Petiolaris Ha412HOChr17 pos3 3000 2000 -Inversion genotype IBD distance Different — Same 100 MB 50 200 150 Ö

Petiolaris Ha412HOChr17 pos3 GSD 1000 IBD[genotype == 2] – IBD[genotype == 0] 4000 3000 -2000 · 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 -0.00 -2_derived 0_derived Genomic region Signal

