

Petiolaris Ha412HOChr01 neg1 PET_12 PET_05 3000 IBD[genotype == 2] - IBD[genotype == 0] 2000 5000 8 4000 inverted normal Genomic region IBD[genotype == different] – IBD[genotype == same] 3000 Inversion Genotype IBD distance 2000 inverted normal Genomic region 1.00 -1000 -0.75 ation counts sign 0_derived Populat 2_derived 0.25 -0 -0.00 -0_derived 2_derived Genomic region

Signal

