Petiolaris Ha412HOChr05 neg1 GSD 3000 -2000 -IBD distance Inversion genotype Different — Same 1000 50 150 100 Ö MB

Petiolaris Ha412HOChr05 neg1 GSD IBD[genotype == 2] – IBD[genotype == 0] normal inverted Genomic region IBD[genotype == different] - IBD[genotype == same] 2000 Inversion Genotype IBD distance inverted normal Genomic region 1000 1.00 -0.75 ation counts sign 0\_derived Populat 2\_derived 0.25 -0.00 -0\_derived 2\_derived Genomic region Signal

