Argophyllus Ha412HOChr10 neg1 ARG\_20 4000 -3000 IBD distance Inversion genotype Different — Same 1000 50 100 150 200 Ö

MB

Argophyllus Ha412HOChr10 neg1 ARG\_20 IBD[genotype == 2] – IBD[genotype == 0] 3000 normal inverted Genomic region IBD[genotype == different] - IBD[genotype == same] 2000 Inversion Genotype IBD distance 2 inverted normal Genomic region 1000 1.00 0.75 ation counts sign 0\_derived Populat 2\_derived 0.25 -0 -0.00 -2\_derived 0\_derived Genomic region Signal

