Niveus Ha412HOChr08 pos1 PET_39 2000 -IBD distance Inversion genotype — Different — Same 1000 -50 100 150 Ö MB

Niveus Ha412HOChr08 pos1 PET_39 IBD[genotype == 2] – IBD[genotype == 0] 3000 --400 normal inverted Genomic region IBD[genotype == different] - IBD[genotype == same] 2000 Inversion Genotype IBD distance 2 inverted 1000 normal Genomic region 1.00 0.75 ation counts sign 0_derived Populat 2_derived 0.25 -0 -0.00 -2_derived 0_derived Genomic region Signal

