

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)

