

Petiolaris Ha412HOChr07 pos1 PET_04 GSD 1000 IBD[genotype == 2] - IBD[genotype == 0] • -1000 · -2000 · inverted normal Genomic region 4000 -IBD[genotype == different] – IBD[genotype == same] Inversion Genotype IBD distance 2000 inverted normal Genomic region 2.0 1.5 Population counts sign 0_derived 2_derived 0.5 0 -0.0 2_derived 0_derived Genomic region Signal

