

1997 年，在中国科学技术大学获生物学学士学位，并获化学物理学辅修学位，论文导师：刘海燕教授、施蕴渝院士。1999 年，在普林斯顿大学化学系获硕士学位。2001 年，在普林斯顿大学化学系获博士学位，是施一公教授培养的的第一个博士。2001-2003 年，在 Memorial Sloan Kettering 癌症中心做博士后，师从 Nikola Pavletich 院士。2003-2008 年在哈佛大学医学院附属儿童医院做博士后，师从贺熹教授。2008 年任上海交通大学教授，独立建立了上海交通大学第一个结构生物学实验室。参与了上海交通大学微生物代谢国家重点实验室的创建。2019 年承办了第六届华东地区结构生物学会会议。现为上海生物物理学会理事。合成生物学重点专项课题组长。

在 *Science*、*Nature*、*Nature Microbiology*、*Nature Communications* (3 篇)、*Molecular Cell* (2 篇)、*Cell Research*、*Nucleic Acids Research*、*Journal of Molecular Cell Biology*、*Cell Discovery*(3 篇)、*mBio*(2 篇)、*Molecular Microbiology* (2 篇) 等发表第一作者获通讯作者论文，被引用 2500 余次。APC-Asef 复合物结构论文被英国皇家学会院士 Mariann Bienz 选入生物医药类论文权威数据库 Faculty of 1000，并在此基础上与同事合作开发抑制结肠癌细胞迁移的化合物 (*Nature Chemical Biology*, 2017)。获教育部自然科学一等奖 (2016)、教育部新世纪优秀人才 (2009)、上海市东方学者 (2008)、上海市东方学者跟踪计划 (2014)、上海市曙光学者 (2008)、上海市浦江人才 (2009)、美国白血病与淋巴瘤学会 Special Fellowship (2005)。指导学生获上海市优秀博士论文 (2014)。

目前主要研究领域为合成生物学与生物物理化学，研究方向包括细胞生命重构、拟蛋白质设计、蛋白质类抗癌药物开发、蛋白质设计、与癌症相关的蛋白质机器的冷冻电镜结构等。

代表性论文 (*表示通讯作者, #表示第一作者):

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3. Liu L, Jiang S, Xing M, Chen C, Lai C, Li N, Liu G, Wu D, Gao H, Hong L, Tan P, Chen S, Deng Z, **Wu G***, Wang L*. Structural analysis of an L-cysteine desulfurase from an Ssp DNA phosphorothioation system. *mBio* 11(2):e00488-20. (2020)
4. Xiong X[#], **Wu G[#]**, Wei Y[#], Liu L, Zhang Y, Su R, Jiang X, Li M, Gao H, Tian X, Zhang Y, Hu L, Chen S, Tang Y, Jiang S, Huang R, Li Z, Wang Y, Deng Z, Wang J, Dedon PC, Chen S, Wang L. SspABCD-SspE is a phosphorothioation-sensing bacterial defence system with broad anti-phage activities. *Nature Microbiology* 5(7):917-928. (2020)
5. Liu G, Fu W, Zhang Z, He Y, Yu H, Wang Y, Wang X, Zhao YL, Deng Z, **Wu G***, He X*. Structural basis for the recognition of sulfur in phosphorothioated DNA. *Nature Communications* 9(1):4689. (2018)

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7. Kulaberoglu Y, Lin K, Holder M, Gai Z, Gomez M, Shifa BA, Mavis M, Hua L, Sharif AAD, Lujan C, Smith EJ, Bjedov I, Tapon N, **Wu G***, Hergovich A*. Stable MOB1 interaction with Hippo/MST is not essential for development and tissue growth control. *Nature Communications* 8:695. (2017)
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10. Zhang Z, Akyildiz S, Xiao Y, Gai Z, An Y, Behrens J*, **Wu G***. Structures of the APC-ARM domain in complexes with discrete Amer1/WTX fragments reveal that it uses a consensus mode to recognize its binding partners. *Cell Discovery* 1:15016. (2015)
11. **Wu G[#]**, Chen D[#], Tang H[#], Ren Y, Chen Q, Lv Y, Zhang Z, Zhao YL*, Yao Y, Xu P*. Structural insights into the specific recognition of N-heterocycle biodenitrogenation-derived substrates by microbial amide hydrolases. *Molecular Microbiology* 91:1009-1021. (2014)
12. Zhao G[#], **Wu G[#]**, Zhang Y, Liu G, Han T, Deng Z, He X*. Structure of the N-glycosidase MilB in complex with hydroxymethyl CMP reveals its Arg23 specifically recognizes the substrate and controls its entry. *Nucleic Acids Research* 42:8115-24. (2014)
13. Zhang Y, Fu L, Qi X, Zhang Z, Xia Y, Jia J, Jiang J, Zhao Y*, **Wu G***. Structural insight into the mutual recognition and regulation between Suppressor of Fused and Gli/Ci. *Nature Communications* 4:2608. (2013, 被引用 34 次)
14. Chen D, Tang H, Lv Y, Zhang Z, Shen K, Lin K, Zhao YL, **Wu G***, Xu P*. Structural and computational studies of the maleate isomerase Iso from *Pseudomonas putida* S16 reveal a breathing motion wrapping the substrate inside. *Molecular Microbiology* 87:1237-1244. (2013, 被引用 10 次)
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16. **Wu G***, Huang H, Abreu JG, He X*. Inhibition of GSK3 phosphorylation of β -catenin via phosphorylated PPPSPXS motifs of Wnt coreceptor LRP6. *PLoS One* 4, e4926. (2009, 被引用 131 次)
17. **Wu G**, He X*. Threonine 41 in β -catenin serves as a key phosphorylation relay residue in β -catenin degradation. *Biochemistry* 45, 5319-5323. (2006, 被引用 40 次)
18. **Wu G**, Liu C, He X*. Ozz, a new name on the long list of β -Catenin's nemeses.

Molecular Cell 13:451-3. (2004)

19. **Wu G**, Xu G, Schulman B, Jeffrey P, Harper JW, Pavletich NP*. Structure of a β -TrCP1-Skp1- β -Catenin complex: destruction-motif binding and lysine specificity of the SCF ^{β -TrCP1} ubiquitin ligase. *Molecular Cell* 11:1445-1456. (2003, 被引用 449 次)
20. **Wu G**[#], Chai J[#], Suber T, Wu JW, Du C, Wang X, Shi Y*. Structural basis of IAP recognition by Smac/DIABLO. *Nature* 408:1008-1012. (2000, 被 Novartis、Genentech 等公司作为设计抗癌药物的依据, 被引用 667 次)
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其他论文 (*表示通讯作者, #表示第一作者) :

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23. Yu H, Li J, Liu G, Zhao G, Wang Y, Hu W, Deng Z, **Wu G**, Gan J, Zhao YL, He X. DNA backbone interactions impact the sequence specificity of DNA sulfur-binding domains: revelations from structural analyses. *Nucleic Acids Research* gkaa574. (2020)
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26. Hu H, Wang L, Wang W, **Wu G**, Tao F, Xu P, Deng Z, Tang H. Regulatory Mechanism of Nicotine Degradation in *Pseudomonas putida*. *mBio* 10(3):e00602-19. (2019)
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 41. Tang H, Yao Y, Wang L, Yu H, Ren Y, **Wu G**, Xu P*. Genomic analysis of *Pseudomonas putida*: genes in a genome island are crucial for nicotine degradation. *Scientific Reports* 2:377. (2012)
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 56. Qin H, Srinivasula SM[#], **Wu G[#]**, Fernandes-Alnemri T, Alnemri ES, Shi Y*. Structural basis of procaspase-9 recruitment by the apoptotic protease-activating factor 1. *Nature* 399:549-557. (1999)
 57. Hua X, Miller Z, **Wu G**, Shi Y, Lodish HF*. Specificity in transforming growth factor β -induced transcription of the plasminogen activator inhibitor-1 gene: interactions of promoter DNA, transcription factor μ E3 and Smad proteins.

Proceedings of National Academy of Science USA 96:13130-13135. (1999)