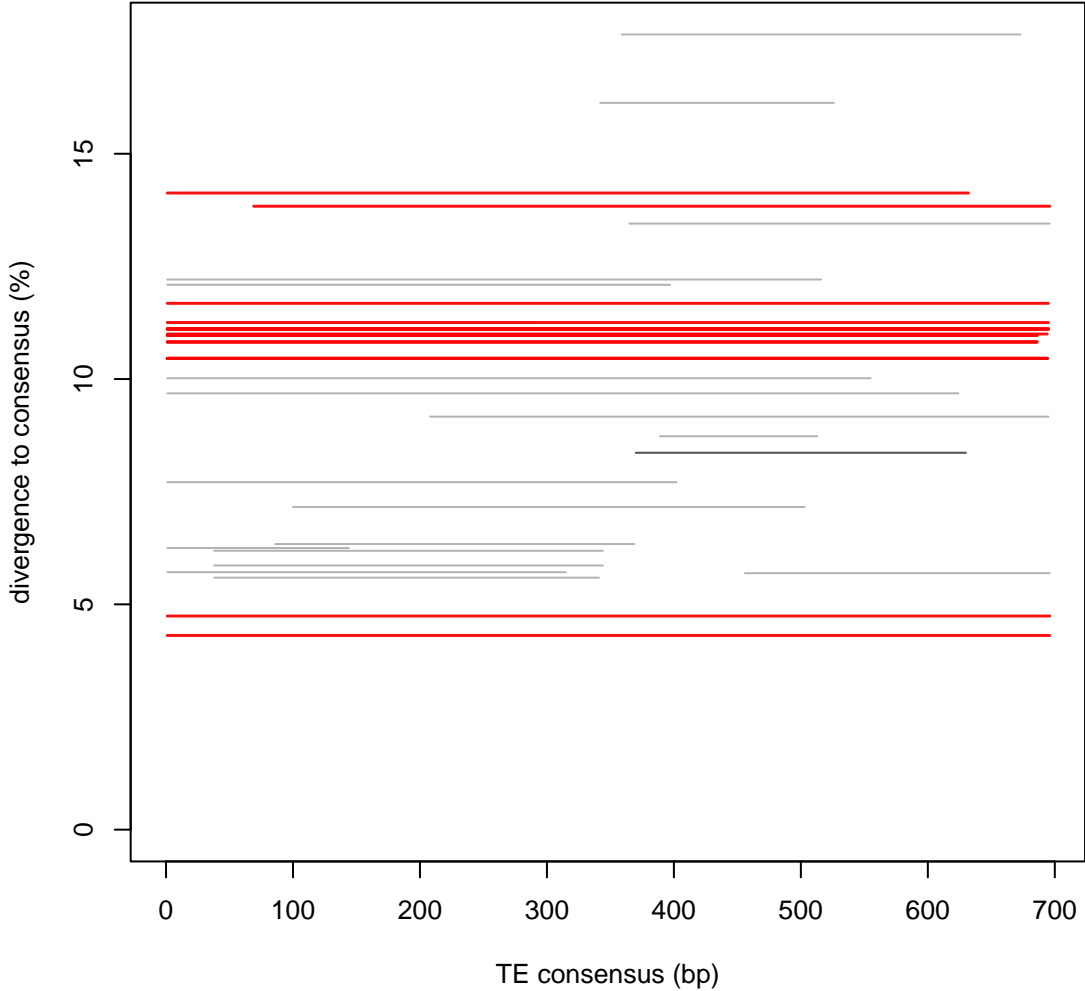
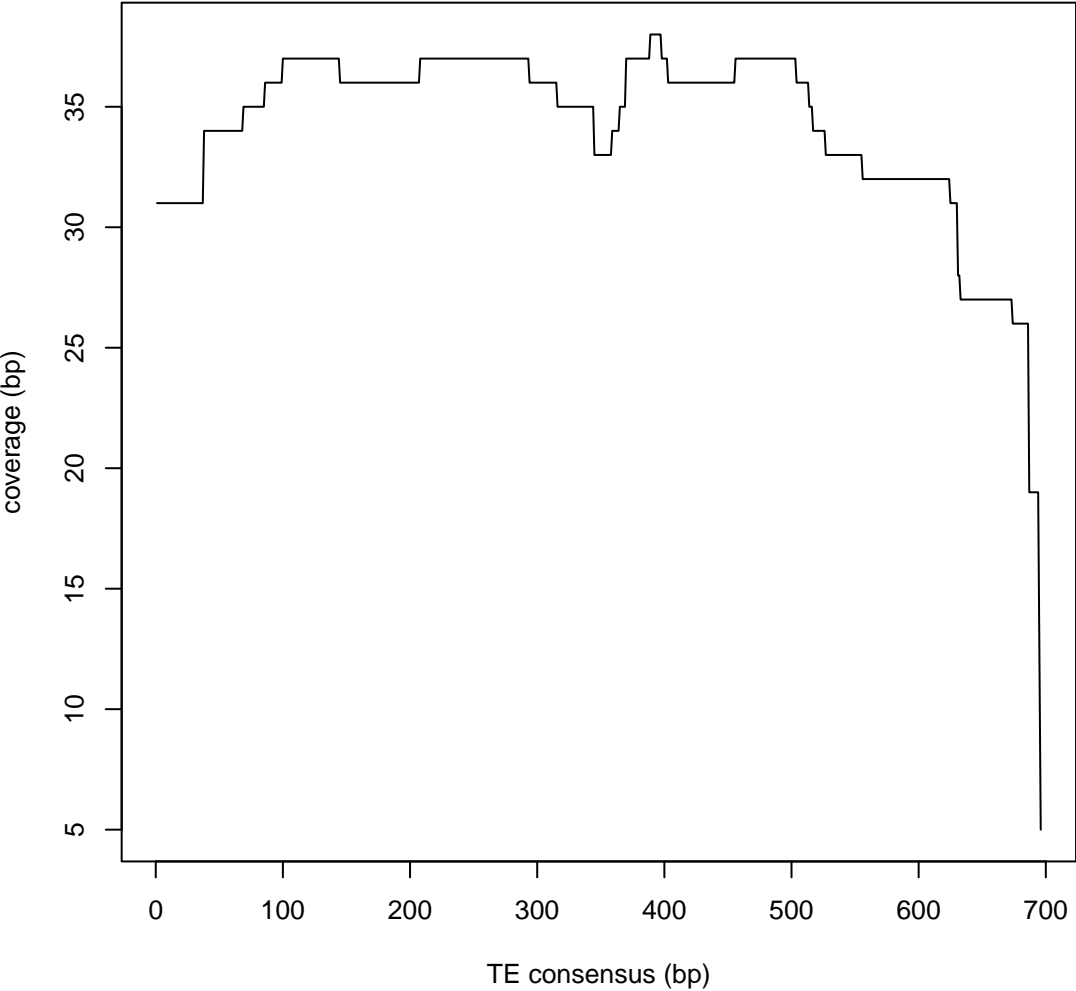


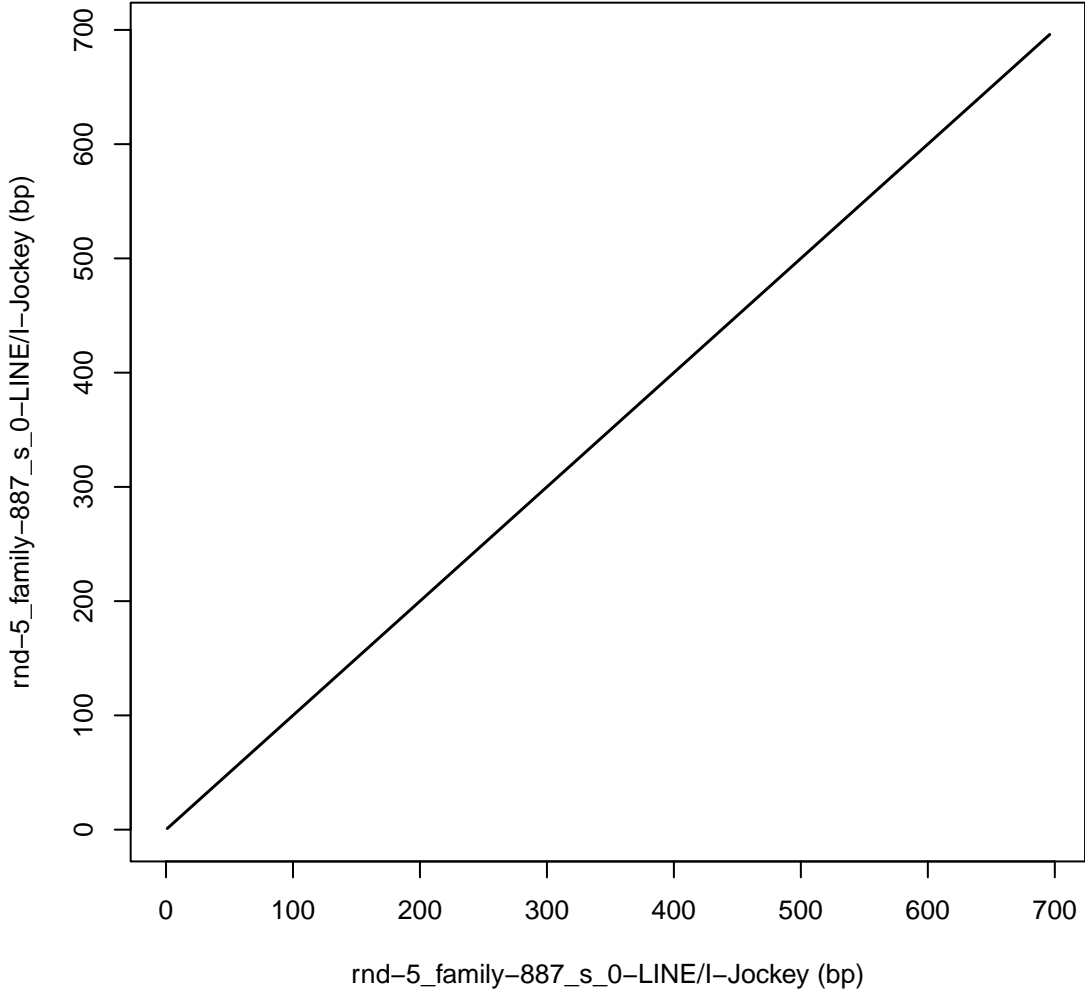
TE: rnd-5_family-887_s_0-LINE/I-Jockey
consensus size: 696bp; fragments: 46; full length: 24 (>=626.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

