

Petiolaris Ha412HOChr11 pos1 Mon GSD IBD[genotype == 2] – IBD[genotype == 0] 1000 -4000 -2000 inverted normal Genomic region IBD[genotype == different] – IBD[genotype == same] IBD distance Inversion Genotype 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

