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学习和工作简历	<p>教育经历：</p> <p>2011-09 至 2014-06 南京农业大学，园艺学院，博士</p> <p>2004-09 至 2007-06 南京农业大学，园艺学院，硕士</p> <p>2000-09 至 2004-06 山西农业大学，园艺学院，学士</p> <p>工作经历：</p> <p>2016-12 至今 南京农业大学，园艺学院，作物遗传与种质创新国家重点实验室，副教授</p> <p>2019-09 至 2020-09 美国康奈尔大学，植物遗传育种学系，访问学者</p> <p>2015-02 至 2016-12 南京农业大学，园艺学院，作物遗传与种质创新国家重点实验室，讲师</p> <p>2007-08 至 2015-02 温州科技职业学院，农业与生物技术系，蔬菜研究所，讲师</p> <p>主要研究方向：重点以萝卜等十字花科蔬菜为研究对象，系统进行种质资源评价与性状鉴定、优异基因发掘与种质创新、育种技术提升与品种选育等相关工作。</p>				
科研项目	<p>国家自然科学基金项目：萝卜耐盐性关键基因鉴定与生物学功能分析 主持</p> <p>中国博士后科学基金面上项目：萝卜耐盐性状关键基因 SOS1 与 NHXs 分子特征及功能解析 主持</p> <p>江苏省重点研发计划专项资金项目：优质晚抽薹萝卜新品种“南春白 8 号”主持</p> <p>江苏省农业科技自主创新资金项目：克服萝卜自交不亲和技术创新 主持</p> <p>农业部重点实验室开放课题：萝卜 Na^+/H^+ 逆向转运蛋白基因分离克隆及其生物学功能验证 主持</p> <p>中央高校基本业务费项目：萝卜资源保存与研究 主持</p>				
发表论文	<p>Wang Yan[#]; Ying Jiali[#]; Zhang Yang; Xu Liang; Zhang Wanting; Ni Meng; Zhu Yuelin[*]; Liu Liwang[*]; Genome-wide identification and functional characterization</p>				

	<p>of the cation proton antiporter (CPA) family related to salt stress response in radish (<i>Raphanus sativus</i> L.), <i>Int J Mol Sci</i>, 2020, 21: 8262.</p> <p>Wang Yan[#]; Song Zhaojun[#]; Zhang Wei[#]; Xu Liang; Su Xiaojun; Muleke Everlyne M'mbone; Liu Liwang[*]; Identification and characterization of expressed TIR-and non-TIR-NBS-LRR resistance gene analogous sequences from radish (<i>Raphanus sativus</i> L.) de novo transcriptome, <i>Sci Hortic</i>, 2017, 216: 284-292.</p> <p>Wang Yan[#]; Xu Liang[#]; Tang Mingjia; Jiang Haiyan; Chen Wei; Zhang Wei; Wang Ronghua; Liu Liwang[*]; Functional and integrative analysis of the proteomic profile of radish root under Pb exposure, <i>Front Plant Sci</i>, 2016, 7: 1871.</p> <p>Wang Yan[#]; Pan Yan[#]; Liu Zhe; Zhu Xianwen; Zhai Lulu; Xu Liang; Yu Rugang; Gong Yiqin; Liu Liwang[*]; De novo transcriptome sequencing of radish (<i>Raphanus sativus</i> L.) and analysis of major genes involved in glucosinolate metabolism, <i>BMC Genomics</i>, 2013, 14:836.</p> <p>Wang Yan[#]; Shen Hong[#]; Xu Liang[#]; Zhu Xianwen; Li Chao; Zhang Wei; Xie Yang; Gong Yiqin; Liu Liwang[*]; Transport, ultrastructural localization, and distribution of chemical forms of lead in radish (<i>Raphanus sativus</i> L.), <i>Front Plant Sci</i>, 2015, 6.</p> <p>Wang Yan, Xu Liang, Chen Yinglong, Shen Hong, Gong Yiqin, Limera Cecilia, Liu Liwang[*]; Transcriptome profiling of Radish (<i>Raphanus sativus</i> L.) root and identification of genes involved in response to Lead (Pb) stress with next generation sequencing. <i>PLoS One</i>, 2013, 8(6): e66539 .</p> <p>Wang Yan[#]; Liu Wei[#]; Shen Hong, Chen Yinglong, Zhai Lulu, Xu Liang, Wang Ronghua, Gong Yiqin, Limera Cecilia, Liu Liwang[*]; Identification of radish (<i>Raphanus sativus</i> L.) miRNAs and their target genes to explore miRNA-mediated regulatory networks in lead(Pb) stress responses by high-throughput sequencing and degradome analysis. <i>Plant Mol Biol Rep</i>, 2014, 33, 358-376.</p> <p>Xu Liang[#]; Wang Yan[#]; Zhang Fei; Tang Mingjia; Chen Yinglong; Wang Jin; Karanja Kinuthia Bernard; Luo Xiaobo; Zhang Wei; Liu Liwang[*]; Dissecting root proteome changes reveals new insight into cadmium stress response in radish (<i>Raphanus sativus</i> L.), <i>Plant Cell Physiol</i>, 2017, 58: 1901–1913.</p> <p>Luo Xiaobo[#]; Xu Liang[#]; Wang Yan; Dong Junhui; Chen Yinglong; Tang Mingjia; Fan Lianxue; Zhu Yuelin; Liu Liwang[*]; An ultra-high density genetic map provides insights into genome synteny, recombination landscape and taproot skin color in radish (<i>Raphanus sativus</i> L.), <i>Plant Biotechnol J</i>, 2020, 18: 274–286.</p> <p>Fan Lianxue, Wang Yan, Xu Liang, Tang Mingjia, Zhang Xiaoli, Ying Jiali, Li Cui, Dong Junhui, Liu Liwang*. Genome-wide association study uncovers critical role of <i>RsPAP2</i> gene in red-skinned <i>Raphanus sativus</i> L., <i>Hortic Res</i>, 2020, 7:</p>
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