Petiolaris Ha412HOChr16 pos2 PET\_14 2000 -IBD distance Inversion genotype — Different — Same 1000 100 150 200 MB

Petiolaris Ha412HOChr16 pos2 PET\_14 1000 -IBD[genotype == 2] – IBD[genotype == 0] 4000 inverted normal 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 -0\_derived 2\_derived

Signal

Genomic region

