

Peak: Chromosome_10_RagTag:18250000–18270000

Model: Overlapping models

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

Overlap signals: Xingu_RTH (low); Xingu_Belem_RTH (high); Tapajos_Xingu_RTH (low); Tapajos_Xingu_JCR (high)

