简　历

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教育经历

- 2007.9.1 – 2013.1.10 **博士 清华大学,** 生物学

联合培养：香港中文大学（2008.9 – 2013.1）

指导教师：辛世文院士，何军贤教授

毕业论文题目：植物氮反应的蛋白质组学研究
答辩委员会：谢道昕，刘栋，刘进元，景海春，郝东云，何军贤，种康（主席）

- 2004.9 – 2007.7 **硕士 吉林大学**，生物化学与分子生物学

指导教师：郝东云教授

- 2000.9 – 2004.7 **本科 吉林大学**，生物技术

毕业设计指导教师：郝东云教授

研究经历

- 2013.10 至今 **博士后研究员** (Postdoctoral Researcher) **瑞士洛桑联邦理工学院**

指导教师 Johan Auwerx 教授：欧洲分子生物学会院士 (EMBO member); 线粒体代谢与系统生物学领域专家 (Web of Science h-index: 116); *Nature* 杂志顾问, *Science*, *Cell* 等杂志编委; 2016 年瑞士最高科学奖 – 马塞尔·本努瓦奖 (Marcel Benoist Prize) 得主。

研究项目：植物细胞线粒体逆境生物学及其与细胞核之间信号转导机制的研究
线粒体功能，代谢类疾病以及衰老的系统遗传学研究

- 2008.9 – 2013.8 **研究助理** (Research Assistant) **香港中文大学**

指导教师辛世文教授：中国工程院院士，农业生物技术专家，专注于提升稻米产量和营养成分。联合导师何军贤教授：植物激素研究领域专家，在对植物类固醇激素油菜素内酯 (Brassinosteroids, BRs) 信号转导调控研究中取得突破性进展。

研究项目：植物氮信号转导的蛋白质组学与磷酸化蛋白质组学研究

课题申请训练

博士后研究期间参与多个项目申请书的撰写，包括欧盟高级资助计划 (ERC advanced grant, 250 万欧元), 瑞士国家自然科学基金 (SNSF, 150 万瑞士法郎)，以及与谷歌，强生，礼来 (Eli Lilly), 安斯泰来制药 (Astellas Pharma) 等高科技公司的合作项目。

学术活动

“Keystone Symposia” Plant Signaling: Molecular Pathways and Network Integration
Granlibakken Tahoe, Tahoe City, California, Jan 21-24, 2018
海报：Systems phytohormone responses to mitochondrial proteotoxic stress.

简　历

EPFL Life Sciences - EMBO Press Symposium 2017, Frontiers in Metabolism - From systems physiology to precision medicine
SwissTech Convention Center- EPFL, Lausanne, Switzerland, Oct 24-26, 2017 出席

3rd International SystemsX.ch Conference on Systems Biology
ETH Zurich, Switzerland, Sep 4-7, 2017
海报：Systems phytohormone responses to mitochondrial proteotoxic stress.

Life Sciences Switzerland LS2 Annual Meeting
Irchel Campus, University of Zurich, Switzerland, Feb 2-3, 2017
报告：Systems approach to understand the impact of genetics and diet on aging.

"Gordon Research Conference" Mitochondria & Chloroplasts
Mount Snow, West Dover, Vermont, Jun 19-24, 2016
海报：Ethylene signaling is involved in the plant retrograde response to mitochondrial stress.

LIMNA (Lausanne Integrative Metabolism and Nutrition Alliance) Symposiums
EPFL, Lausanne, Switzerland, 2014-2017 系列会议，出席

审稿经历

ACS 杂志社 *Journal of Proteome Research, Journal of Agriculture and Food Chemistry* 并共同审阅 *Cell, Molecular Cell, PLOS Genetics* 等杂志论文以及国际基金申请书

发表论文

Web of Science 统计: h-index: **11**, 总引用数: **470**, 每篇平均引用数: **21**. (2017/12/30)

第一作者 (* 共同第一作者)

[1] **Wang X** & Auwerx J. Systems phytohormone responses to mitochondrial proteotoxic stress. ***Molecular Cell*** 2017; 68: 540-51. (IF: 14.713)

[2] Li H*, **Wang X***, Rukina D, Huang Q, Lin T, Sorrentino V, Zhang H, Sleiman MB, Arends D, McDaid A, Luan P, Ziari N, Velázquez-Villegas LA, Gariani K, Kutalik Z, Schoonjans K, Radcliffe RA, Prins P, Morgenthaler S, Williams RW, Auwerx J. An integrated systems genetics and omics toolkit to probe gene function. ***Cell Systems*** 2017 (In Press) (IF: 8.406)

[3] **Wang X***, Ryu D*, Houtkooper RH, Auwerx J. Antibiotic use and abuse: a threat to mitochondria and chloroplasts with impact on research, health, and environment. ***BioEssays*** 2015; 37(10):1045-53. (IF: 4.441, cites: 10)

[4] **Wang X***, Bian Y*, Cheng K, Gu LF, Ye M, Zou H, Sun SSM, He JX. A Large-scale protein phosphorylation analysis reveals novel phosphorylation motifs and phosphoregulatory networks in *Arabidopsis*. ***Journal of Proteomics*** 2013; 78:486-98. (IF: 3.914, cites: 38)

[5] Wang YD*, **Wang X***, Ngai SM, Wong YS. Comparative proteomics analysis of selenium responses in selenium-enriched rice grains. ***Journal of Proteome Research*** 2013; 12:808-20. (IF: 4.268, cites: 11)

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[7] **Wang X**, Bian Y, Cheng K, Zou H, Sun SSM, He JX. A comprehensive differential proteomic study of nitrate deprivation in *Arabidopsis* reveals complex regulatory networks of plant nitrogen responses. ***Journal of Proteome Research*** 2012; 11:2301-15. (IF: 4.268, cites: 37)

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其他论文 (* 共同第一作者)

[9] Dahlmans D, Houzelle A, Andreux P, Jörgensen J A., **Wang X**, de Windt L J., Schrauwen P, Auwerx J, Hoeks J. An unbiased silencing screen in muscle cells identifies miR-320a, miR-150, miR-196b, and miR-34c as regulators of skeletal muscle mitochondrial metabolism. ***Molecular Metabolism***. 2017; 6(11):1429-1442. (IF: 6.799)

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[11] Stein S*, Lemos V*, Xu P, Demagny H, **Wang X**, Ryu D, Jimenez V, Bosch F, Lüscher TF, Oosterveer MH, Schoonjans K. Impaired SUMOylation of nuclear receptor LRH-1 promotes nonalcoholic fatty liver disease. ***Journal of Clinical Investigation***. 2017; 127(2):583-592. (IF: 12.784)

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[14] Jha P, **Wang X**, Auwerx J. Analysis of mitochondrial respiratory chain supercomplexes using blue native polyacrylamide gel electrophoresis (BN-PAGE). ***Current Protocols in Mouse Biology***. 2016; 6(1):1-14. (cites: 10)

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