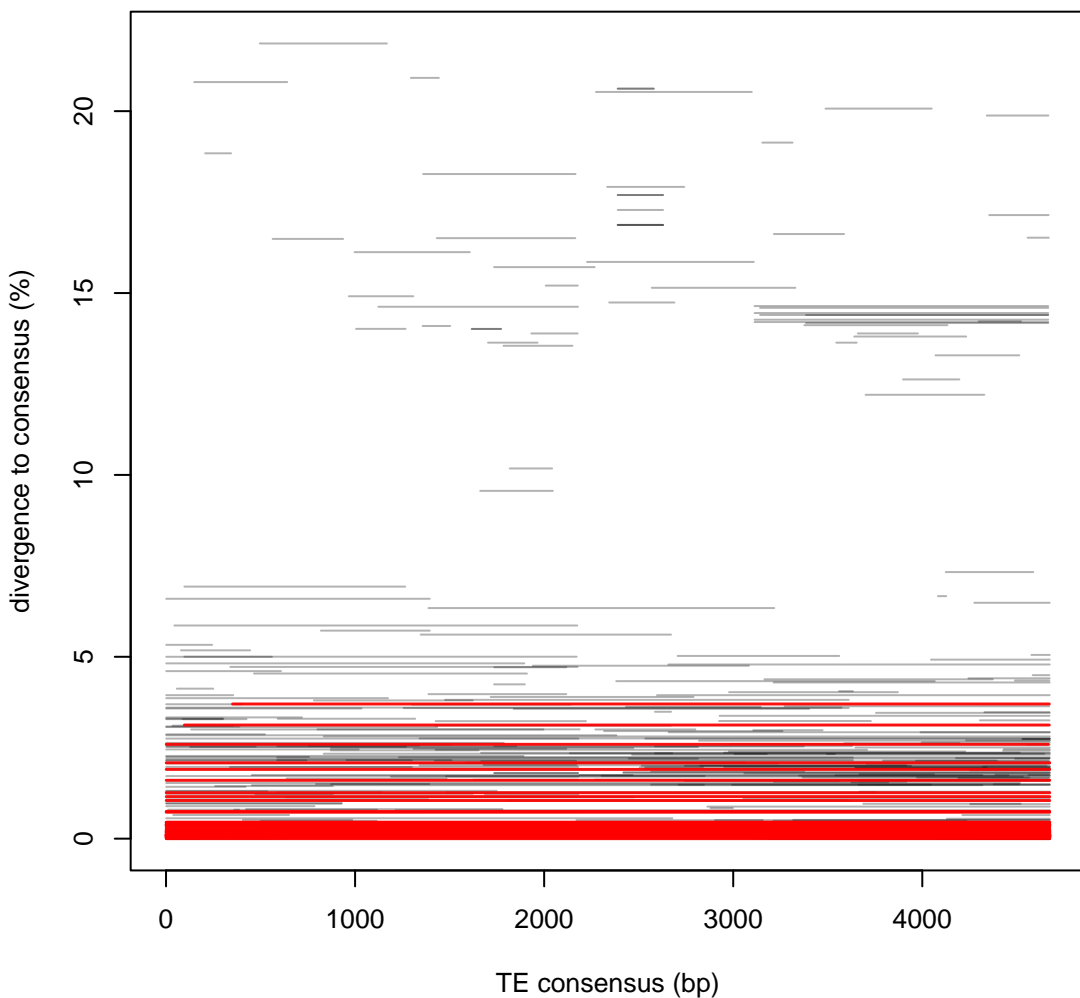
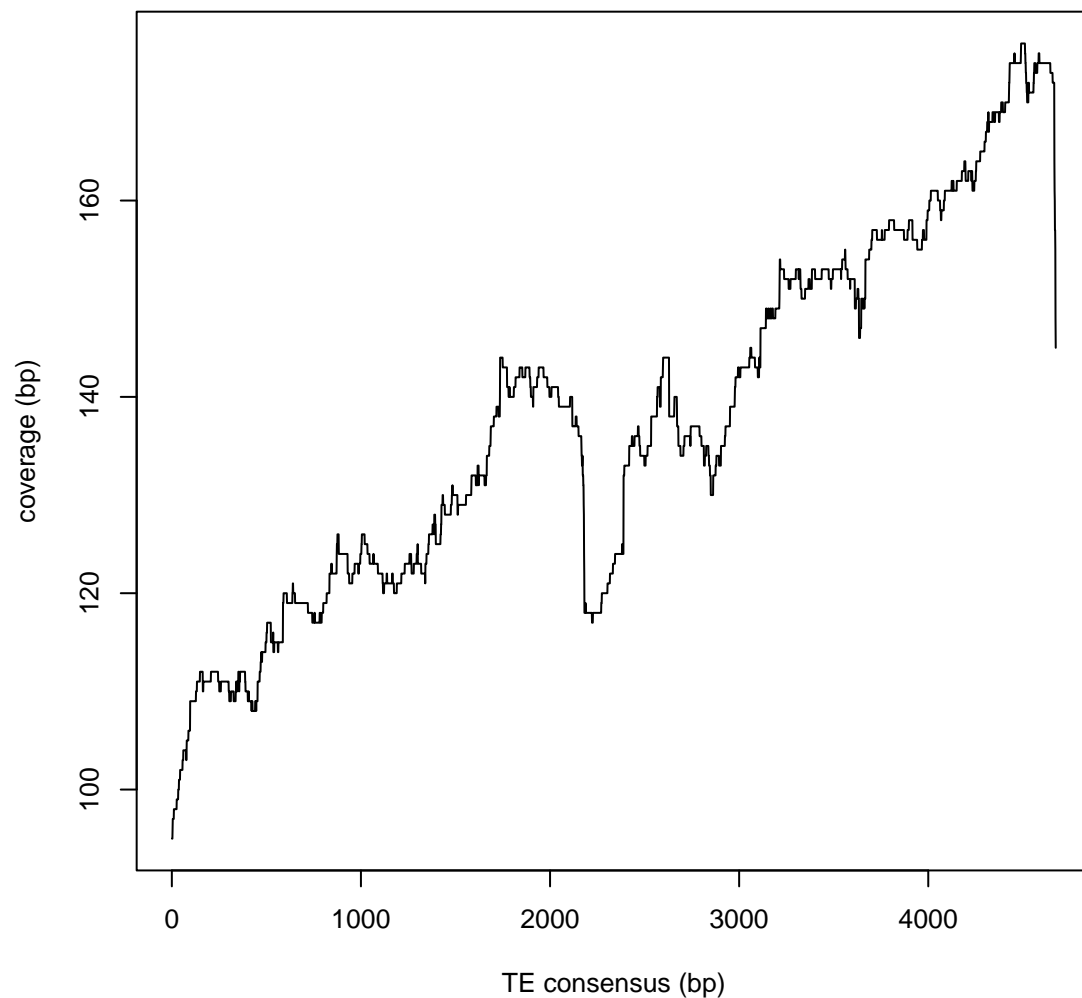


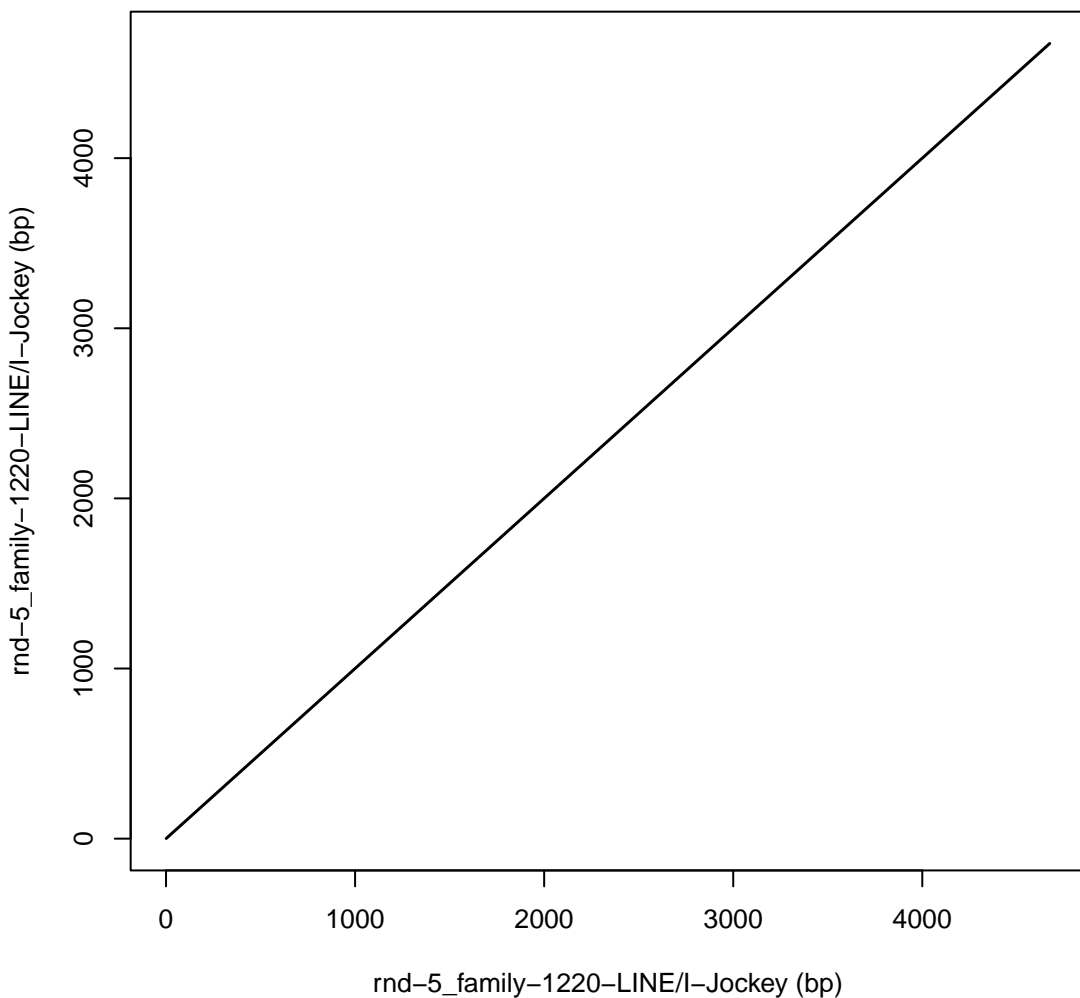
TE: rnd-5_family-1220-LINE/I-Jockey
consensus size: 4674bp; fragments: 392; full length: 51 (>=4206.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

