

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	李利 Li Li	导师类别 Supervisor Level	博导 <input checked="" type="checkbox"/> 硕导 <input checked="" type="checkbox"/> Doctor <input checked="" type="checkbox"/> Master <input checked="" type="checkbox"/>
最后学历 Highest Degree	博士研究生 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	动物遗传育种与繁殖 Animal genetics, breeding and reproduction		
邮箱 Email	lily@sicau.edu.cn		
出国经历 Experience Abroad	2009.09—2010.09 澳大利亚悉尼大学访问学者 Sep. 2009—Sep. 2010, Sydney University(AUS)		
研究方向 Research Fields	1.羊的遗传多样性研究 Genetic diversity in Goat & Sheep 2.羊重要经济性状的遗传机制 Genetic mechanism of some important economic traits in Goat & Sheep		
代表性成果 (10 项以内) Publications	<p>Li L, Jiang J, Wang LJ, Zhong T, Chen B, Zhan SY, Zhang HP, Du LX. Expression patterns of peroxisome proliferator-activated receptor gamma 1 versus gamma 2, and their association with intramuscular fat in goat tissues. <i>Gene</i>. 2013,528(2): 195-200.(IF=2.196)</p> <p>Li L, Su HW, Wei CH, Ren HX, Xu LY, Zhao FP, Zhang L, Wang LJ, Zhong T, Zhang HP, Liu WZ, Du LX .Comparison of Muscle Characteristics and Underpinning Mechanisms between Texel and Ujumqin Sheep Aged from Day 70 to 135 of Gestation. <i>Livestock Science</i>,2016,192:8-16. (IF=1.293)</p> <p>Zhan SY, Dong Y, Zhao W, Guo JZ, Zhong T, Wang LJ, Li L , Zhang HP. Genome-wide Identification and Characterization of Long Non-coding RNAs in Developmental Skeletal Muscle of Fetal Goat. <i>BMC Genomics</i>,2016,17:666 (Corresponding author, IF=3.986)</p> <p>Zhan SY, Zhao W, Song TZ, Dong Y, Guo JZ, Cao JX, Zhong T, Wang LJ, Li L, Zhang HP.Dynamic transcriptomic analysis in hircine <i>longissimus dorsi</i> muscle from fetal to neonatal development stages. <i>Funct Integr Genomics</i>,2017,DOI 10.1007/ s10142- 017-0573-9. (Corresponding author, IF=3.496)</p> <p>Yang L, Xu LY, Zhou Y, Liu M, Wang L, Kijas J W, Zhang HP, Li L, Liu GE. Diversity of copy number variation in a worldwide population of sheep.<i>Genomics</i>, 2017, http://dx.doi.org/ 10.1016 /j.ygeno.2017.09.005. (Corresponding author, IF= 2.801)</p> <p>Yang CT, Li L, Zhong T, Wang L.J, Zhang HP. Characterization of the</p>		

complete mitochondrial genome sequence of Ujumuin sheep (*Ovis aries*). *Mitochondrial DNA* , *Mitochondrial DNA* , 2017,28(3): 315-316.(Corresponding author, IF=1.209)

Xu LY, Zhao FP, Ren HX, **Li Li** ,Lu J, Liu JS, Zhang SF, Liu GE, Song JZ, Zhang L, Wei CH, Du LX. Co-expression analysis of fetal weight-related genes in ovine skeletal muscle during mid and late fetal development stages. *Int J Biol Sci*,2014,10(9):1039-1050. (Co-first author, IF=4.372)

Wei CH, **Li Li**, Su HW, Xu LY, Lu J, Zhang L, Liu WZ ,Ren HX, Du LX. Identification of the crucial molecular events during the large-scale myoblast fusion in sheep. *Physiological Genomics*,2014,46(12): 429-440. (Co-first author, IF=2.812)

Wang LJ, **Li L**, Jiang J, Wang Y, Zhong T, Chen Y, Wang Y , Zhang HP. Molecular characterization and different expression patterns of the FABP gene family during goat skeletal muscle development. *Mol Biol Rep*, 2015,42(1),201-207.(Co-first author, IF=1.958)

Cao JX, Wei CH, Liu DM, Wang HH, Wu MM, Xie ZY, Capellini TD., Zhang L, Zhao FP, **Li L**, Zhong T, Wang LJ, Lu J, Liu RZ, Zhang SF, Du YF, Zhang HP, Du LX. DNA methylation Landscape of body size variation in sheep. *Scientific Reports*, 2015(5):13950. (IF=5.578)