Sunday, July 15

Late-Breaking Original Research Chair: Anthony V. Capuco, Bovine Functional Genomics Lab 122AB

3:00 PM	LB1	Identification of a genomic region associated with severe combined immunodeficiency in pigs. E. H. Waide*¹, C. K. Tuggle¹, D. M. Thekkoot¹, N. Boddicker¹, R. R. R. Rowland², C. R. Wyatt², and J. C. M. Dekkers¹, ¹lowa State University, Ames, ²Kansas State University, Manhattan.
3:15 PM	LB2	Characterization of genetic variation within the somatotropic axis in DNA pools of beef and dairy cattle divergent for milk production, size, fertility, and immune response. M. P. Mullen*1,2, C. Creevey³, D. P. Berry⁴, M. S. McCabe², D. J. Howard¹, D. A. Magee⁵, M. C. Lucy⁶, D. E. MacHugh⁵,7, and S. M. Waters², ¹Animal and Bioscience Research Department, Animal and Grassland Research and Innovation Centre, Teagasc, Athenry, Co. Galway, Ireland, ²UCD School of Veterinary Medicine, University College Dublin, Belfield, Dublin 4, Ireland, ³Animal and Bioscience Research Department, Animal and Grassland Research and Innovation Centre, Teagasc, Grange, Co. Meath, Ireland, ⁴Animal and Bioscience Research Department, Animal and Grassland Research and Innovation Centre, Teagasc, Moorepark, Fermoy, Cork, Ireland, ⁵UCD School of Agriculture and Food Science, University College Dublin, Belfield, Dublin 4, Ireland, ⁶Department of Animal Sciences, University of Missouri, Columbia, ¬UCD Conway Institute of Biomolecular and Biomedical Research, University College Dublin, Belfield, Dublin 4, Ireland.
3:30 PM	LB3	Imputation of microsatellite alleles from dense SNP genotypes for paternal verification. M. C. McClure ¹ , T. S. Sonstegard ¹ , G. R. Wiggans ¹ , A. Van Eenennaam ² , K. Weber ² , C. Penedo ² , and C. P. Van Tassell* ¹ , 1USDA-ARS, Beltsville, MD, ² University of California, Davis.
3:45 PM	LB4	Improvement of feed efficiency through diet and breed-dependent genetic polymorphisms. N. V. L. Serão,* J. E. Beever, D. B. Faulkner, and S. L. Rodriguez-Zas, <i>University of Illinois at Urbana-Champaign</i> .
4:00 PM	LB5	Mixed rumen microbes respond to excess carbohydrate by synthesizing glycogen and spilling energy. T. J. Hackmann,* K. L. Backus, and J. L. Firkins, <i>The Ohio State University, Columbus</i> .
4:15 PM	LB6	Fine mapping and discovery of recessive mutations that cause abortions in dairy cattle. P. M. VanRaden¹, D. J. Null*¹, T. S. Sonstegard², H. A. Adams³, C. P. Van Tassell², and K. M. Olson⁴, ¹Animal Improvement Programs Laboratory, ARS, USDA, Beltsville, MD, ²Bovine Functional Genomics Laboratory, ARS, USDA, Beltsville, MD, ³Institute for Genomic Biology, University of Illinois, Urbana, ⁴National Association of Animal Breeders, Columbia, MO.
4:30 PM	LB7	Maternal dietary energy source during gestation affects expression of imprinted genes in fetal tissues in sheep. X. Lan, R. Gambra, M. A. Berg, E. J. Cretney, H. Khatib, and A. E. Radunz,* <i>University of Wisconsin-Madison, Madison.</i>
4:45 PM	LB8	Potential for post-extraction algal residue to replace cottonseed meal as a protein supplement to grazing cattle. M. L. Drewery,* J. E. Sawyer, and T. A. Wickersham, <i>Texas A&M University, College Station.</i>