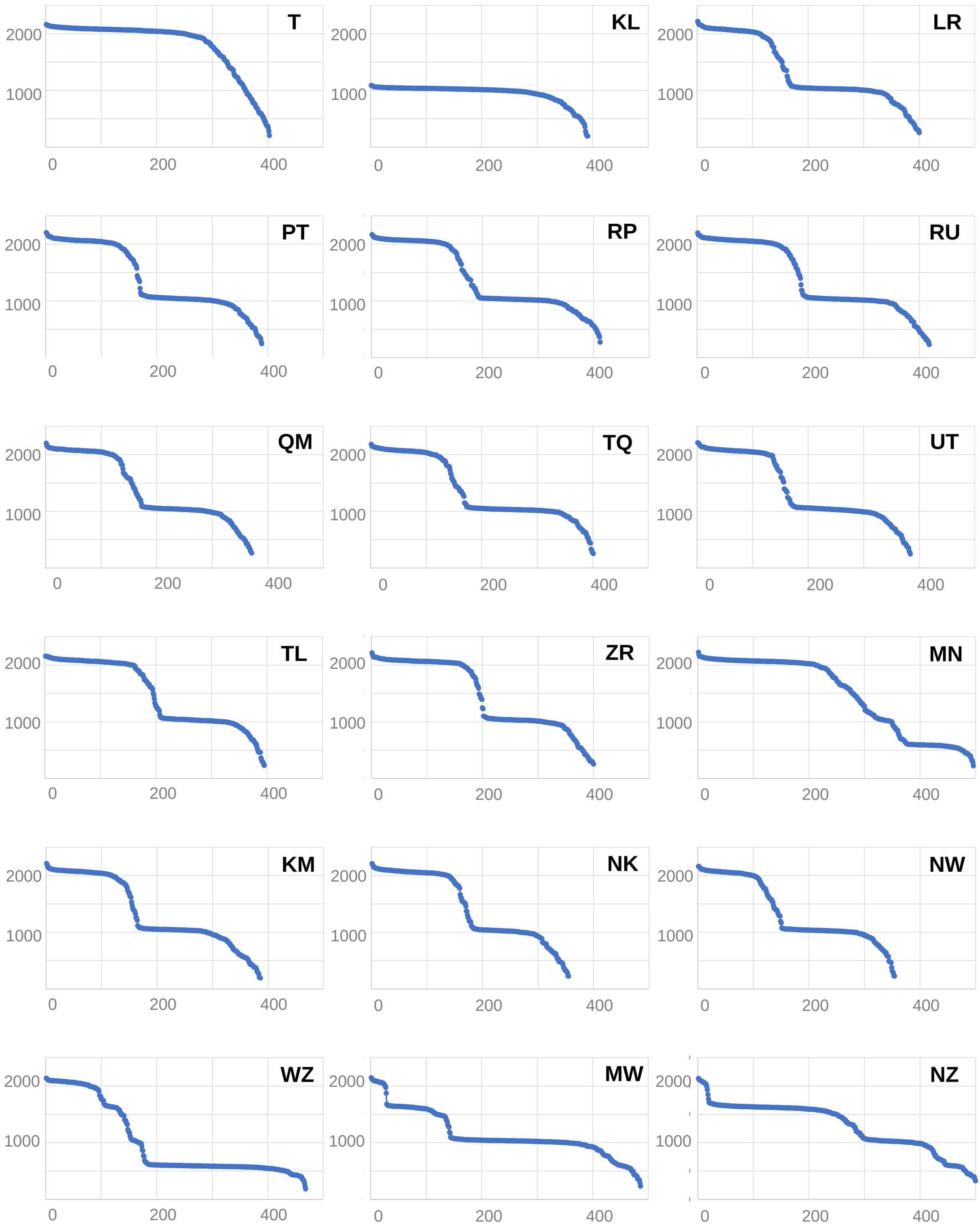
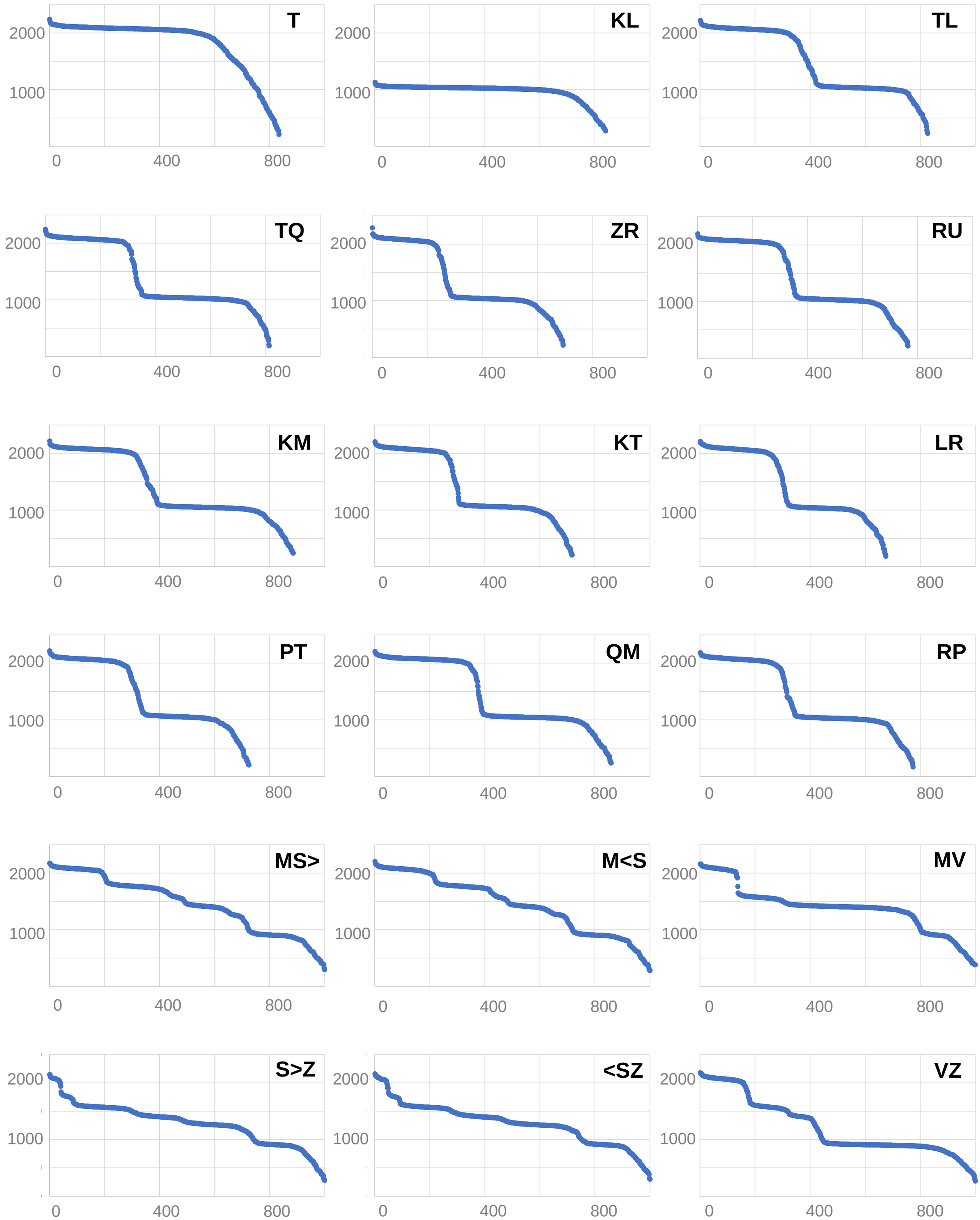
Quantification of *L.sativa* mitochondrial genome junctions



X axis - PacBio reads aligned to 2000 bp junction query and sorted by alignment length
Y axis - alignment length to 2000 bp junction query

S3A Figure

Quantification of *L.saligna* mitochondrial genome junctions



X axis - PacBio reads aligned to 2000 bp junction query and sorted by alignment length
Y axis - alignment length to 2000 bp junction query

S3B Figure