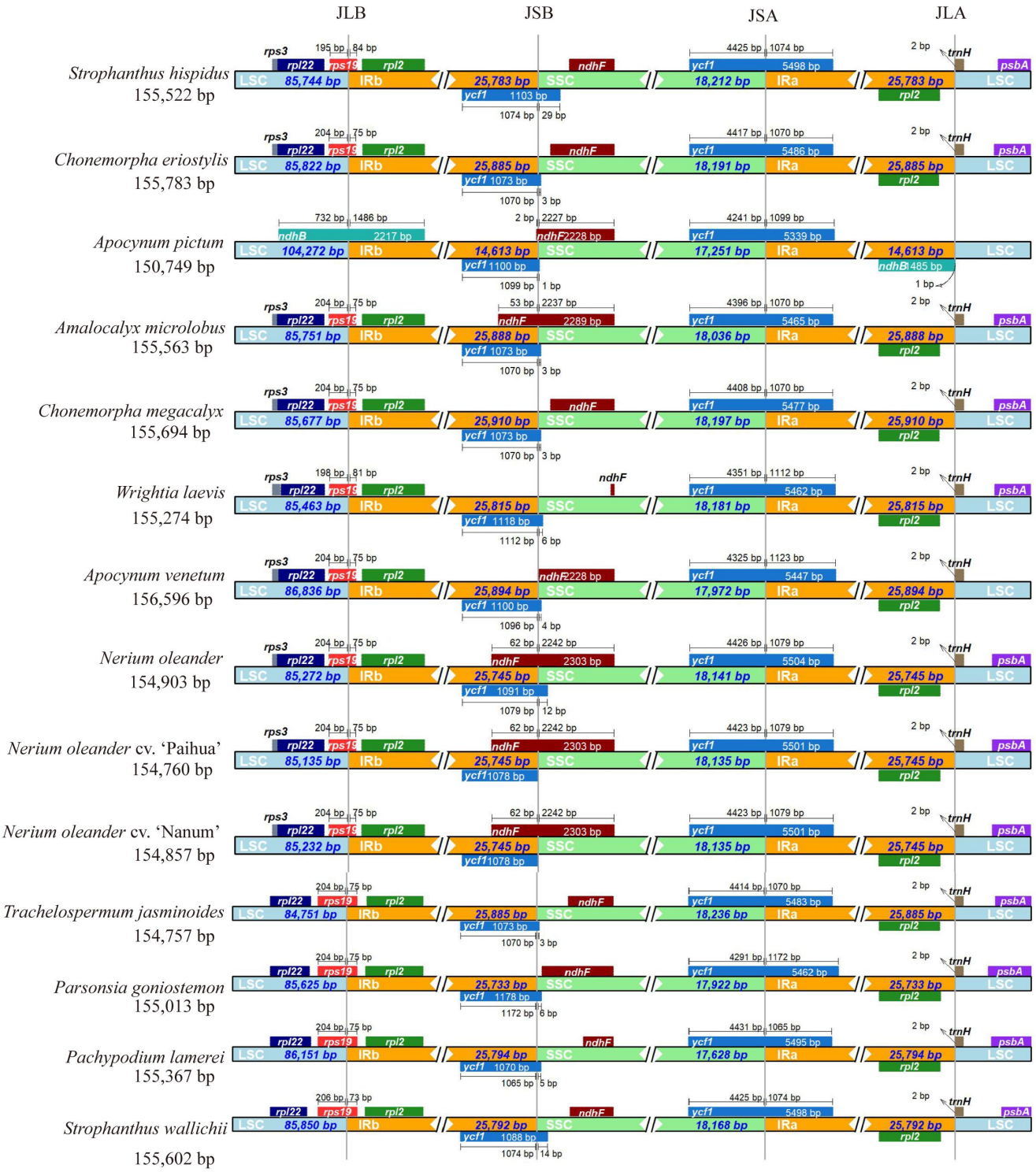

Figure S3. Comparison of LSC, SSC, and IR boundary regions among 14 Apocynoideae chloroplast genomes. Genes are represented by colored boxes. Numbers above gene features indicate the distance between gene ends and boundary sites.