

Argophyllus Ha412HOChr16 pos3 ARG_15 ARG_13 5000 IBD[genotype == 2] – IBD[genotype == 0] 2000 4000 -3000 -2000 2000 -4000 inverted normal Genomic region 4000 IBD[genotype == different] - IBD[genotype == same] 1000 3000 2000 IBD distance Inversion Genotype ARG_20 ARG_18 1000 2 5000 4000 inverted normal Genomic region 3000 2000 -Population counts sign 0_derived 2_derived 1000 0 -2_derived 0_derived Signal Genomic region

