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专家人才

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简历:

1988-6/1992 北京大学 学士
 1994-1/2000 美国罗格斯大学 (Rutgers University) 博士
 2000 - 2/2000 美国罗格斯大学 (Rutgers University) 博士后
 2000 - 8/2004 美国 Tularik 公司, 科学家, 助理科学家, 资深研究助理
 2004 - 10/2005 美国 Amgen 公司, 资深科学家
 2005 - 现在 中国科学院广州生物医药与健康研究院, 研究员

研究领域:

以结构生物学为主要手段, 紧密结合新药研发, 着眼于与人类重大疾病和重要生理功能相关的蛋白质及其复合体的结构与功能的关系, 阐明生物大分子与小分子化合物在结构与功能方面的分子作用机理, 为药物的分子设计提供思路; 同时, 通过解析生物大分子及其复合物的三维结构, 为药物设计提供一个分子水平上的模板。

承担科研项目情况:

1. 国家重点研发计划 (2017年), 项目负责人。
2. 973 (2012年, 2005年), 项目负责人。
3. 国家自然科学基金面上项目 (2018年、2016年、2012年、2009年), 项目负责人。
4. 国家自然科学基金联合基金项目 (2013年), 项目负责人。
5. 广东省自然科学基金团队项目 (2015), 项目负责人。
6. 广州市健康医疗协同创新重大专项 (2014), 项目负责人。

社会任职:

呼吸疾病国家重点实验室 副主任
 广东省生物医药计算重点实验室 副主任

获奖及荣誉:

代表论著:

1. Xu T.*, Gan Q., Wu B., Yin M., Xu J., Shu X., **Liu J.***. Molecular Basis for PI(3,5)P2 Recognition by SNX11, a Protein Involved in Lysosomal Degradation and Endosome Homeostasis Regulation. *J Mol Biol* 2020, 432(16): 4750-4761.
2. Xu J.*#, Tang X.#, Zhu Y., Yu Z., Su K., Zhang Y., Dong Y., Zhu W., Zhang C., Wu R., **Liu J.***. Structural studies reveal flexible roof of active site responsible for omega-transaminase CrmG overcoming by-product inhibition. *Commun Biol* 2020, 3(1): 455.
3. Makafe, G. G.#, Hussain, M.#, Surineni, G., Tan, Y., Wong, N. K., Julius, M., Liu, L., Gift, C., Jiang, H., Tang, Y., Liu, J., Tan, S., Yu, Z., Liu, Z., Lu, Z., Fang, C., Zhou, Y., Zhang, J., Zhu, Q., **Liu, J.*** and Zhang, T.* Quinoline Derivatives Kill Mycobacterium tuberculosis by Activating Glutamate Kinase. *Cell Chem Biol* 2019, 26, 1187-1194 e1185
4. Hussain, M.#, Adah, D.#, Tariq, M., Lu, Y., Zhang, J.* and **Liu, J.*** CXCL13/CXCR5 signaling axis in cancer. *Life Sci* 2019, 227, 175-186
5. Xu T, Zhou CZ, Xiao J.*, **Liu J.*** Unique Conformation in a Natural Interruption Sequence of Type XIX Collagen Revealed by Its High-Resolution Crystal Structure. *Biochemistry* 2018, 57(7): 1087-1095
6. Wang N, Zou Q, Xu J, Zhang J, **Liu J.*** Ligand binding and heterodimerization with retinoid X receptor alpha (RXRalpha) induce farnesoid X receptor (FXR) conformational changes affecting coactivator binding. *The Journal of biological chemistry* 2018, 293(47): 18180-18191
7. Xu T, Xie C., Yao D., Zhou CZ., **Liu J.***. Crystal structures of Aflatoxin-oxidase from *Armillariella tabescens* reveal a dual activity enzyme. *Biochemical and biophysical research communications* 2017, 494(3-4): 621-625
8. Xu J#, Zhang L#, Ye Y, Shan Y, Wan C, Wang J, Pei D, Shu X.*, **Liu J.*** SNX16 Regulates the Recycling of E-Cadherin through a Unique Mechanism of Coordinated Membrane and Cargo Binding. *Structure* 2017, 25(8): 1251-1263 e1255
9. Hou S, Xu T, Xu J, Qu L, Xu Y, Chen L, **Liu J.***. Structural basis for functional selectivity and ligand recognition revealed by crystal structures of human secreted phospholipase A2 group II-E. *Scientific reports* 2017, 7(1): 10815
10. Zhu Y#, Xu J#, Mei X#, Feng Z, Zhang L, Zhang Q, Zhang G, Zhu W*, **Liu J***, Zhang C*. Biochemical and Structural Insights into the Aminotransferase CrmG in Caerulomycin Biosynthesis. *ACS chemical biology* 2016, 11(4): 943-952
11. Hussain M, Lu Y, Liu YQ, Su K, Zhang J, **Liu J***, Zhou GB*. Skp1: Implications in cancer and SCF-oriented anti-cancer drug discovery. *Pharmacological research* 2016, 111: 34-42
12. Zhu L, Yang J, Li H, Sun H, **Liu J***, Wang J*. Conformational change study of dengue virus NS2B-NS3 protease using (19)F NMR spectroscopy. *Biochemical and biophysical research communications* 2015, 461(4): 677-680

13. Liu YQ[#], Wang XL[#], Cheng X[#], Lu YZ[#], Wang GZ[#], Li XC, Zhang J, Wen ZS, Huang ZL, Gao QL, Yang LN, Cheng YX*, Tao SC, **Liu J** *, Zhou GB*. Skp1 in lung cancer: clinical significance and therapeutic efficacy of its small molecule inhibitors. *Oncotarg* et 2015, 6(33): 34953-34967
14. Guo S[#], Xu J[#], Pavlidis IV, Lan D, Bornscheuer UT, **Liu J** *, Wang Y*. Structure of product-bound SMG1 lipase: active site gating implications. *The FEBS journal* 2015, 282(23), 4538-4547
15. Xu T, Xu J, Ye Y, Wang Q, Shu X, Pei D, **Liu J** *. Structure of human SNX10 reveals insights into its role in human autosomal recessive osteopetrosis. *Proteins* 2014, 82(12): 3483-3489
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17. Li H, Zhu L, Hou S, Yang J, Wang J*, **Liu J** *. An inhibition model of BPTI to unlinked dengue virus NS2B-NS3 protease. *FEBS letters* 2014, 588(17): 2794-2799
18. Xu J, Xu T, Wu B, Ye Y, You X, Shu X, Pei D, **Liu J** *. Structure of sorting nexin 11 (SNX11) reveals a novel extended phox homology (PX) domain critical for inhibition of SNX10-induced vacuolation. *The Journal of biological chemistry* 2013, 288(23): 16598-16605
19. Xu T[#], Liu L[#], Hou S, Xu J, Yang B, Wang Y*, **Liu J** *. Crystal structure of a mono- and diacylglycerol lipase from *Malassezia globosa* reveals a novel lid conformation and insights into the substrate specificity. *J Struct Biol* 2012, 178(3): 363-369
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22. Zhang Y, Xu T, Chen Q, Wang B, **Liu J** *. Expression, purification, and refolding of active human and mouse secreted group IIE phospholipase A. *Protein Expr Purif* 2011, 80(1): 68-73
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24. Lu N, Wang W, **Liu J** *, Wong CW*. Protein kinase C epsilon affects mitochondrial function through estrogen-related receptor alpha. *Cell Signal* 2011, 23(9): 1473-1478
25. Liu S, Dong J, Mei G, Liu G, Xu W, Su Z, **Liu J** *. Crystallization and preliminary crystallographic studies of a cysteine protease inhibitor from the human nematode parasite *Ascaris lumbricoides*. *Acta Crystallogr Sect F Struct Biol Cryst Commun* 2011, 67(Pt 2): 228-230
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30. Wang Z[#], Benoit G[#], **Liu J**, Prasad S, Aarnisalo P, Liu X, Xu H, Walker NP*, Perlmann T*. Structure and function of Nurr1 identifies a class of ligand-independent nuclear receptors. *Nature* 2003, 423(6939): 555-560.
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