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EDUCATION/TRAINING

2008	Postdoc	Molecular Biology, Harvard Medical School, USA
2003	Ph.D.	Plant Pathology, Kansas State University, USA
1998	M.S.	Genetics, Chinese Academy of Sciences, China
1993	B.S.	Plant Genetics & Breeding, China Agricultural University, China

POSITIONS & EMPLOYMENT

2009 – Present	Assistant (2009-2013)/Associate (2013-2017) /Professor (2017-) Department of Biochemistry and Biophysics Institute for Plant Genomics and Biotechnology Faculty of Molecular & Environmental Plant Sciences Texas A&M University, College Station, TX

OTHER POSITIONS AND PROFESSIONAL MEMBERSHIPS

2016 – Present	Senior Editor, Molecular Plant-Microbe Interactions
2016	Editor, Plant Pattern Recognition Receptor Methods and Protocols Methods in Molecular Biology
2012 – 2017	Editorial Board, Molecular Plant Pathology

HONORS & AWARDS

- Vice chancellor's Award in Excellence for Research, Texas A&M University, 2018
- National Science Foundation Career Award, 2013
- American Society of Plant Biologists Early Career Award, 2008

SELECTED PUBLICATIONS

- Hou, S., Jamieson, P., and **He, P.** (2018) The cloak, dagger, and shield: proteases in plant-pathogen interactions. ***Biochemical Journal*** 475: 2491-2509.
- Jamieson, P.A., Shan, L., and **He, P.** (2018) Plant cell surface molecular cypher: receptor-like proteins and their roles in immunity and development. ***Plant Science*** 274: 242-251.
- Shan, L., and **He, P.** (2018) Pipped at the post: pipecolic acid derivative identified as SAR regulator. ***Cell*** 173:286-287.
- Mang H., Feng, B., Hu, Z., Boisson-Dernier, A., Franck, C.M., Meng, X., Xu, G., Wang, T., Shan, L., and **He, P.** (2017) Differential regulation of two-tiered plant immunity and sexual reproduction by ANXUR receptor-like kinases. ***The Plant Cell.*** 29: 3140–3156.
- Li, F., Li, M., Wang, P., Cox, K. L., Duan, L., Dever, J.K., Shan, L. Li, Z., and **He, P.** (2017) Regulation of cotton drought responses by a MAP kinase cascade-mediated phosphorylation of GhWRKY59. ***New Phytologist*** 215: 1462-1475.
- Feng, B., Liu, C., Shan, L. and **He, P.** (2016) ADP-ribosylation takes control in plant-bacterium interactions ***PLoS Pathog***. 12: e1005941.
- Feng, B., Ma, S., Chen, S., Zhu, N., Zhang, S., Yu, B., Yu, Y., Le, B., Chen, X., Dinesh-Kumar, S.P., Shan, L., and **He, P.** (2016) PARylation of the forkhead-associated domain protein DAWDLE regulates plant immunity. ***EMBO Reports*** 17:1799-1813.
- Li, B., Meng, X., Shan, L., and **He, P.** (2016) Transcriptional regulation of plant pattern-triggered immunity. ***Cell Host & Microbe*** 19:641-650.
- de Oliveira, M.V.V., Xu, G., Li, B., de Souza Vespoli, L., Meng, X., Chen, X., Yu, X., de Souza, S.A., Intorne, A.C., de A. Manhães, A.M.E., Musinsky, A.L., Koiwa, H., de Souza Filho, G.A., Shan, L., and **He, P.** (2016). Specific control of Arabidopsis BAK1/SERK4-regulated cell death by protein glycosylation. ***Nature Plants*** 2:15218.
- Feng, B., Liu, C., Oliveira, M.V.V., Intorne, A.C., Li, B., Babilonia, K., Filho, G.A.S., Shan, L., and **He, P.** (2015) Protein poly(ADP-ribosyl)ation regulates *Arabidopsis* immune gene expression and defense responses. ***PLoS Genetics*** 11(1): e1004936. doi:10.1371/journal.pgen.1004936.

- Li, F., Cheng, C., Cui, F., Oliveira, M.V.V., Yu, X., Meng, X., Intorne, A.C., Babilonia, K., Li, M., Chen, X., Ma, X., Xiao, S., Zeng, Y., Fei, Z., Metz, R., Johnson, C.D., Koiwa, H., Sun, W., Li, Z., Filho, G.A.S., Shan, L., and **He, P.** (2014) Modulation of RNA polymerase II phosphorylation downstream of pathogen perception orchestrates plant immunity. ***Cell Host & Microbe*** 16:748-758.
- Gao, X., Chen, X., Lin, W., Lu, D., Niu, Y., Li, L., Cheng, C., McCormack, M., Sheen, J., Shan, L., and **He, P.** (2013) Bifurcation of *Arabidopsis* NLR Immune Signaling via Ca2+-Sensor Protein Kinases. ***PLoS Pathog* 9**, e1003127.
- Cheng, C, Gao, X, Feng, B, Sheen, J, Shan, L, and **He, P**. (2013) Plant immune response to pathogens differs with changing temperatures. ***Nat Commun.*** 4:2530. doi: 10.1038/ncomms3530
- Gao, X., Wheeler, T., Li, Z., Kenerley, C., **He, P.,** and Shan, L. (2011) Silencing GhNDR1 and GhMKK2 compromised cotton resistance to *Verticillium wilt*. ***The Plant Journal.*** 66:293-305.
- Lu, D., Lin, W., Gao, X., Wu, S., Cheng, C., Avila, J., Heese, A., Devarenne, T., **He, P.***, and Shan, L.* (2011) Direct ubiquitination of pattern recognition receptor FLS2 attenuates plant innate immunity. ***Science.*** 332:1439-1442. (*Co-corresponding authors).
- Lu, D., Wu, S., Gao, X., Zhang, Y., Shan, L. and **He, P.** (2010) A receptor-like cytoplasmic kinase BIK1 associates with flagellin receptor complex to initiate plant innate immunity. ***Proc Natl Acad Sci U S A***. 107:496-501.
- Boudsocq, M., Willmann M. R., McCormack, M., Lee, H., Shan L., **He P.,** Bush, J., Cheng, S., and Sheen J., (2010) Differential innate immune signalling via Ca^{2+} sensor protein kinases. ***Nature.*** 464: 418-422.
- Shan, L., **He, P**., Li, J., Heese, A., Peck, S. C., Nurnberger, T., Martin, G. B., and Sheen, J. (2008). Bacterial effectors target the common signaling partner BAK1 to disrupt multiple MAMP receptor-signaling complexes and impede plant immunity. ***Cell Host Microbe*** 4:17-27.
- Sheen, J., and **He, P.** (2007) Nuclear actions in innate immune signaling. ***Cell*** 128:821-823.
- **He, P.,** Shan, L., Lin, N.-C., Martin, G., Kemmerling, B., Nurnberger, T., and Sheen, J. (2006). Specific bacterial suppressors of MAMP signaling upstream of MAPKKK in Arabidopsis innate immunity. ***Cell*** 125:563-575.
- **He, P.,** Chintamanani, S., Chen, Z., Zhu, L., Kunkel, B. N., Alfano, J. R., Tang, X., and Zhou, J.-M. (2004). Activation of a COI1-dependent pathway in Arabidopsis by Pseudomonas syringae type III effectors and coronatine. ***The Plant Journal*** 37:589-602.
- **He, P.,** Friebe, B. R., Gill, B. S., and Zhou, J.-M. (2003). Allopolyploidy alters gene expression in the highly stable hexaploid wheat. ***Plant Molecular Biology*** 52:401-414.
- **He, P.,** Li, J., Zheng, X., Shen, L., Lu, C., Chen, Y., and Zhu, L. (2001). Comparison of molecular linkage maps and agronomic trait loci between DH and RIL populations derived from the same rice cross. ***Crop Science*** 41: 1240-1246.
- **He, P.,** Li, S., Qian, Q., Ma, Y., Li, J., Wang, W., Chen, Y., and Zhu L. (1999). Genetic analysis of rice grain quality. ***Theor. Appl. Genet.*** 98: 502-508.
- **He, P.,** Shen, L., Lu, C., Chen Y., and Zhu L. (1998). Analysis of quantitative trait loci which contribute to anther culturability in rice (Oryza sativa L.). ***Molecular Breeding*** 4: 165-172.