



Candidate gene region for control of rib eye area in Canchim beef cattle

S.L. Meirelles¹, G.V. Gouveia¹, G. Gasparin², M.M. Alencar³,
J.J.S. Gouveia⁴ and L.C.A. Regitano³

¹Departamento de Genética e Evolução,
Universidade Federal de São Carlos, São Carlos, SP, Brasil

²Escola Superior de Agricultura “Luiz de Queiroz”, Universidade de São Paulo,
Piracicaba, SP, Brasil

³Embrapa Pecuária Sudeste, São Carlos, SP, Brasil

⁴Universidade Federal do Vale do São Francisco, Petrolina, PE, Brasil

Corresponding author: L.C.A. Regitano
E-mail: luciana@cnpse.embrapa.br

Genet. Mol. Res. 10 (2): 1220-1226 (2011)

Received November 26, 2010

Accepted March 3, 2011

Published June 21, 2011

DOI 10.4238/vol10-2gmr1175

ABSTRACT. Investigation of molecular marker effects on production traits is essential to define marker assisted selection strategies in beef cattle. We looked for a possible association of molecular markers and backfat thickness (BFT) and rib eye area (REA) in Canchim (5/8 Charolais + 3/8 Zebu) and MA (offspring of Charolais bulls and 1/2 Canchim + 1/2 Zebu cows) animals raised exclusively on pasture. Traits were measured on 987 individuals from seven herds from two Brazilian States (São Paulo and Goiás), in March and April from 2005 to 2007, when animals were, on average, 19 months of age. Five microsatellite markers lying in QTL regions for BFT and REA (BMS490 and ETH10 on chromosome 5, INRA133 and ILSTS090 on chromosome 6, and BMS2142 on chromosome 19) were genotyped and association analyses were performed under an animal model using the restricted

maximum likelihood method. After correction for multiple tests, a significant effect of microsatellite BMS490 on REA was observed, suggesting that at least one QTL affecting carcass traits in this region of the BTA5. No significant effect on BFT was observed for these markers.

Key words: Association study; Carcass traits; Cattle; Microsatellite markers