFT: How Good Are the Functionals?" in "DFT Based Descriptors of Reactivity: Concepts And Applications", Cracow, Dec. 3-5, 1998 (invited talk).

- 148. Dong-Qing Wei "Recent Advances on the Molecular Theories of Electrolyte Solutions: Equilibrium Structures, Thermodynamics and Dynamics", "The Problems of Solvation and Complex Formation in Solutions", Ivanovo, Russia, 1998 (plenary talk).
- 149. Dong-Qing Wei, "Ab Initio MD and Its Application", "The Second International Conference on Frontiers of Physics" and "Joint Meeting of Chinese Physical Societies", on the Occasion of Professor Dayou Wu's 90th birthday, Taipei, 1997.
- 150. Dong-Qing Wei and D.R. Salahub, "DFT Ab Initio Molecular Dynamics and Combined DFT and Molecular Dynamics Simulations", "214th American Chemical Society National Meeting", Las Vegas, September, 1997.
- 151. Dong-Qing Wei and D.R. Salahub, "Hydrated Proton Clusters: Structure, Spectroscopy and Ab Initio Dynamics", "Gordon Research Conference", NH, USA, Aug 1996.
- 152. Dong-Qing Wei "Theory and Simulation of Strongly Interacting Dipolar Fluids: Ferroelectric Liquid Crystals, Ferrofluids and Electrorheological Fluids", "The First International Conference on Frontiers of Physics: Looking to the 21st Century", Shantou, 1995.
- D. Salahub, Dong-Qing Wei, M. Leboeuf, Hong Guo, V. Malkin, O. Olga, T. Woolf and B. Roux, "Chemical Reactivity in Complex Environments Studies With Density Functional Theory and Other Tools", oral presentation in the Scientific Session, Network of Centres of Excellence in Molecular and Interfacial Dynamics, Victoria, B.C., May4-7, 1994.
- 154. Dong-Qing Wei, A. Chandra and G. Patey, "Ion Solvation Dynamics", "The Second Canadian Computational Chemistry Conference", Kingston, May 21-25, 1994.
- Dong-Qing Wei and G.N. Patey,"Ferroelectric Liquid Crystals: a Computer Simulation Study", 34th IUPAC Congress, Beijing, August 1993, "76th Canadian Society for Chemistry Conference and Exhibition", June, 11, Sherbrooke, Quebec, Canada, the Scientific Session, 1992, Network of Centers of Excellence in Molecular and Interfacial Dynamics, Vancouver, B.C., Canada.
- 156. Dong-Qing Wei and G.N. Patey, "Dynamics in a Ferroelectric Nematic Phase", "The Gorden Conference on Water and Aqueous Solution", August 1992, New London, NH, USA.
- 157. Chandra, Dong-Qing Wei and G.N. Patey, "Dielectric Relaxation of Electrolyte Solutions", "The Gorden Conference on Water and Aqueous Solution", August 1992, New London, NH, USA.
- 158. Dong-Qing Wei and G.N. Patey, "The Double Layer Structure in a Model Electrolyte Solution of Polarizable Anion", "The Gorden Conference on Water and Aqueous Solution", August, 1992, New London, NH, USA.
- 159. Dong-Qing Wei, G.N. Patey, "Dielectric Relaxation of Molecular Liquids", "The Gorden Conference on Water and Aqueous Solution", August 1990, New London, NH, USA.
- 160. Dong-Qing Wei, J.J. Zhu, J.S. Lu and L. Blum, "Thermodynamic Behavior of Salt in Mixed Solvents", The Gorden Conference on Water and Aqueous Solution", August 1986, New London, NH, USA.
- 161. Dong-Qing Wei and J.S. Lu, "The Salt Effect on the Gas-liquid Equilibrium in Mixed Solvents", "The Second National Conference on Thermodynamics, Thermochemistry and Thermoanalysis", September 1984, Wuhan, China.
- 162. Dong-Qing Wei, "Time Oscillation Study of a Two Molecule Three Intermediate Reaction", "The Fourth National Conference on Non-equilibrium Statistical Mechanics", October 1984, Guilin, China.

163. Dong-Qing Wei, "Markov Processes in Meteorology", "Symposium on the Application of Statistical Physics on Meteorology", September 1983, Xinjiang, China.

Membership and Professional Affiliations

Fellow, The Royal Society of Chemistry, since 2022;

Editor-in-Chief, "Interdisciplinary Sciences - Computational Life Sciences", snice 2009

Editor-in-Chief, "Current Computer Aided Drug Design", snice 2022;

Co-Editor (Bioinformatics), "Current Chinese Sciences", since 2020;

Editorial Advisor: The Royal Society of Chemistry book series on Theoretical & Computational Chemistry.;

Editorial Board: "Molecular Simulation", "Journal of Molecular Modeling and Graphics", "Scientific Reports", "Biomolecules", "Protein & Peptide Letters", "Journal of Atomic and Molecular Physics (In Chinese: "Chinese Journal of High Pressure Physics", "J. Biomedical Research").

Referees of the following journals

Phys. Rev. Letters, Phys. Reviews, JACS, Bioinformatics, Briefing in Bioinformatics, J. Chem. Inf. Model., JCTC, J. Med. Chem., Biophys. J., J. Biophys., BMC Systems Biology, J. Theor.

Bio., Scientific Report, Current Med. Chem., Med. Chem., J. Chem. Phys. Chem. Phys. Lett., J. Phys. Chem., PCCP, J. Comp. Chem., Chem. Phys., BBRC, Mol. Phys., Mol. Simulation, BMC Microbiol., BBA, Amino Acids, Int. J Bio Macromol, Int. J. Infectious Diseases, Biotech. Prog., IEEE Trans. Comp. Bio. and Bioinformatics, App. Biochem. and Biotech., Cancer Lett., Bioorg. Med. Chem., Protein Peptide Lett., Pharmacogenetics, J. Mol. Graph and Modeling, Acta Chem. Sin., Chinese Physics Letter, J. Energetic Materials, Chem. J. of Chinese Universities, Chinese Science Bull., Comp. Phys., PloS One, PLoS Comp Bio., Nature Communication, App Phys., App Phys. Lett., Mol. Liquids.

Chairman, International Association of Scientists in the Interdisciplinary Areas (IASIA);

Vice Chairman, International Society of Bioinformatics (ISB);

Vice Chairman and Chief of Standing Committee for the Division of Bioinformatics, Chinese Society of Interdisciplinary Sciences;

Vice Chairman, Division of the Assessment and Protection of Health & Medicine Big Data Application, Chinese Association of Health Information;

Vice Chairman and Standing Committee Member of the Division of Functional Genomics Bioinformatics and System Biology, Chinese Cell Biological Society;

Standing Committee Member of the Division of Biological Mathematics, Chinese Mathematics Society;

Executive Standing Committee Member of the Computational System Biology Division, Chinese Society of Operational Research;

Standing Committee Member of the Subdivision for Dynamic Response of Weaponry Materials, Chinese Association of Military Industries;

Ex-Member, IEEE and Engineering in Medicine and Biology Society;

Ex-Member, Biophysical Society

Ex-Member, American Chemical Society

Member, the Virtual Laboratory for Computational Chemistry of CNIC and Supercomputing Center of CNIC, Chinese Academy of Sciences

International Conferences Involved

Chaairman, AI Drug Developemt Forum, 15th "China Pharmaceutical Strategy Conference", March 18-20, 2023, Shijiazhuang, China.

Member of the Organizing Committee, "International Conference on AI and Precision Medicine Informatics & 8th National Conference on Computational Biology and Bioinformatics", May 12-15, 2023, Xuzhou, China.

Member of the Organizing Committee, "10th National Conference on System Biology and Bioinformatics", July 22-25, 2022, Guangzhou, China.

Organizing Committee Member, "The IEEE International Conference on Bioinformatics and Biomedicine (BIBM)", Dec. 9-12, 2021, Houston, TX, USA.

Chairman of the Organizing Committee, "International Conference on AI and Precision Medicine Informatics & 7th National Conference on Computational Biology and Bioinformatics", July 16-20, 2021, Yantai, China.

Organizing Committee Member and Session Chair, "The IEEE International Conference on Bioinformatics and Biomedicine (BIBM)", Dec. 16-20, 2020, Seoul and On-Line, Korea.

Organizing Committee Member, The 13th International Conference on Computational Systems Biology (ISB 2020), September 18-21, 2020, Haikou, China.

Member of Advisory Board, "5th Conference of Theory and Applications of Computational Chemistry (TACC2020)", September 7-12, 2020 at Hokkaido University, Sapporo, Japan.

General Chair, 16th International Symposium on Bioinformatics Research and Applications (ISBRA), December 1-4(2020), Moscow, Russia.

Organizing Committee Member, "The IEEE International Conference on Bioinformatics and Biomedicine (BIBM)", November 18-21, 2019, San Diego, CA, USA.

Organizing Committee Member, "The IEEE International Conference on Bioinformatics and Biomedicine (BIBM)", Dec 3-6, 2018, Madrid, Spain.

Member of the Program Committee, "The 13th International Conference on System Biology (ISB)", August 18-21, 2018, Guiyang, China.

Member of Organizing Committee, "17th Chinese National Conference of Interdisciplinary Sciences", Aug. 01-05, 2018, Hulunbeier, China.

Member of Organizing Committee, "10th Edition of International Conference on Structural Biology 2018", March 15-16, 2018 Barcelona, Spain.

Organizing Committee Member, "The IEEE International Conference on Bioinformatics and Biomedicine (BIBM)", November 17, 2017, Kansas City, Missouri, USA.

Organizing Committee Member, "International Conference on Proteomics (ICP-2017)", November 20-21, 2017, Rome, Italy.

Member of the Program Committee, "GIW / BIOINFO 2017 (International Joint Conference on Genome Informatics Workshop (GIW) and BIOINFO of KSBi 2017)", Oct 31-Nov 3, 2017, Seoul, South Korea.

Committee of Scientific Advisors, "The Baltic Conference Series", 08-11 October 2017, Sweden. Chairman of Publication Committee, "Big Data Conference of Chinese Computation Federation (CCF)", October 14-16, 2017, Shenzhen, China.

Member of the Program Committee, "The 16th International Conference on Bioinformatics (InCoB 2017)", September 20-22, 2017, Shenzhen, China.

Co-Chair and Chairman of Technical Committee, "6th International Conference on Bioinformatics and Biomedical Science (ICBBS 2017)", June 22-24, 2017, Singapore.

Member of the Program Committee, "8th International Conference on Proteomics and Bioinformatics" May 22-24, 2017, Osaka, Japan.

Member of the Program Committee, "7th National Conference of Bioinformatics and System Biology", October 19-23, 2016, Chengdu, China.

Member of the Program Committee, "15th annual InCoB (International Conference on Bioinformatics)", September 21-23,2016, Singapore.

Member of the International Advisory Committee, "Theory and Applications of Computational Chemistry"-TACC 2016, Sept. 2016, Seattle, USA.

Member of Advisory Committee, "Symposium of Interdisciplinary Studies of Mathematics, Computer and Life Sciences", May 21-22, 2016, Beijing, China; Chairman, "deMon Workshop", May 4-8, Zhengzhou, China.

Chairman, "Symposium of Big Data and Precision Medicine", March 25-27, 2016, Shanghai, China. **Member of the International Advisory Committee**, "International Conference on Bioinformatics and Systems Biology (BSB)", March 4-5, Allahabad, India.

Member of the Program Committee, "The 26th annual GIW and 14th annual InCoB conference", September 9-11, 2015, Tokyo, Japan.

Member of the Program Committee, "The 9th International Conference on System Biology (ISB)", August 21-24, 2015, Luoyang, China.

Member of the Program Committee, "International Conference on Biological Engineering and Gene Technology", July 18-19, 2015, Shanghai, China.

Member of the International Advisory Board, "2nd World Congress on Biotechnology", June 2015, Hyderabad city, India.

Member of the Program Committee, The 8th International Conference on Systems Biology and the 4th Translational Bioinformatics Conference, October 25-27, 2014, Qingdao, China.

Chairman, The Oriental Forum, Sept. 28-30, 2014, Shanghai, China.

Member of Organizing Committee, The 6th National Conference of Bioinformatics and System Biology, Oct. 6-9, 2014, Nanjing, China.

Member of the Program Committee, International Conference on Bioinformatics, July 31-Aug.4, 2014, Sydney, Australia.

Member of the Program Committee, 13th International Conference on Bioinformatics of the Asia-Pacific Bioinformatics Network, July 31- Aug. 2, 2014, Sydney, Australia.

Member of the Program Committee, IEEE BIBM 2013 (The IEEE International Conference on Bioinformatics and Biomedicine), Dec. 18-21, 2013, Shanghai, China.

Chairman of the Advisory Committee, International Conference on Computational and System Biology, Nov. 14-16, 2013, Shengzhen, China, Sponsored by the International Association of Scientists in the Interdisciplinary Areas (IASIA), South University of Science and Technology of China (SUSTC) and the 2nd Hospital of Shenzhen;

Organizing Committee Member, Cell Science-2013, November 20 – 22, 2013. Baltimore, USA, organized by OMICS Publishing Group.

Member of the Program Committee, The 12th International Conference on Bioinformatics, Sept. 20-23, 2013, Taicang, Suzhou, China.

Chairman, National Symposium on Deep Carbon Circulation, July 19-24, 2013, Lushan, Jiangxi, China, Sponsored by Chinese National Geophysical Society and International Association of Scientists in the Interdisciplinary Areas (IASIA);

Member of the Program Committee, 1st International Conference on Translational Biomedical Informatics (ICTBI 2012), December 8-10, 2012, Taicang, Suzhou, China.

Chairman, International Conference on Computational and System Biology, Oct. 12-14, 2012, Shanghai, China, Sponsored by The IEEE Engineering in Medicine and Biology Society (EMBS), and International Association of Scientists in the Interdisciplinary Areas (IASIA);

Session Chair, Symposium of High-Pressure Physics, Sept. 25-27, Wuhan, China.

Vice Chairman and Session Chair, Theory and Application of Computational Chemistry, Sept. 2-7, 2012, Pavia Italy;

Session Chair, The Worldwide Chinese Computational Biology and Molecular Simulation Conference, Aug. 9-12, 2012, Dalian.

Session Chair, the 5th National Conference of Bioinformatics, Harbin, Aug. 7-10, 2012.

Chairman, 13th deMon Developer Workshop, May 11-15, 2012, Shanghai, China;

Member of Program Committee, The IEEE International Conference on Bioinformatics and Biomedicine (BIBM), Oct. 4-7, 2012, Philadelphia, USA.

Member of Program Committee, 5th National Bioinformatics Conference, Aug. 7-10, 2012, Harbin, China.

Chairman, International Conference on Computational and System Biology, Oct. 12-14, 2011, Shanghai, China, Sponsored by The IEEE Engineering in Medicine and Biology Society (EMBS), and International Association of Scientists in the Interdisciplinary Areas (IASIA);

Member of the program committee, 2011 Asian Congress of Biotechnology, May 11-15, 2011, Shanghai, China;

Chairman, International Conference on Computational and System Biology, Oct. 22-24, 2010, Hangzhou, China, Sponsored by The IEEE Engineering in Medicine and Biology Society (EMBS), and International Association of Scientists in the Interdisciplinary Areas (IASIA);

Program Chair, International Conference on Computational System Biology, Suzhou, China, Sept. 9-11, 2010, co-sponsored by National Natural Science Foundation of China (NSFC), Academy of Mathematics and Systems Sciences of CAS (AMSS), Shanghai Institutes for Biological Sciences of CAS (SIBS), Shanghai Jiao Tong University, Soochow University, Computational Systems Biology Society of ORSC, Systems Biology Technical Committee of IEEE SMC Society, and also technically sponsored by IEEE SMC Society;

Member of the program committee, The 2010 IEEE International Conference on Bioinformatics and Biomedicine, Dec. 19-22, 2010, Hong Kong;

Chairman, International Conference on Computational and System Biology, Oct. 9-11, 2009, Shanghai, China, Sponsored by The IEEE Engineering in Medicine and Biology Society (EMBS), and International Association of Scientists in the Interdisciplinary Areas (IASIA);

Member of organizing committee, The 7th International Bioinformatics Workshop, June 19th-21st, 2009, Soochow University, Suzhou, China;

Member of organizing committee, The 8th International Bioinformatics Workshop, 2010, Wuhan, China;

Chairman, Theory and Application of Computational Chemistry, Sept. 23-27, 2008, Shanghai, China, one of the largest theoretical chemistry conferences attended by 1000 scientists with 50 plenary talks;

Chairman, 2nd IEEE Conferences on Bioinformatics and Biomedical Engineering, May 16-20, 2008, Shanghai, China;

Member of the program committee, International Conference on Intelligent Data Engineering and Automated Learning (IDEAL), 2003-present

Recent Grants and Awards

- 1. Prediction of Drug-Target Interactions Using Graph Deep Learning and Multi-Label Learning, National Science Foundation of China, 32070662, 2021.1-2023.12, 580,000, PI.
- Precise Clinical Positioning Decision System of Wendan Decoction, University Star Project, Shanghai Jiaotong University, The Medical-Engineering Research Funds, YG2021ZD02 2021.1-2023.12, 500,000 RMB, Co-PI.
- 3. Artificial Intelligence-guided Directed Evolution for Enzyme Engineering, National Science Foundation of China, 32030063, 2021.1-2024.12, 2,920,000, Co-PI.
- 4. Experimental Evaluation and Artificial Intelligence Design of Anti Covid-19 Drug and Vaccine, Strategic Research Funding, Guangdong Province, 2020.5-2021.12, 10 million, PI.
- Construction of Industrial Yeast Using Artificial Intelligence Technology, Major Project, State Key Lab. Of Microbial Metabolism, Shanghai Jiaotong University, 2020.5-2022.12, 0.5 million, Co-PI.
- 6. Shanghai Jiaotong University(Shanghai)-Islamabad-Belgrade Antibiotics Drug Resistance Joint Creative Research Center, Science and Technology Commission of Shanghai Municipality (Grant: 19430750600), 2020.1-2023.12, 2Million, PI.
- 7. Fundamental Theory and Methods of High Throughput Proteomics Computation, Major Grant, National Foundation, 2.81 million, project PI, Project No: 61832019, 2019-2023.
- 8. The Interfacial Features and Recognition Mechanism of Protein-Protein Interactions, The Priority Area of Research, Ministry of Science and Technology, Project No.: 2016YFA0501703, 2016-2021, 4,4 million RMB, project leader.
- 9. Computer Aided Drug Design, First Class Course Building, Shanghai Jiaotong University, 2019.
- 10. Pharmacology Research and New Drug Development of Mangostin Against AD, Joint Medical-Engineering Research Funds, Shanghai Jiao Tong University, Project No. YG2017ZD14, 0.7 million, project co-leader.
- 11. Phase Transition of Energetic Molecular Crystals, The Challenge Research, Chinese Academy of Engineering Physics, 2016-2021, 2.8 million, project leader.
- 12. P450 Enzyme and Efficient Biological Synthesis of phenylpropanoic compounds, Major Project of the National Key Lab. of Microbial Metabolism, 2016-2018, 1 million, project leader.
- 13. The first-class awards of the national classical publication fund for <Translational Medicine Informatics>, National Funds Publication, 2016-2016, 160,000RMB, project leader.
- 14. The first-class international journal drives for <Interdisciplinary Sciences: Computational Life Sciences>, Shanghai High Level Journal Support Project, 2016-2016, 300,000RMB, project leader.
- 15. Microbial Metabolism in The Waste Treatment of Pig Farms, The National Key Technology Support Program, 2014-2016, project leader.
- 16. Graduate Students' Forum of Computational and System Biology, Shanghai Graduate Degree Commission, 2013-2014, project leader.
- 17. Software Development of Personalized Drug Design and Virtual Screening Service, National Innovation Funds, 2012-2015, project leader, project. No.238312C26213202383.
- 18. Fund for International Journal, Shanghai Jiao Tong University, 2013-2014, project leader.

- 19. Software Development For The Property Computation of Energetic Materials, Software Center, Chinese Academy of Engineering Physics, 2013-2014, project leader.
- 20. Joint Biophysics Lab., Shanghai Bureau of Foreign Expert, 2012-2013, project leader, project No. B2012-093.
- 21. Ab Initio Molecular Dynamics Simulation of Typical Explosives, National Science Foundation of China, 2012-2015, Project Leader, Project No. 11174201.
- Multi-targets Drug Screening Technology Based on The Chinese Traditional Medicine Database,
 The National Research Foundation for the Doctoral Program of Higher Education of
 China under Grant No.20120073110057.
- 23. Drug Screening Technology Unifying Western and Chinese Medicine, The National High Technology Research and Development Program of China(863), 2012-2016, PI, project No.2012AA020307.
- 24. Fundamental and Key Problem of Synthetic Biology, The National Basic Research Program of China (973), Ministry of Science and Technology, 2012-2015, PI, project No. 2012CB721000.
- 25. The Major Project, Shanghai Commission of Science and Technology on The Structure of BK Channel And Molecular Mechanism Regulated by The Long Chain Fatty Acids, Prof. No: 11JC1406400,2011-2014, PI.
- 26. The Interdisciplinary Research Project of Shanghai Jiao Tong University on The Multi-scale Research of The Molecular Mechanism of antimicrobial peptide Mimics, 2011-2013, project leader, Project No. AE0800006.
- 27. Research and Development of Unified Software for SNPs of Drug-Metabolic Enzyme and Drug Response, The National High Technology Research and Development Program of China(863), 2008-2012, project leader, project No. 2007AA02Z333.
- 28. The National Basic Research Program of China (973) on Fundamental Durability Problems of Aerospace Devices and Equipment, Ministry of Science and Technology, 2011-2014, PI, project No. 2011CB707500.
- 29. The National Basic Research Program of China (973) on Fundamental and Key Problem of Artificial Vision, Ministry of Science and Technology, 2005-2014, PI, project No. 2005CB724303 and 2011CB707500.
- 30. Molecular Dynamics Simulation of Typical Explosives, National Key Lab. on Explosive Science and Technology, 2009-2011, Project Leader, Project No. KFJJ09-02.
- 31. Molecular Dynamics Simulation of Typical Explosives, National Key Lab. on Explosive Science and Technology, 2012-2014, Project Leader, Project No. KFJJ12-02.
- 32. Drug Discovery Technology Based on Effective Components and Multi-Targets, Ph.D. Research Funds, Ministry of Education, Project Leader, 2012-2014, project No.: 20120073110057.
- 33. Biophysics Joint Lab., Shanghai Bureau of Experts, 2012-2013, Project Leader.
- 34. Funding For International Journal of "985" Third Phase, Shanghai Jiao Tong University, 2013-2014, Project Leader.
- 35. Publishing Funding for "Molecular Simulation and Computer Aided Drug Design", National Publishing Funds, 2012-2013, Project Leader.
- 36. Major Foreign High-Level Expert Funding, National Expert Bureau, 2011-2013, Project Leader.
- 37. Conference Grant for the International Conference on Computational and System Biology (ICCSB, 2009), Shanghai, China, Oct. 11-13, National Science Foundation.

- 38. Design and Screening of Leading Compounds for Anti-Alzheimer Disease, and Relevant Theoretical Studies, National Science Foundation, 2009-2011, project leader, project No 30870476.
- 39. Theoretical and Computational Method and its Application of Chemical Reactions in Complex Environments, National Science Foundation, 2008-2010, project leader, project No. 20773085.
- 40. National Comprehensive Technology Platforms For Innovative Drug R&D, as a PI, 2009-2010, Project No. 2009ZX9301-007.
- 41. Conference Grant for the Theory and Application of Computational Chemistry (TACC 2008), Shanghai, China, Sept. 23-27, National Science Foundation, project leader, project No. 20810302012, 2008.
- 42. Conference Grant for the IEEE Bioinformatics and Biomedical Engineering, Shanghai, China, May 16-18, National Science Foundation, project leader, 2008.
- 43. Funding for International Journals by "985" funds from Jiao Tong University, project leader, 2008.
- 44. Study on the aggregation mechanism of amyloid fibrils, National Science Foundation, 2008-2011, PI, project No. 30770502.
- 45. Theoretical Studies of The HMX Explosives, Chinese Academy of Engineering Physics and also National Key Lab. on Explosives, 2007-2009, project leader.
- 46. The New Approach in Drug Design, Discoveries and Optimization of New Antibiotics Drugs, 2003-2005, Major Grant, The Tianjin Commission of Science and Technology, project leader, Project No. 033801911.
- 47. Discoveries and Optimization of General Anti-virus Drugs Based on Traditional Chinese Medicine Database, Major and Priority Grant from The Tianjin Commission of Science and Technology, project leader, Project No. 043185111-4, 2004-2006.
- 48. Design, Screening, Synthesis and Optimization of Inhibitors Against SARS, The Tianjin Commission of Education, project leader, Project No. 20030001, 2004-2006.
- 49. 3-D Structure Determination Using Bioinformatics, The Tianjin Commission of Science and Technology, project leader, Project No. 023618211, 2002-2005.
- 50. Ab Initio Molecular Dynamics Simulation of Energetic Liquids, Chinese National Science Foundation, project leader, Project No. 10376024, 2004-2006.
- 51. Unified Hydrophobic and Hydrophilic Potential and Its Application in The Drug Design, Chinese National Science Foundation, PI, Project No. 20373048, 2004-2006.

International Exchange Funding from Shanghai Jiao Tong University The Grand Master Awards and Honorary Professorship on behalf of the following leading scientists:

2007, Timothy A. Springer, Harvard University, Fellow of American Academy of Sciences Gert Lubec, Vienna Medical University, Fellow of UK Academy of Sciences

2008, Martin Karplus, Harvard University, Fellow of American Academy of Sciences

Rudolph A. Marcus, Cal. Tech, 1992 Nobel Prize in Chemistry

2010, Dennis Salahub, Univ. of Calgary, Fellow of Canadian Royal Society,

Luc Montagnier, Pasteur Institute, 2008 Nobel Prize in Medicine and Physiology

Overseas Outstanding Scholars

2010, Kuo-Chen Chou, Gordon Life Science Institute, US (ranked No. 1 in terms of hot papers) Heping Zhang, Yale University, Enrico Clemmenti, Italy.

Awards

- 1. The Scopus Highly Cited Scholar, 2020 and 2021.
- 2. The Outstanding Member Award of Shanghai Microbial Society, 2020.
- 3. Nomination for The First-Class Award of Natural Sciences, The Shanghai Municipal Government, 2019.
- 4. The Second-Class Award of the Chinese Computer Federation, 2018.
- 5. The Second-Class Award of Chinese Medical Society, 2017.
- 6. The First-Class Award of Scientific and Technological Advancement by Shanghai Municipal Government, 2017.
- 7. The Outstanding Ph.D. Supervisor by the Shanghai Education Commission, 2016.
- 8. The Science and Technology Award of Shanghai Jiao Tong University, 2013.
- 9. 横山亮次(Yokoyama Ryōji) Awards, 2011.
- 10. The Fifth MMPH Research Award, 2009
- 11. 2nd Prize of Scientific Advancement by CAMP, 2012 due to Screening and Drug Binding Mechanism of Anti-AIDS Drugs.
- 12. 2nd Prize, Excellent Undergraduate Textbook Award, Shanghai Jiao Tong University, "Molecular Simulation and Computer Aided Drug Design", 2015.
- 13. 2011-2012 Excellent Teacher, Shanghai Jiao Tong University
- 14. Nominated for the 10 Most Important Scientific Advancements 2010-2011, Ministry of Education.
- 15. Nominated for the Achievement in Asia Award (Robert T. Poe Prize)
- 16. Research Council UK for China Summer School Competition
- 17. Award from K.C.WONG EDUCATION FOUNDATION, Hong Kong for international conferences
- 18. Shanghai Mengminwei Award for 2009.
- 19. Shanghai Mengminwei Award for 2008.
- 20. 2012 "Authors Contributed the Most" to the Chinese Science Bulletin.
- 21. The best paper award of 2006(the first time) by <Acta Pharmaceutica Sinica> for "HIV Inhibitor Screening Based on The Traditional Chinese Medicines Database", Acta Pharmaceutica Sinica, 41(3), 241-246 (2006)(Figure was placed on the cover of the issue).
- 22. Shanghai Baiyulan Award for 2009.
- 23. Shanghai Baiyulan Award for 2006.
- 24. Shanghai Mengminwei Award for 2006.
- 25. Honorable Mention for a poster titled: "Ab Initio Molecular Dynamics Simulations of Molecular Collisions of Nitromethane", at the 12th Biennial International Conference of the APS Topical Group on Shock Compression of Condensed Matter, Atlanta, June 2001.

Major Awards received by students

- 26. 1. The best thesis award for undergraduate students, Beijing University to Li Lin, July 2007.
- 27. 2. The Morgan Stanley Award administrated by the Shanghai Jiao Tong University to Li Lin, Oct. 2007.
- 28. 3. The national first prize of mathematics modeling to students Liang Jianyi and Shang Yuan, Nov. 2007.
- 29. The National Awards For the Excellent Graduate Students awarded to Gu Ruoxiu, Dec. 2008, The DuPont Award For Graduate Students, awarded to Gu Hui, Dec. 2008.
- 30. The National Awards For the Excellent Graduate Students awarded to Li Li, Dec. 2010, The Jienengke 3rd Prize For Graduate Students, awarded to Chen Qi, Dec. 2009.

- 31. The Qiushi graduate student awards to Lian Peng, The Jienengke 2nd prize for WangYing, The Jenengke 3rd Prize for Li Li, Oct. 2011.
- 32. Mulan 1st Prize awarded to Gu Ruoxu, Guanghua 1st Prize to Chen Qi, Jienengke 2nd to Lian Peng, Jienengke 3rd to Li Li, Oct. 2012.
- 33. Shanghai Jiao Tong University-AMD High Performance Computational Awards to YukunWang. 8, 2013.
- 34. 9. The Outstanding Ph.D. Student to Ruoxu Gu by the Shanghai Educational Commission, 2016.
- 35. The Outstanding Graduate Student awarded by the Shanghai Jiaotong University, Yanjing Wang Ph.D. student, Yanjing Wang, 2018.
- 36. The Outstanding Graduate Student awarded by the Shanghai Jiaotong University, Yanjing Wang Ph.D. student, Chengdong Li, 2019.
- 37. 10.The Globalink Research Award to post-doc, Dr. Gurudeeban Selvaraj and Dr. Satyavani Kaliamurthi by the Mitacs(a national, not-for-profit organization in Canada), 2019.
- 38. The National Graduate Student Award to Ph.D. student, Yanyi Chu, 2020.
- 39. The National Graduate Student Award to Ph.D. student, Yanyi Chu, 2021.
- 40. The Outstanding Graduate Student awarded by the city of Shanghai to Ph.D. student, Yanyi Chu, 2022.
- 41. The Outstanding Undergraduate Student awarded by the city of Shanghai to Zilin Cai
- 42. The super postdoc awarded to Dr. Abbas Khan, 2022.
- 43. The city's New Researcher Award, City of Marseille and the Aix-Marseille University.

Patents

- 44. Dong-Qing Wei, K.C. Chou, Yiru Gan and Qishi Du, "A Polypeptide and Its Derivatives as Inhibitors Against SARS", Patent No: CN 200410018679.3, 2005-01-05.
- 45. Dong-Qing Wei and Yu-Kun Ma, "Chemical synthetic methods for potential preventive and anti-Alzheimer Disease- wgx50, wgx51, wgx52, wgx180", Patent No. ZL20111 0439656.X.
- 46. Dong-Qing Wei and Heng Wang, "New usage of wgx-50, an chemical extracted from Zanthoxylum Bungeanum", 2020-06-17, Patent No. ZL202010551170.4, June 17, 2020.
- 47. Dong-Qing Wei and Heng Wang, "Mechanism of skin aging reversal by active ingredient WGX-50 of Zanthoxylum bungeanum Maxim", Patent No.: 2021104838, Aug. 02, 2021.
- 48. Dong-Qing Wei and Heng Wang, "Extraction method of active ingredient WGX-50 from Zanthoxylum bungeanum and its application in cosmetics", Patent No.: 2021105282, Aug. 11, 2021.
- 49. Dong-Qing Wei and Heng Wang, "Multi-effect Cosmetic Composition Containing Suberosin and Preparation Method thereof", Patent No.: 2021105695, Aug. 17, 2021.
- 50. Software Copy Rights
- 51. Molecule search software based on Maccskey V1.0, 2009SR030823, 2009.08.05
- 52. Database of small molecules from national products or Traditional Medicines, 2009SR042815, 2009.09.27.
- 53. Drug metabolism prediction software based on neural networks V1.0, 2009SR056211, 2009.12.02.
- 54. SNPs prediction software of CYP 450 based on SVM V1.0, 2010SR019961, 2010.05.01.
- 55. DNA binding sites based on protein properties, 2010R11L055771, 2010.08.09.
- 56. Drug metabolism prediction software based on SVM, 2010R11L055833, 2010.08.09.

- 57. Software obtaining the active sites of proteins, 2010R11L055755, 2010.08.09.
- 58. Software searching the active sites of proteins based on the Convex Hull, 2010R11L055766, 2010.08.09.
- 59. The database processing software to predict the SNPs based on sequences, 2010R11L055742, 2010.08.09.
- 60. Molecule finger-print search software based on drug molecule database, V1.0, 2010SR042160, 2010.08.18.
- 61. Databases of SNPs and enzymatic properties of CYP450 enzymes V1.0, 2010SR042161, 2010.08.18.
- 62. Database of drugs-targets and screening platform based on networks V1.0, 2010SR042163, 2010.08.18.
- 63. deMon Gui, V1.0, 2012SR090355.
- 64. Anti-cancer Vaccine Scanner (ACVA: precision based approach for cancer treatments, 2020SR0370047).
- 65. CytoMegaloVirus DataBase, 2020SR0249161.
- 66. weiBI Server, 2020SR0602725.
- 67. weiDOCK Server.

Highly Qualified Personal Trained

Currently supervising 6 post-doctoral fellows, 14 Ph.D. students, 9 master students; graduated 39 Ph.D. students, 61 master students, 25 post-docs. Some alumni have become quite distinguished, including Ruoxu Gu, a Tenured Associated Prof. at Shanghai Jiaotong University, Prof. Tao Zhang at Tianjin Medical University, Dr. Mingzhu Zhao and Yi Xiong at Shanghai Jiao Tong University, Peng Lian at Oakridge National Lab. Li Li at Harvard Medical School, Qi Chen at East Huadong University.

Graduate Students Supervising & Post-Doc.

Name	University	Date	Project	Status
Heqi Sun	Shanghai Jiao Tong University	2022.9	AI Drug	Ph.D. in progress
Yufang Zhang	Shanghai Jiao Tong University	2021.9	BioStat.	Ph.D. in progress
Qiankun Wang	Shanghai Jiao Tong University	2020.9	MD simulation	Ph.D. in progress
Aamir Mehmood	Shanghai Jiao Tong University	2020.9	System Biology	Ph.D. in progress
Munawar Abbas	Henan Institute of Technology	2020.9	MD simulation	Ph.D. in progress
Yitian Fang	Shagnhai Jiaotong University	2019.9	Deep learning models of diseases and biomarkers	Ph.D. in progress
Xueying Mao	Shanghai Jiao Tong University	2019.9	Deep Learning and Protein interactions	Ph.D. in progress
Arif Ali	Shanghai Jiao Tong University	2019.9	Deep Learning and Network Pharm.	Ph.D. in progress
Athar Shafiq	Shanghai Jiao Tong University	2019.9	Drug design	Ph.D. in progress

Fahad	Shanghai Jiao Tong	2018.9	AI and Mol. Machine	Ph.D. in progress
Humayun	University			1 0
Qiuying Dai	Shanghai Jiao Tong University	2018.9	Protein-Protein Interactions	Ph.D. in progress
Yanyi Chu	Shanghai Jiao Tong University	2017.8	Cancer Immunology	Ph.D. in progress
Ashma Sindhoo	Shanghai Jiao Tong University	2015.9	Hadoop and Network Pharmacology	Ph.D. in progress
Juan Huang	Shanghai Jiao Tong University	2015.9	Molecular Motors	Ph.D. in progress
Shenggeng Lin	Shanghai Jiao Tong University	2021.9	Deep Learning	Master in progress
Shifeng Jiang	Shanghai Jiao Tong University	2021.9	Vaccine design and AI	Master in progress
Shah Zeb Khan	Shanghai Jiao Tong University	2021.9	Vaccine design	Master in progress
Bowen Zhao	Shanghai Jiao Tong University	2020.9	Machine learning	Master in progress
Weizhi Chen	Shanghai Jiao Tong University	2020.9	MD simulation	Master in progress
Mingming Jiang	Shanghai Jiao Tong University	2020.9	Deep Learning	Master in progress
Jing Zhao	Shanghai Jiao Tong University	2020.9	Machine learning	Master in progress
Chujun Lv	Shanghai Jiao Tong University	2020.9	Machine learning	Master in progress
Zhili Zhang	Shanghai Jiao Tong University	2018.9-	Personalized Drug	Master in progress
Yatong Liu	Shanghai Jiao Tong University	2018.9	Protein Interactions	Master in progress

Students Graduated since 2005

Name	University	Date Graduated	Project	Status	Current Position
Wei Wang	Shanghai Jiao Tong University	2021.11	Research of Deep Forest Model to the Non-Coding RNA Omics Data	Ph.D.	Takeda Pharma.
Abbas Khan	Shanghai Jiao Tong University	2021.11	Structure-based approaches to understand the molecular mechanism	Ph.D.	Shenzhen institute of Advance Technology,

			of pathogenesis and development of therapeutics database for human respiratory viruses		Chinese Academy of Sciences
Taimoor Khan	Shanghai Jiao Tong University	2021.11	Development of biological databases and its application in designing therapeutics against human pathogenic viruses	Ph.D. in progress	Shenzhen institute of Advance Technology, Chinese Academy of Sciences
Yanjing Wang	Shanghai Jiao Tong University	2019.9-	Molecular Simulation, Drug Design, Molecular Docking	PhD	Post-doc at Shanghai Jiao Tong University
Mahamma d Junaid	Shanghai Jiao Tong University	2020.3	Deep Learning an Cancer Immunity	PhD	Shenzhen institute of advance technology
Chengdong Li	Shanghai Jiao Tong University	2020.3	MD Simulations of Membrane Proteins	Ph.D.	Fujian medical University
Fang Li	Shanghai Jiao Tong University	2019.12-	MD Simulations of Protein Clusters	Ph.D.	Shanghai Jiao Tong University
Yuxin Du	Shanghai Jiao Tong University	2019.9-	Acute Myeloid Leukemia	Ph.D.	Najing Hospital
Kening Li	Shanghai Jiao Tong University	2019.9-	Acute Myeloid Leukemia	Ph.D.	Nanjing medical University
Xiaoqing Guan	Shanghai Jiao Tong University	2019.6	Membrane Permeation	Ph.D.	Shanghai University of Chinese Traditional Medicine
Chunwei Leng	Wuhan Institute of Tech.	2019.7	Energetic Materials	Ph.D.	Henan Institute of Technology
Xiuqing Zhang	Sichuan University	2018.7	High-energetic co-crystal explosive CL-20/TNT	Ph.D.	Zhongbei University

Huiyuan Zhang	Shanghai Jiao Tong University	2018.7	Simulation of Membrane and Protein	Ph.D.	Hebei University of Railways
Yanzhi Bai	Shanghai Jiao Tong University	2018.7	High-pressure Solid CO2 Single-crystal Diffraction and Theoretical Studies	Ph.D.	Post. doc Shanghai Jiao Tong University
Huaimeng Fan	Shanghai Jiao Tong University	2017.7	AD and Wgx-50	Ph.D.	Suzhou Investment Co.
Hao Dai	Shanghai Jiao Tong University	2017.6	Statistical Modeling of CYP 450 Metabolism	Ph.D.	Shanghai Academy of Biological Sciences
Yuanyuan Qi	Sichuan University	2016.7	Condensed Matter Physics	Ph.D.	Henan University of Technology
Yukun Wang	Shanghai Jiao Tong University	2015.7-	New Algorithm of Free Energy Calculation Membrane Permeation Mechanism of Antimicrobial Peptides	Ph.D.	Post.doc at Yale University John Hopkins University
Li Zhang	Beijing Institute of Technology	2015.7	Applied Physics	Ph.D.	Post-doc at Beijing Institute of Technology
Shouxin Cui	Shanghai Jiao Tong University	2015.5-	Theoretical Study of TiZr Alloys	Ph.D.	Liaocheng University
Nina Ge	Sichuan University	2011.7-	Energetic Materials	Ph.D	Xinan University of Science and Technology
Kai Xu	Sichuan University	2010.7-	Energetic Materials	Ph.D.	Henan University Hydro-Power
Peng Lian	Shanghai Jiao Tong University	2013.11	Hybrid Quantum Mechanics and Molecular Mechanics; Drug Metabolism and Cytochrome P450 enzymes; The Catalytic Mechanism of Cellulase	Ph.D.	Fudan University
Ruoxu Gu	Shanghai Jiao Tong University	2013.11	Molecular Dynamics Simulation of M2 Channel	Ph.D.	Post Doc. Oakridge National Lab. USA

Chen Qi	Shanghai Jiao Tong University	2013.11	Impact of resistance mutations on inhibitor binding to HIV-1 integrase	Ph.D.	East China Universi ty of Science and T echnology
Li Li	Shanghai Jiao Tong University	2013.6	Study of SNPS in Human Cytochrome P450 and Protein- Chemical Interactions	Ph.D.	Post-doc in Harvard Medical School
Mingzhu Zhao	Shanghai Jiao Tong University	2013.6	Study on Predictions of Drug-Target Interactions And Drug Combinations	Ph.D.	Shanghai Jiao Tong University
Ying Wang	Shanghai Jiao Tong University	2012.6	Study on Structure and Function of Two Kinds of Important Proteins Based on MD Simulation	Ph.D.	Shanghai Academy of Agricultural Sciences
Xiaoli Yuan	Sichuan University	2012.5	Theoretical Study on Elastic, Electronic Structure and Thermodynamic Properties of Zr Alloy	Ph.D.	Hehai University
Hongling Cui	Sichuan University	2011.7	Ab Initio Studies of HMX	Ph.D.	Henan Industrial University
Tao Zhang	Shanghai Jiao Tong University	2011.7	CYP-nsSNP Database and Theoretical Study of Cyp-mediated Drug Metabolism	Ph.D.	Tianjin Medical University
Yi Xiong	Wuhan University	2011.5	The Study of Characterization and Prediction of Binding Sites on Proteins Based on Machine Learning Methods	Ph.D.	Assistant Professor at Shanghai Jiao Tong University, China
Jing Chang	Sichuan University	2010.10	First-principles MD Simulations of Solid NM and β-HMX	Ph.D.	Sichuan Normal University
Yanli Zhang	Southwest Jiao Tong University	2010.7	Mesoscopic Simulation of Aggregate Behavior of Fluoropolymers in the TATB-Based PBX	Ph.D.	Leshan College
Hong Liu	Southwest Jiao Tong University	2009.7	Ab Initio MD Simulations of Nitromethane Liquids	Ph.D.	China Earthquake Research Institute

Laiyu Lu	Sichuan University	2009.7	Ab Initio Studies of HMX	Ph.D.	Sichuan Normal University
Jingfang Wang	Shanghai Institutes for Biological Sciences	2009.7	Molecular Dynamics Simulations of CYP450 SNPs	Ph.D.	Shanghai Jiao Tong University
Xiaoli Guo	Shanghai Jiao Tong University	2008.7	Cleavage Mechanism of the H5N1 Hemagglutinin by Trypsin and Furin	Ph.D.	Shanghai Jiao Tong University
Zhiwen Shi	Shanghai Jiao Tong University	2021.7-	Protein Machine	M.Ms.	A company in Chengdu
Bin Han	Guangzhou University of Chinese Medicine	2020.7	Gene Expression	M.Sc.	A Guangzhou company
Yufang Zhang	Shanghai Jiao Tong University	2019.7	Deep learning	M.Sc.	Shanghai Investment company
Xiangen Wang	Shanghai Jiao Tong University	2019.7	Muti-label	M.Sc.	PhD at City university of Hongkong
Xiaoqi Shan	Shanghai Jiao Tong University	2019.7	Drug Combination	M.Sc.	Qunar.com
Ali Arif	Shanghai Jiao Tong University	2019.7	Design of validated vaccines against Epstein-Barr virus (EBV)	M.Sc.	Shanghai Jiao Tong University
Qiankun Wang	Shanghai Jiao Tong University	2019.7	MD Simulations of Proteins	M. Sc.	Shanghai Jiao Tong University
Siying Qu	Shanghai Jiao Tong University	2019.7	Drug Design	M.Sc.	Huawei
Abbas Khan	Shanghai Jiao Tong University	2018.9	Identification of novel drug targets against Diamond-Blackfan anemia	M.Sc.	Shanghai Jiao Tong University
Shuang Hou	Shanghai Jiao Tong University	2015.7-	Inhibition of β-amyloid Channels	M.Sc.	PhD in Tongji university

Zhenhua Li	Shanghai Jiao Tong University	2016.9	MD Simulations of Membrane Proteins	M.Sc.	An accounting firm
Liyue Bai	Shanghai Jiao Tong University	2015.7-	Drug Combinations by an Improved Na we Bayesian Algorithm	M.Sc	An investment Company
Qian Xu	Shanghai Jiao Tong University	2018.9-	Drug Combination	M.Sc.	Lily Shanghai
Haifeng Yang	Shanghai Jiao Tong University	2018.7	Drug Combinations	M.Sc.	Huawei, Hangzhou
Pan Tan	Shanghai Jiao Tong University	2018.7	MD Simulations	M.Sc.	Shanghai Jiao Tong University
Yuxi Zheng	Shanghai Jiao Tong University	2016.9-	Personal Drug	M.Sc.	Self-employed
Yiqing Wei	Shanghai Jiao Tong University	2015.7-	Genetic Islands	M.Sc.	Sina.com
Tangzhen Zhao	Shanghai Jiao Tong University	2014.7-	MD Simulation of Membrane Proteins	M.Sc.	Ph.D. Studies at Shanghai Jiao Tong Univ.
Yifan Sun	Shanghai Jiao Tong University	2014.7	Drug Combination Prediction	M.Sc.	Novartis
Jianping Lv	Shanghai Jiao Tong University	2014.7	Molecular Dynamics Simulation on Proteins	M.Sc.	GM Shanghai
Lin Huang	Shanghai Jiao Tong University	2013.7	Molecular Dynamics Simulation of M2 from Influenza A Virus	M.Sc.	Royal Institute of Technology Roslagstullsbacken 15 SE-10691 Stockholm, Sweden
Chaohui Jin	Shanghai Jiao Tong University	2013.4	Integration of Cyp450 Metabolism Database and Online Prediction of Compound Adme	M.Sc.	Self-employed
Shigao Chen	Shanghai Jiao Tong University	2012.7	Virtual Screening for New Drug Candidates Against Alzheimer's Disease Based on Stitch Database	M.Sc.	Patent Bureau Suzhou

Peisi He	Shanghai Jiao Tong University	2012.7	The Molecular Dynamics Simulation Study of Influenza Virus B Proton Channel	M.Sc.	Siemens Power
Quanyi Li	Shanghai Jiao Tong University	2012.7	High-pressure Solid CO2 Single-crystal Diffraction and Theoretical Studies	M. Sc.	
Juan Liang	Shanghai Jiao Tong University	2012.7	MD Simulations	M.Sc.	Novartis Shanghai
Yanyan Qin	Shanghai Jiao Tong University	2012.7	Quantum Studies of Nitromethane	M.Sc.	
Yufang Wang	Shanghai Jiao Tong University	2012.7	Study of the Functional Consequences of Single Amino-acid Substitution in Human Cytochrome p450	M.Sc.	Stockholm University
Qiang Zhou	Shanghai Jiao Tong University	2012.7	Prediction of Protein- ligand Interaction Based on Chemical Preference & Construction of Human Cytochrome P450 Substrate Database (CYP-Meta)	M.Sc.	Sina, Beijing
Xiaobing Li	Wuhan University	2012.5	MD Simulations	M.Sc.	
Yilei Wen	Institute of Animal Science, Academia Sinica	2012.5	Statistical Modeling	M.Sc.	
Chaoqun Xu	Liaoning University	2012.4	Hypervelocity Impact Simulation of TC4 Alloy Based on Material Point Method	M.Sc.	Dongling Vibration Test Instrument Co., Ltd
Jian Yang	Liaoning University	2012.4	Energetic Materials	M.Sc.	
Jiao Zhang	Liaoning University	2012.4	Ferroelectric Liquid Crystals	M.Sc.	
Xiaolei Cui	Liaoning University	2011.12	Energy Materials	M.Sc.	

			1		
Jing He	Shanghai Jiao Tong University	2011.7	Prediction of the Protein-Ligand Binding Sites Based on Geometric Algorithm and Evolutionary Sequence Conservation	M.Sc.	Shanghai Data
Jue Li	Shanghai Jiao Tong University	2011.7	Investigations of Drug- metabolized Enzyme Cytochrome p450 and their Implications for Personalized Drug— Drug-Metabolized Mechanism of CYP2E1	M.Sc.	Shanghai Auto Desk
Huimin Lv	Shanghai Jiao Tong University	2011.7	Free Energy Calculations and Binding Analysis of Two Potential Anti- Influenza Drugs with Polymerase Basic Protein-2 (pb2)	M.Sc.	Pfizer Shanghai
Zhaobin Xu	Shandong University	2011.7	Zinc Finger Stimulation and Optimization	M.Sc.	Villanova University
Liwei Yan	Liaoning University	2011.7	Ferroelectric Liquids	M.Sc.	
Yu Yao	Shanghai Jiao Tong University	2011.7	Mutation Probability of Cytochrome p450 Based on Ga-svm	M.Sc.	Pfizer Shanghai
Lin Gao	Liaoning University	2010.7	Ferroelectric Liquids	M.Sc.	
Shuo Liu	Liaoning University	2010.7	Studies of HMX Using Different Force Fields	M.Sc.	
Zhiyuan Xie	Shanghai Jiao Tong University	2010.7	The Computational Model to Predict Accurately Inhibitory Activity for Inhibitors towards CYP3A4.	M.Sc.	BIG
Congying Dai	Henan Normal University	2010.6	Theoretical Studies of the Rate of Excited State Proton Transfer	M.Sc.	Nantong Cambridge International Exam Center
Hui Gu	Shanghai Jiao Tong	2009.7	Research on HIV-1 Protease Drug	M.Sc.	Ph.D in Rutgers University, USA

	University		Resistance and Virtual screening for Possible Drug Candidates for Alzheimer's Disease.		
Xinchi Hou	Liaoning University	2009.7	CYP 450 Database	M.Sc.	University of British Columbia
Jihe Hu	Liaoning University	2009.7	Ferroelectric Liquids	M.Sc.	Philips
Dan Lian	Liaoning University	2009.7	LDA Studies of HMX	M.Sc.	
Xijun Wang	Shanghai Jiao Tong University	2009.7	Detonation Mechanism of Energetic Materials Using First Principle Modeling	Researche r/M.Sc.	Concordia University, Canada
Bei Tang	Shanghai Jiao Tong University	2008.7	MD Simulations	M.Sc.	Siemens Hong Zhou
Lu Wang	Liaoning university	2007.8	Ab Initio Studies of Al Clusters	M.Sc.	Hong Poly-tech University
Yingjie Wang	Liaoning university	2007.8	Effect of Dipole Elongation on the Ferroelectric Phases of Polar Liquids	M.Sc.	
Yun Li	Tianjin Normal University	2007.7	3-D QSAR of Anti- Bacterial Molecules	M.Sc.	Tianjin Police School
Huachun Wei	Tianjin Normal University	2007.7	Molecular Insights of SAH Enzyme Catalysis and Implication for Inhibitor Design	M.Sc.	Tianjin Industrial University
Rui Zhang	Tianjin Normal University	2007.7	Drug Design against SARS	M.Sc.	University of Calgary
Huiqin Zheng	Tianjin Normal University	2007.7	Screening for New Agonists against Alzheimer's Disease	M.Sc.	Henan Educational College
Chunfang Wang	Tianjin Normal University	2007.7	Structure and Vibrational Frequencies of Ph3PCl2 With Discrete Solvent Molecules and in Gas Phase	M.Sc.	Tianjin High School

Hui Gao	Tianjin Normal	2006.7	QSAR of HIV Drug Candidates from the	M.Sc.	Zhejiang University
Weina Gao	University Tianjin Normal University	2006.7	Traditional Medicine Agaritine and Its Derivatives Are Potential Inhibitors against HIV Proteases	M.Sc.	
Shuqing Wang	Tianjin Normal University	2006.7	Cleavable Peptides of SARSCoV Mpro and Chemical Modification of Octapeptides	M.Sc.	Tianjin Medical University
Wenkang Di	Shanghai Jiao Tong University	2021.7	Deep learning and microbiology	B.S.	Shanghai Jiao Tong University
Tianhang Chen	Shanghai Jiao Tong University	2021.7	Type IV secreted effectors	B.S.	Shanghai Jiaotong University
Lifeng Zhang	Ottawa University	2020.8	Summer Student	B.S.	Oaatwa Univ.
Tongwei Dai	Shanghai Jiaotong University	2020.7	HLA epitope	B.S.	Shanghai Jiaotong University
Yi Fang	Shanghai Jiaotong University	2020.7	Antimicrobial peptides	B.S.	Shanghai Jiaotong University
Kejia Liu	University of California, Santa Barbara	2020.8	Stat. and diseases	Summer students	University of California, Santa Barbara
Mengyang Li	Shanghai Jiaotong Univ.	2020.8	Deep Learning	B.S.	Shanghai Jiaotong Univ.
Yi Lai	Chinese Univ. of Agriculture	2019.8	Stat. Models	Summer student	University of Wisconsin
Zhennan Peng	Lanzhou University	2019.8	Deep Learning	Summer student	Zhongshan University
Vicent Clark	Ottawa University	2018.9	MD simulations of antimicrobiald peptides	Summer students	Ottawa Univ.
Wenhan Chang	Shanghai Jiao Tong University	2014.7-	MD Simulations	Undergra duate	Ph.D. in US
Zhuofei Meng	Shanghai Jiao Tong University	2014.7-	MD Simulations	Undergra duate	Ph.D. in US

Junqiang Jiang	Zhengzhou Information Engineering University	2012.9	DeMon GUI	B.Sc.	A company in Wuhan
Songyao Ma	Shanghai Jiao Tong University	2013.7	Construction and Analysis of Extremophile Gene Database	B.Sc.	Graduate Studies in US
Xiaolin Hu	Shanghai Jiao Tong University	2013.7	The mechanism of Pyrococcus furiosus RecJ interac with ssDNA/ssRNA	B.Sc.	Shanghai Jiao Tong University
Yiwei Zhou	Shanghai Jiao Tong University	2013.7	Virtual Screening of New Drug Target on G- Protein Coupled Receptor Based on Statistical Models	B.Sc.	Academia Sinica
Huiying Yan	Shanghai Jiao Tong University	2013.7	MD Simulation Study of the M2 Proton Channel of Influenza Virus B	B.Sc.	Shanghai Jiao Tong University
Qian Cheng	Shanghai Jiao Tong University	2012.7	Metabolic Database of CYP450	B.Sc.	University of Michigan
Hao Zhang	Shanghai Jiao Tong University	2012.7	MD simulation of Amyloids	B.Sc.	Duke University
Qing Zhao	Shanghai Jiao Tong University	2011.7	Bioinformatics Studies of CYP 450	B.Sc.	Yale University
Detian Deng	Shanghai Jiao Tong University	2011.7	The Application of Bayesian Classification Method to Clinical Diagnosis and Prognosis	B.Sc.	Johns Hopkins University
Jiaqi Wu	Shanghai Jiao Tong University	2011.7	Active ingredients based on traditional Chinese medicine and multi-target drug molecule screening	B.Sc.	Taiwan
Leyan Chu	Shanghai Jiao Tong University	2010.10	Multi-Scale Modeling of Mucosal Cancer and Photodynamic Therapy	B.Sc.	MF in progress

Ge Gong	Shanghai Jiao Tong University	2010.7	MD simulations of FXA	B.Sc.	University of Southern California
Pengfei Liu	Wuhan University of Technology	2010.3	Modulation of the Spontaneous Curvature and Bending Rigidity of Lipid Membranes by Interfacially Adsorbed Amphipathic Peptides	B.Sc.	Suzhou Feng hua Environmental Engineering Co.,Ltd.
Roujie Chen	Shanghai Jiao Tong University	2009.7	CYP450 and Absorption, Distribution, Metabolism and Excretion (ADME) of drug molecules	B.Sc.	Columbia University
Yuan Shang	Shanghai Jiao Tong University	2009.7	Research of Substrate Selectivity A Mutant of Candida Antarctica Lipase B	B.Sc.	Hong Kong University of Science and Technology
Jianyi Liang	Shanghai Jiao Tong University	2009.7	Chinese Traditional Medicine Database	B.Sc.	Shanghai Jiao Tong University
Runfa Wu	Shanghai Jiao Tong University	2009.7	DNA-S Modification	B.Sc.	Graduate Studies in US
Yao Zhou	Shanghai Jiao Tong University	2009.7	Theoretical Studies of CYP450 Enzyme	B.Sc.	Shanghai Jiao Tong University
Binglin Yue	Shanghai Jiao Tong University	2009.7	MD Studies of Anti- Bacteria Peptides	B.Sc.	Graduate Studies in US
Jingyi Yan	Shanghai Jiao Tong University	2009.7	MD Simulation of CYP1E2	B.Sc.	University of British Columbia
Chengchen g Zhang	Shanghai Jiao Tong University	2009.7	MD Simulation of CYP1D6	B.Sc.	University of British Columbia
Yuqing Zhong	Zhengzhou University	2009.7	Biomedical Engineering, Electro- Neurophysiology	B.Sc.	University of North Texas
Mengchen Pu	Shanghai Jiao Tong University	2008.7	SNP Prediction of CYP450	B.Sc.	Graduate studies in Europe

Jisi Tang	Shanghai Jiao Tong University	2008.7	MD Simulation of CYP3A4	B.Sc.	Boston University
Shuhao Yu	Shanghai Jiao Tong University	2008.7	Neural Network Modeling and QSAR	B.Sc.	Institute of Biological Sciences, Academia Sinica
Yuanli Zhang	Northwest University	2008.7	QM/MM Studies of Enzyme Catalysis	B.Sc.	University of Oklahoma
He Zhang	Shanghai Jiao Tong University	2007.7	MD Simulations of Water Channels	B.Sc.	University of Wisconsin
Tianhang Chen	Shanghai Jiao Tong University	2021.9	Machine learnii	ıg	undergraduate student

Visiting Scholars/Post-Doc.

Name	Institutions	Duratio n	Field of Studies	Degree	Professional Title
Xuan Xiao	Jingdezhen College	2007- 2008	Bioinformatics	Ph.D.	Professor
Aixiu Li	Chinese Air Force Medical University	2007- 2008	Computer Aided Drug Design	Ph.D.	Professor
Yongxiang Shi	Shandong University	2009- 2010	MD Simulation of Protein-DNA Interactions	Ph.D.	Associate Professor
Wencheng Li	South China Institute of Technology	2007- 2008	Bioinformatics	Ph.D.	Post-doc Fellow
Bin Wang	Anhui Medical University	2012- 2013	Biostatistics	Ph.D.	Associate Professor
Guangfu Ji	Chinese Academy of Engineering Physics	2012- 2013	Condensed Matter Physics	Ph.D.	Professor
Fenglei Yang	Shanghai University	2012- 2013	Bioinformatics	Ph.D.	Lecturer
Linjing Zhao	Shanghai Technical University	2012- 2013	Bioinformatics	M.Sc.	Lecturer
Tiantian Li	Guizhou College of Chinese Medicine	2012- 2013	Computer Aided Drug Design	M. Sc.	Lecturer
Yonghong Zhang	Chongqing Medical College	2014- 2015	Computer Aided Drug Design	Ph.D.	Lecturer

Bo Zhou	Guizhou Medical College	2014- 2015	Computer Aided Drug Design	Ph.D.	Associated Professor
Diaxi Li	Shanghai University of Science and Technology	2015- 2016	MD Simulation of Thermo-stability of Protein	Ph.D.	Associated Professor
Meena Kumari	Shanghai Jiao Tong University	2015- 2017	Drug Combination	Ph.D.	Post-Doc
Faez Iqbal Khan	Henan Institute of Technology	2015- 2017	MD Simulation of Protein Catalysis	Ph.D.	Post-Doc
Yongkai Wei	Henan Institute of Technology	2015-	Ab initio MD of Energetic Materials	Ph.D.	Research Assistant Professor
Yuanyuan Qi	Henan Institute of Technology	2016-	Thermal Transportation	Ph.D.	Research Assistant Professor
Kotb Attia	Shanghai Jiao Tong University	2015- 2016	Personalized Drug Design	Ph.D.	Post-Doc
Ran Huang	The University of Akron	2015- 2020	Polymer Modeling	Ph.D.	Post-Doc
Gurudeeban Selvaraj	Henan Institute of Technology	2017-	Cancer Immunology	Ph.D.	Post-Doc
SatyavaniKaliam urthi	Henan Institute of Technology	2017-	Cancer Immunology	Ph.D.	Post-Doc
Aman Chandra Kaushik	Shanghai Jiao Tong University	2017- 2021	Protein-Protein Interaction	Ph.D.	Post-Doc.
Muhammad Tahir Khan	Shanghai Jiao Tong University	2018- 2020	Microbiology and Tuberculosis	Ph.D.	Post-Doc.
Heng Wang	Shanghai Jiao Tong University	2018- 2021	Synthetic Biology	Ph.D.	Post-Doc
Sathishkumar Chinnasamy	Shanghai Jiao Tong University	2018- 2021	CADD	Ph.D.	Post-Doc
Yanjing Wang	Shanghai Jiao Tong University	2019- 2021	CADD	Ph.D.	Post-Doc
Xue Jiang	Shanghai Jiaotong University	2020-	Single cell biology	Ph.D.	Post-Doc
Abbas Khan	Institute of AdvancedStudies, Chinese Academy of Sciences	2021-	AI drugs	PhD	Post-doc
Taimoor Khan	Institute of AdvancedStudies, Chinese Academy of Sciences	2021-	AI cancer	Ph.D.	Post-doc

Muhammad Shanghai Jiaotong 2019- Suleman University 2022 Covid-19 Ph.D. Post-doc
