

Petiolaris Ha412HOChr05 neg1 GSD PET_23 2000 6000 IBD[genotype == 2] - IBD[genotype == 0] 1000 4000 -3000 inverted normal 2000 Genomic region 2000 IBD[genotype == different] – IBD[genotype == same] IBD distance Inversion Genotype PET_24 PET_30 2 inverted normal 4000 Genomic region Population counts 2000 sign 0_derived 2_derived 0 -0 -0_derived 2_derived Genomic region Signal

