

Argophyllus Ha412HOChr03 pos1 ARG\_21 ARG\_07 IBD[genotype == 2] – IBD[genotype == 0] 2000 -4000 inverted normal Genomic region 5000 IBD[genotype == different] - IBD[genotype == same] 3000 3000 IBD distance Inversion Genotype 2000 -1000 2000 -1000 normal inverted Genomic region 1.00 -1000 -0.75 ttion counts sign 0\_derived Populat 2\_derived 0.25 -0.00 -0\_derived 2\_derived

Signal

Genomic region

