

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)

