

Petiolaris Ha412HOChr06 neg2 PET_03 Mon PET_02 6000 IBD[genotype == 2] - IBD[genotype == 0] 4000 2000 inverted normal Genomic region PET_04 PET_08 PET_11 6000 IBD[genotype == different] - IBD[genotype == same] 2000 4000 -IBD distance Inversion Genotype 1000 -2000 inverted normal PET_43 Genomic region 6000 4000 -Population counts sign 0_derived 2_derived 2000 -0_derived 2_derived

Genomic region

Signal

