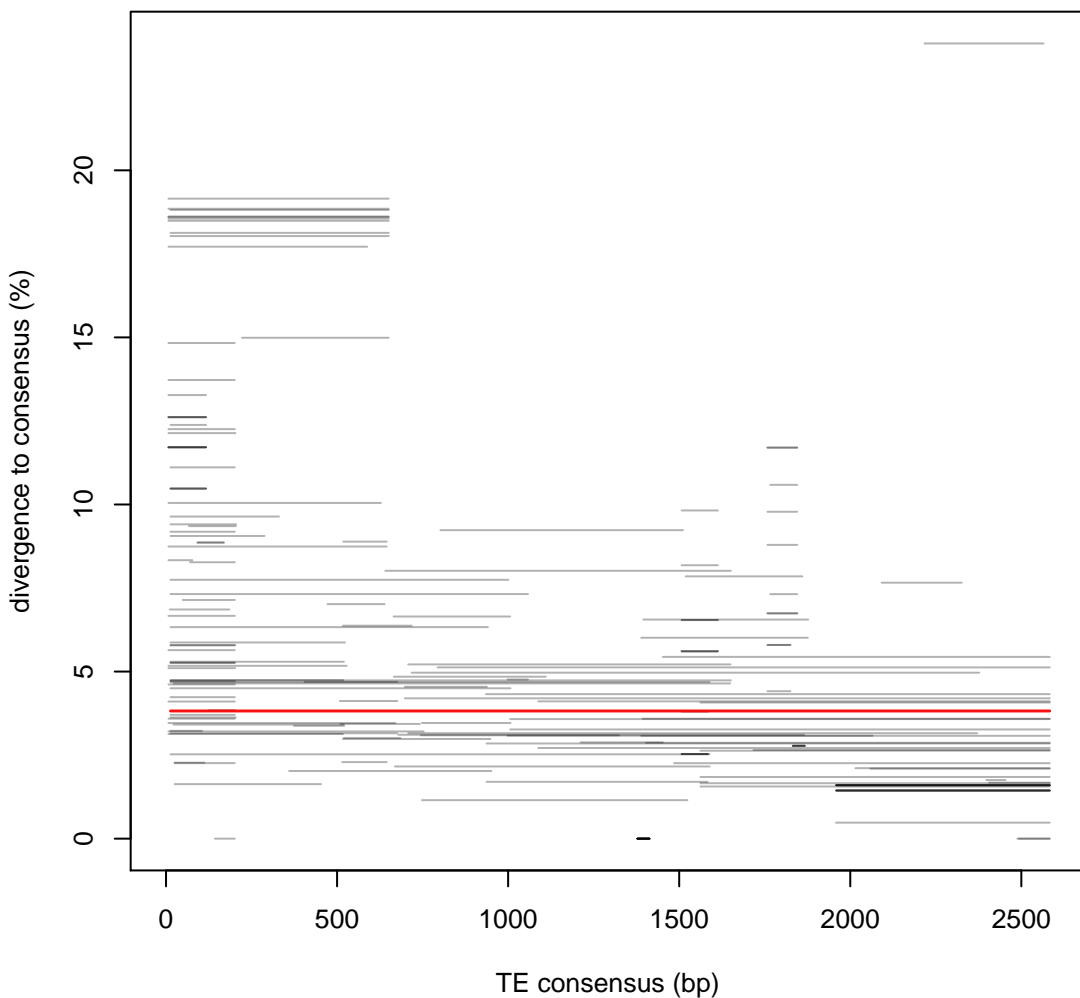
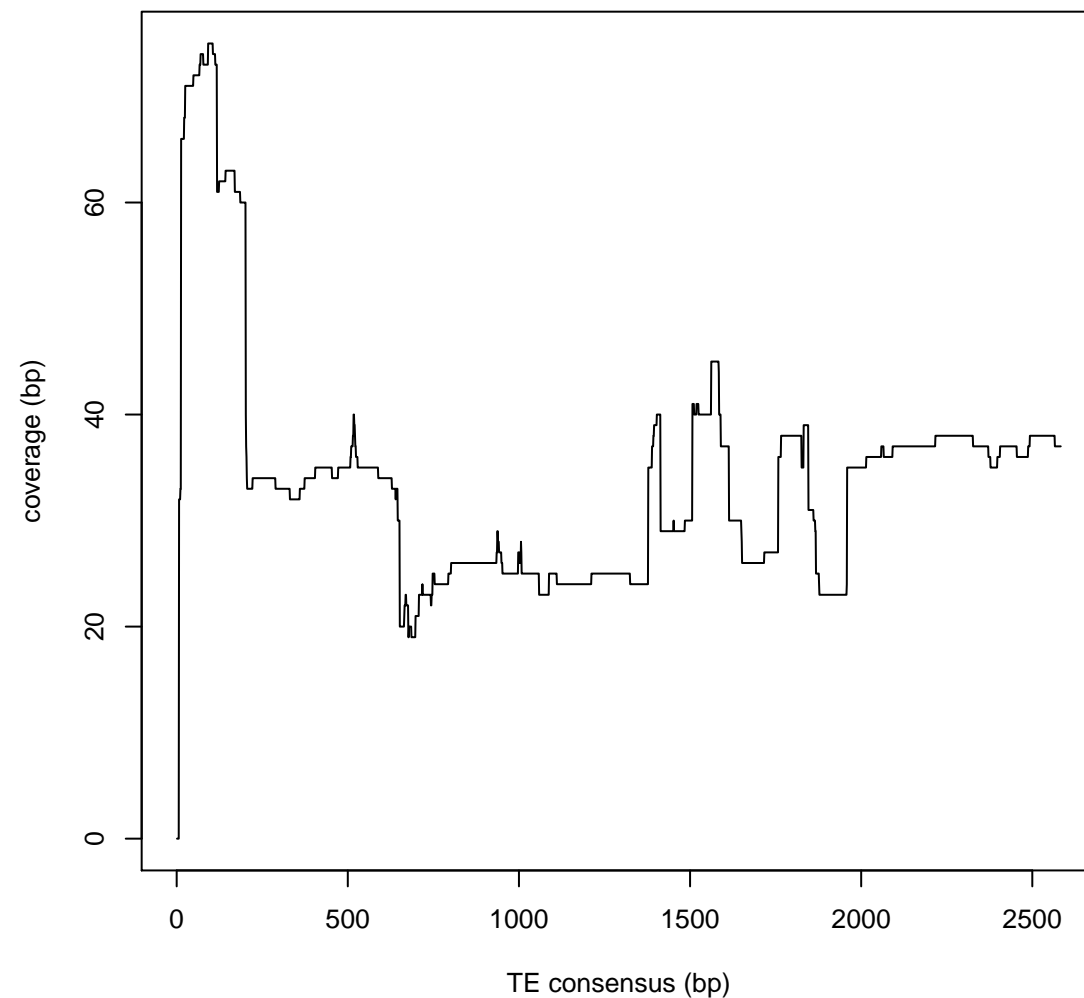


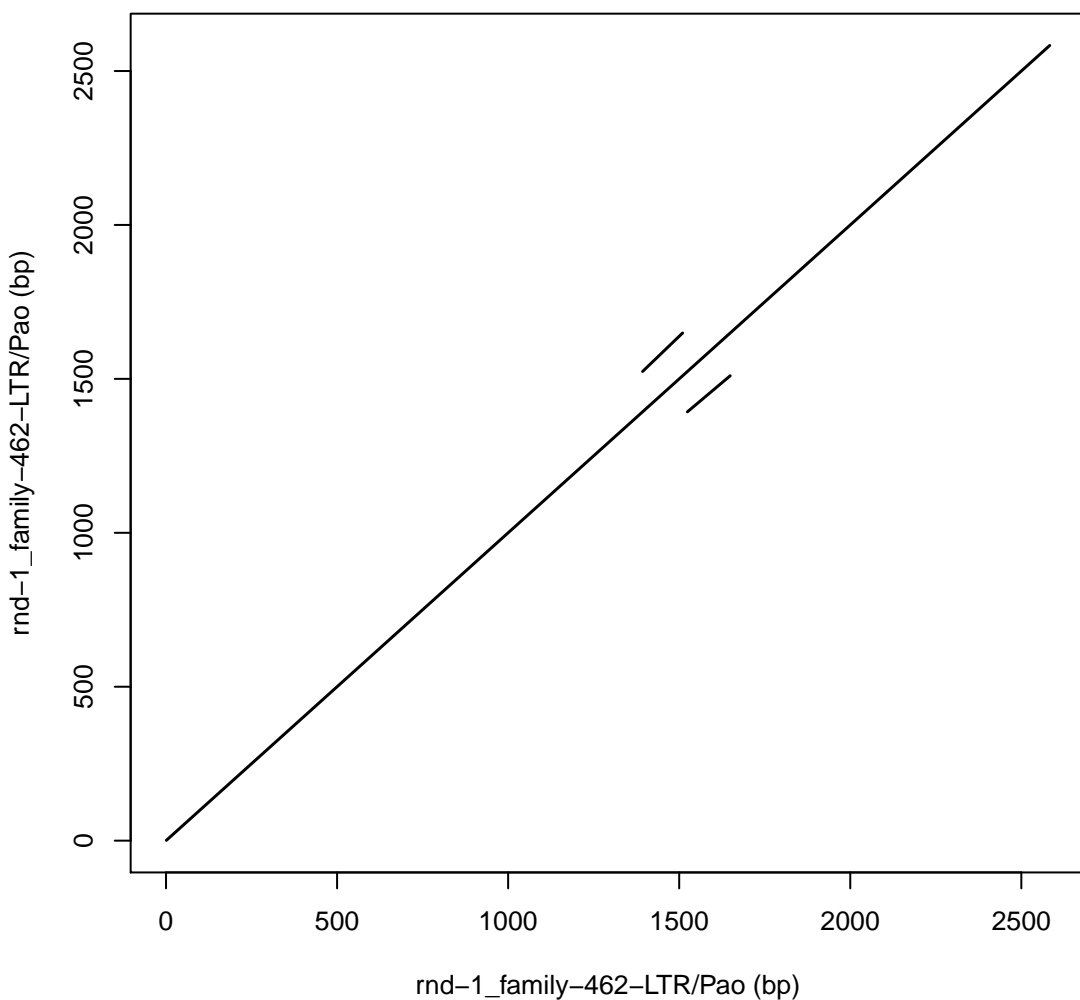
TE: rnd-1_family-462-LTR/Pao
consensus size: 2583bp; fragments: 186; full length: 1 (>=2324.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

