Petiolaris Ha412HOChr09 neg1 IBD distance Inversion genotype Different — Same 1000 100 MB 50 150 200 Ö

Petiolaris Ha412HOChr09 neg1 GSD 2000 IBD[genotype == 2] – IBD[genotype == 0] 4000 3000 normal inverted Genomic region IBD[genotype == different] – IBD[genotype == same] IBD distance Inversion Genotype inverted normal Genomic region 1.00 -1000 0.75 ation counts sign 0\_derived Populat 2\_derived 0.25 -0.00 -0\_derived 2\_derived Genomic region Signal

