

Petiolaris Ha412HOChr09 neg1 GSD PET_17 6000 • IBD[genotype == 2] – IBD[genotype == 0] 1000 4000 -2000 inverted normal 2000 Genomic region IBD[genotype == different] – IBD[genotype == same] IBD distance Inversion Genotype PET_18 PET_21 4000 inverted normal Genomic region Population counts 2000 sign 0_derived 2_derived 0 -2_derived 0_derived Genomic region Signal

