Niveus Ha412HOChr06 neg3 PET\_40 2000 -IBD distance Inversion genotype — Different — Same 1000

100

MB

150

50

Ö

Niveus Ha412HOChr06 neg3 PET\_40 500 IBD[genotype == 2] – IBD[genotype == 0] 3000 –1000 · -1500 · inverted normal Genomic region IBD[genotype == different] - IBD[genotype == same] 2000 Inversion Genotype IBD distance inverted normal Genomic region 1000 1.00 0.75 tion counts sign 0\_derived Populat 2\_derived 0.25 -0.00 -2\_derived 0\_derived Genomic region

Signal

