

Peak: Chromosome_3_RagTag:36680000–36730000

Model: Overlapping models

Overlapping models: Selection–bottleneck + Introgression + Deep lineage sorting

Overlap signals: Belem_enrichment (high); Belem_RTH (low); Xingu_Belem_RTH (high); Tapajos_Xingu_JCR (high)

