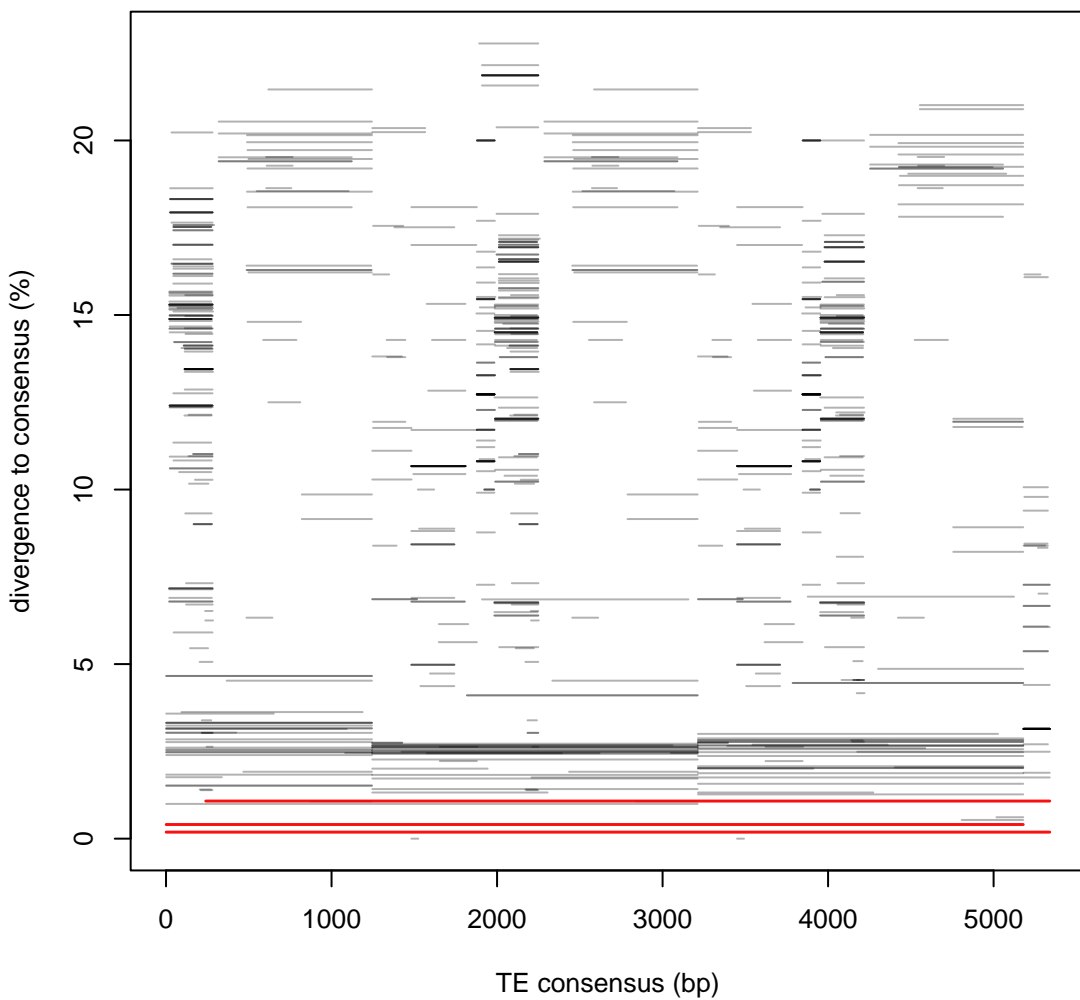
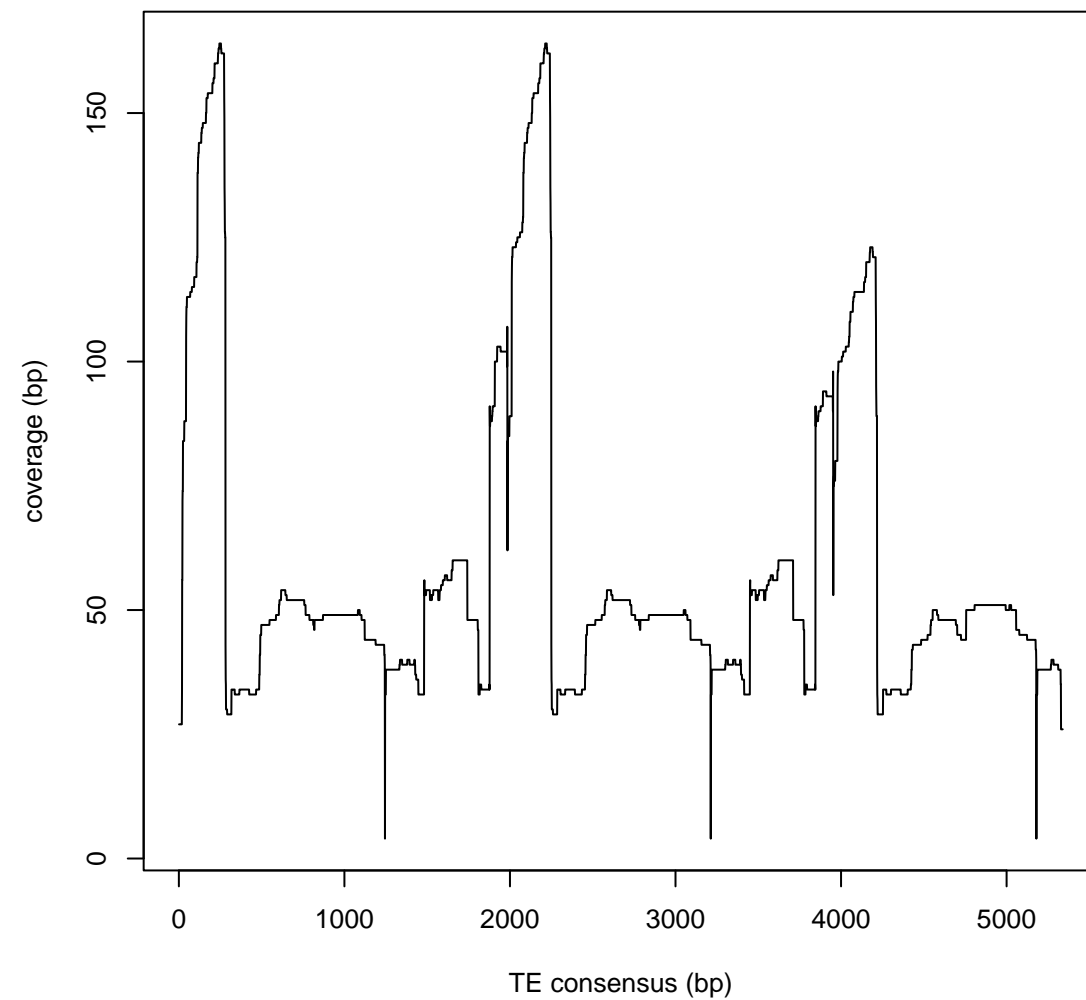


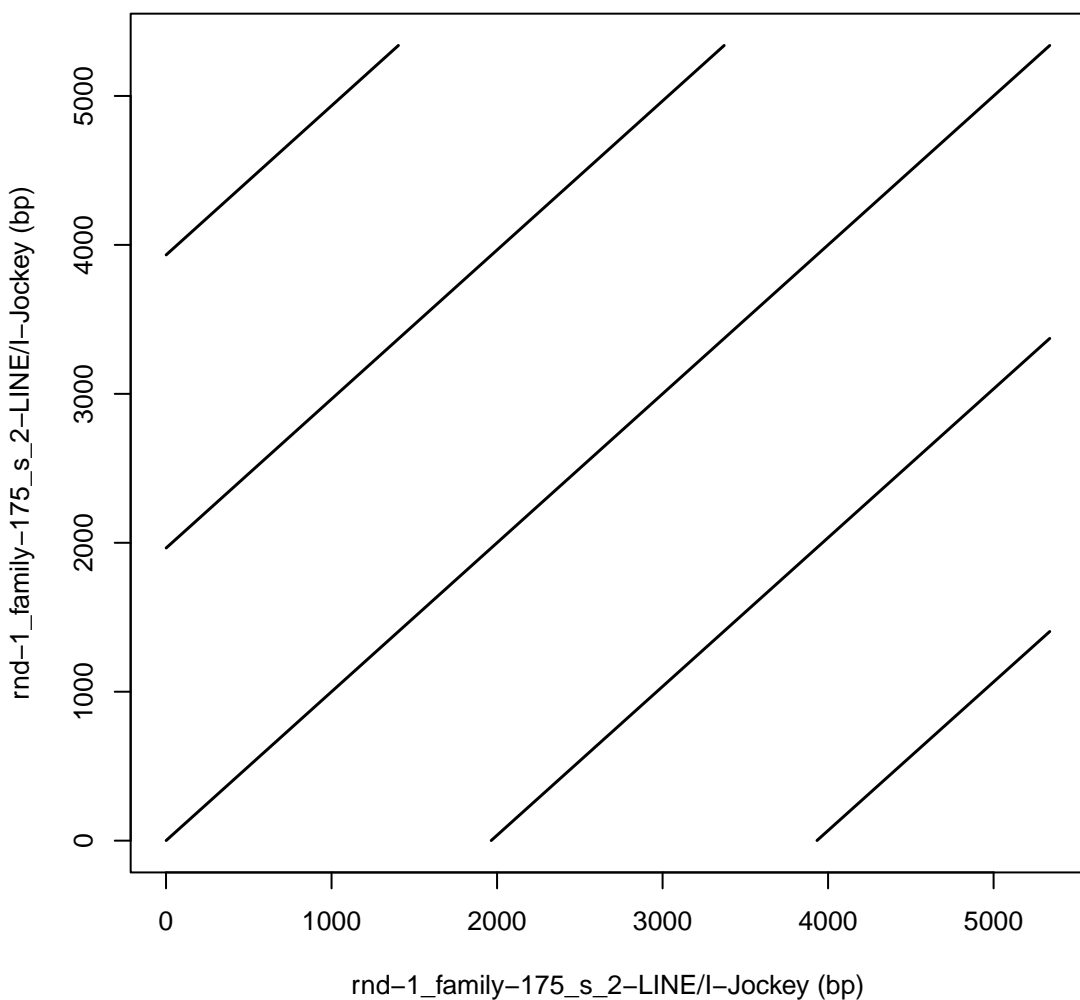
TE: rnd-1_family-175_s_2-LINE/I-Jockey
consensus size: 5339bp; fragments: 808; full length: 3 (>=4805.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

