P0581 Validation of a Low Density SNP Panel for Breed Certification Testing in Brazilian Sheep (*Ovis aries*) Breeds as a Tool for Flock Genetic Management

Fabio Danilo Vieira , Embrapa Informática Agropecuária, Campinas, SP, Brazil
Michel E Beleza Yamagishi , Embrapa Informática Agropecuária, Campinas, SP, Brazil
Thaisa S. Lacerda , Universidade de Brasilia, Programa de Pós-Gradução em Ciências Animais, Brasilia, Brazil
Carolina Vasconselos , Universidade de Brasilia, Programa de Pós-Gradução em Ciências Animais, Bedro Tanno , Embrapa Recursos Genéticos e Biotecnologia, Brasilia, Brazil
Roberto Hiroshi Higa , Embrapa Informática Agropecuária, Campinas, SP, Brazil
Concepta M. McManus , Universidade Federal do Rio Grande do Sul, Faculdade de Agronomia, Porto Alegre, Brazil
Paulo Luis Carneiro , Universidade Estadual do Sudoeste da Bahia
Hymerson C Azevedo , Embrapa Tabuleiros Costeiros
Olivardo Faco , Embrapa Caprinos e Ovinos
Carlos J H Souza , Embrapa Pecuaria Sul
Adriana M Araujo , Embrapa Meio Norte
Vera MV Martins , Universidade Estadual de Santa Catarina, Lages
Alexandre Caetano , EMBRAPA Recursos Geneticos e Biotecnologia, Brasilia, Brazil
The International Sheep Genomics Consortium , ISGC

Samuel Rezende Paiva , EMBRAPA Recursos Geneticos e Biotecnologia, Brasilia, Brazil

Sheep production activities are growing rapidly in Brazil and are based on locally adapted breeds in addition to commercial breeds available internationally. Contemporary methods for genetic management of flocks are in high demand and motivated the work for development of a SNP-based marker panel useful for breed certification, parentage testing and individual identification purposes. A total of 17 markers, derived from a group of 49,034 SNPs genotyped with the Illumina SheepSNP50 Bead Chip (Illumina Inc., San Diego, CA) in three Brazilian breeds (Brazilian Creole, Morada Nova and Santa Ines) by the International Sheep Genome Consortium, were used to produce a GoldenGateTM VeraCodeTM assay. The criteria used for selecting markers were as follows: fixation confidence 20.9; genomic distance >3 Mbp; and no deviation from Hardy-Weinberg equilibrium. A total of 467 samples from six different breeds were tested (Creole, n=300; Bergamacia, n=24; Corridale, n=28; Pantaneira, n=50; Rabo Largo, n=20; Santa Ines, n=45) to validate the assay. Two SNPs did not produce consistent genotyping results in the tested platform and were excluded from further analysis. The remaining markers were used to perform an allocation test using the program Structure where five repetitions were performed using a total of 250k permutations each. Results indicate the marker panel is efficient in distinguishing five of the tested breeds. All samples were correctly assigned to the respective groups with the exception of the Pantaneira and Creole samples, which were grouped together. Other studies suggest these two groups are highly related and should probably be classified as two ecoptypes of the same breed. Therefore, the proposed reduced panel represents a useful tool for breed-certification of live animals and derived products.

Financial support: EMBRAPA, CNPq

Back to: Poster Abstract

<< Previous Abstract | Next Abstract >>

1 of 1 23-01-2012 09:33