

Petiolaris Ha412HOChr13 pos6 PET_17 PET_19 PET_16 6000 2000 IBD[genotype == 2] - IBD[genotype == 0] 4000 -2000 normal inverted 2000 Genomic region 1500 -IBD[genotype == different] - IBD[genotype == same] IBD distance Inversion Genotype PET_28 PET_24 inverted normal 4000 -Genomic region Population counts 2000 sign 0_derived 2_derived 0 -0_derived 2_derived Genomic region Signal

