

Petiolaris Ha412HOChr08 pos2 PET\_10 GSD 2000 6000 IBD[genotype == 2] – IBD[genotype == 0] 8 inverted normal Genomic region 4000 2000 IBD[genotype == different] - IBD[genotype == same] 1000 IBD distance Inversion Genotype -1000 · 2000 inverted normal Genomic region 1.00 -0.75 ation counts sign 0\_derived Populat 2\_derived 0.25 -0 -0.00 -0\_derived 2\_derived Genomic region Signal

