

Petiolaris Ha412HOChr09 pos1 PET_18 PET_21 PET_17 6000 IBD[genotype == 2] – IBD[genotype == 0] inverted normal Genomic region 4000 2000 IBD[genotype == different] – IBD[genotype == same] Inversion Genotype IBD distance 2000 inverted normal Genomic region Population counts sign 0_derived 2_derived 2_derived 0_derived Genomic region Signal

