

外国留学生研究生导师情况表

Resume of Supervisor (中英文版)

导师姓名 Name of Supervisor	马啸 Xiao Ma	导师类别 Supervisor Level	博导 ■ 硕导 ■ Doctor Master
最后学历 Highest Degree	博士 PhD	职称 Professional Title	教授 professor
院所 College/Institute	动物科技学院 College of Animal Science and Technology		
学科 Discipline	草学 Grassland science		
邮箱 Email	maroar@126.com		
出国经历 Experience Abroad	2010.09-2010.12 国际家畜研究所 (ILRI) 牧草中心访问学者 (2010.09-2010.12 As a visiting scholar at Forage genebank of International Livestock Research Institute, Addis Ababa, Ethiopia) 2011.04-2012.04 日本北海道大学访问学者 (2011.04-2012.04 As a visiting scholar at Field Science Center for Northern Biosphere in Hokkaido University, Sapporo, Japan)		
研究方向 Research Fields	牧草遗传育种 Genetics and breeding of Forage/Fodder crops		
代表性成果 (10 项以内) Publications	[1] Ma X , SY Chen, XQ Zhang, et al. Assessment of Genetic Diversity of Siberian wildrye using Gliadin Markers. <i>Molecules</i> , 2012. 17(4):4424-34. [2] Ma X , XQ Zhang, YH Zhou, SQ Bai, et al. Assessing genetic diversity of <i>Elymus sibiricus</i> populations from Qinghai-Tibet Plateau by ISSR markers. <i>Biochem Syst Ecol</i> , 2008, 36: 514 - 522. [3] Zhang C, Zhang J, Y. Fan, X. Ma* . 2017. Genetic Structure and Eco-Geographical Differentiation of Wild Sheep Fescue in Xinjiang, Northwest China. <i>Molecules</i> , 22(8):1316. (corresponding author) [4] Fan Y, Zhang C, Wu W, Ma X* . Analysis of Genetic Diversity and Structure Pattern of <i>Indigofera pseudotinctoria</i> in Wushan Mountains Using AFLP Markers. <i>Molecules</i> 2017, 22, 1734. (corresponding author) [5] Xiao-yan Gu, Zhi-hui Guo, Xiao Ma* , et al. Population genetic variability and structure of <i>Elymus breviaristatus</i> endemic to QT Plateau inferred from SSR markers. <i>Biochem Syst Ecol</i> , 2015,58:247 - 256.(corresponding author) [6] Guo ZH, Fu KX, Zhang XQ, X. Ma* , et al. SSRs transferability and genetic diversity of three allogamous ryegrass species. <i>Comptes Rendus Biologies</i> , 2016,39(2): 60 - 67. (corresponding author) [7] Guo ZH, Fu KX, Zhang XQ, Bai SQ, X. Ma* , et al. Molecular insights into the genetic diversity of <i>Hemarthria compressa</i> germplasm collections native to southwest China. <i>Molecules</i> 2014, 19(12), 21541-59. (corresponding author) [8] Fu K., Guo Z., Zhang X., Xiao Ma* et al. Insight into the genetic variability analysis and cultivar identification of tall fescue by using SSR markers. <i>Hereditas</i> , 2016,153(1):9-15. (corresponding author) [9] Sun M, Zhang CL, Zhang XQ, Ma Xiao* et al. AFLP assessment of Genetic variability and relationships in an Asian wild germplasm collection of <i>Dactylis glomerata</i> L. <i>Comptes Rendus Biologies</i> , 2017, 40(3):145-155. (corresponding author) [10] Ma X , Chen SY, Bai SQ, Zhang XQ, et al. RAPD analysis of genetic diversity and population structure of <i>Elymus sibiricus</i> (Poaceae) native to the southeastern Qinghai-Tibet Plateau, China. <i>Genetics and Molecular Research</i> , 2012, 11(3):2708-2718.		