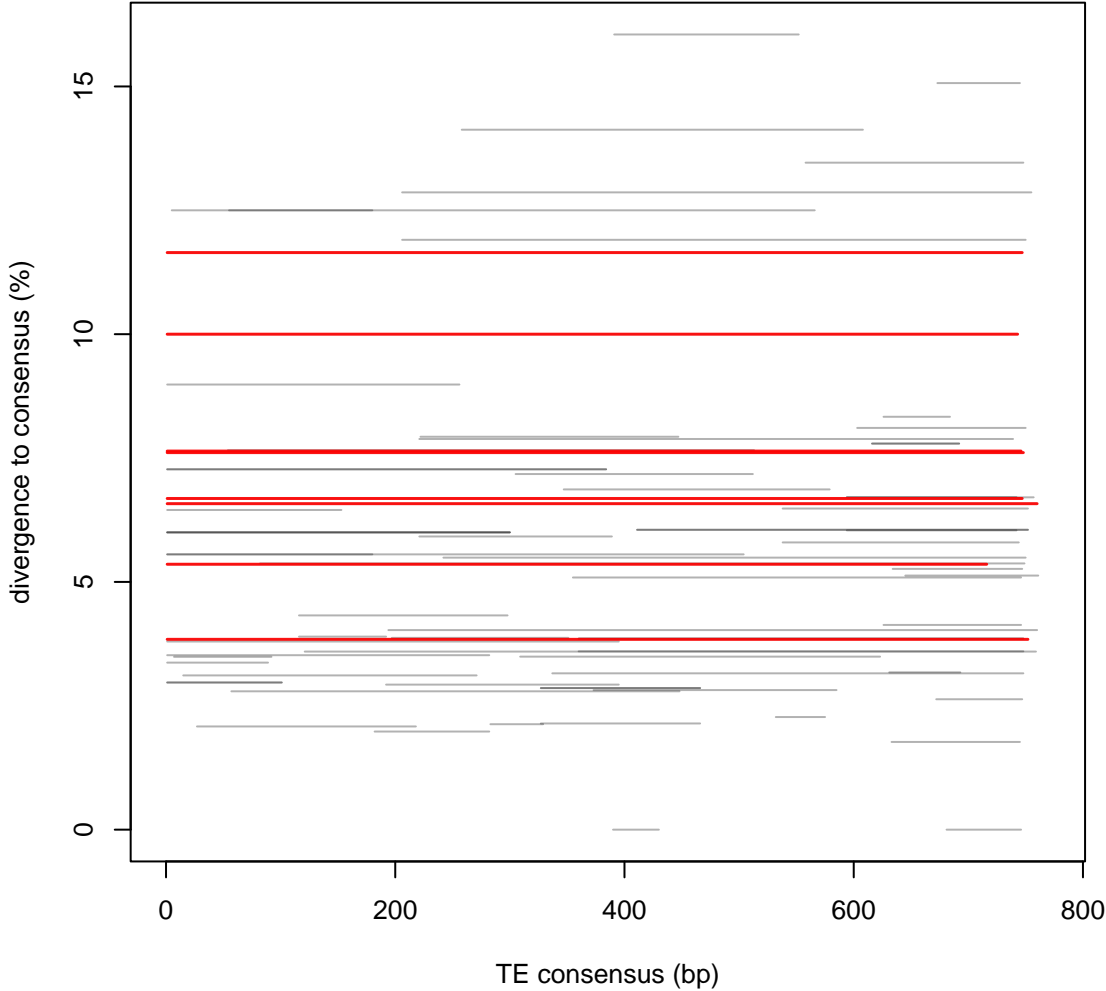
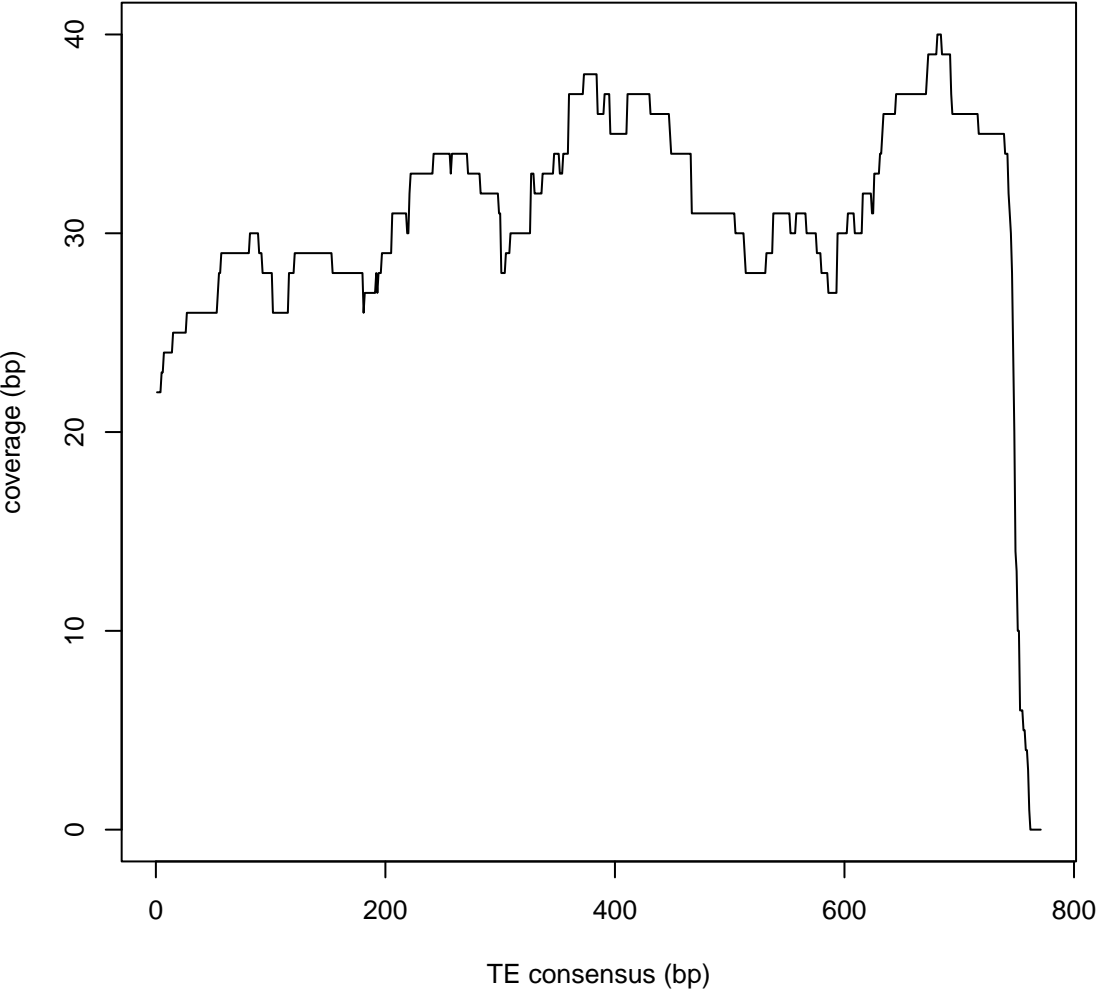


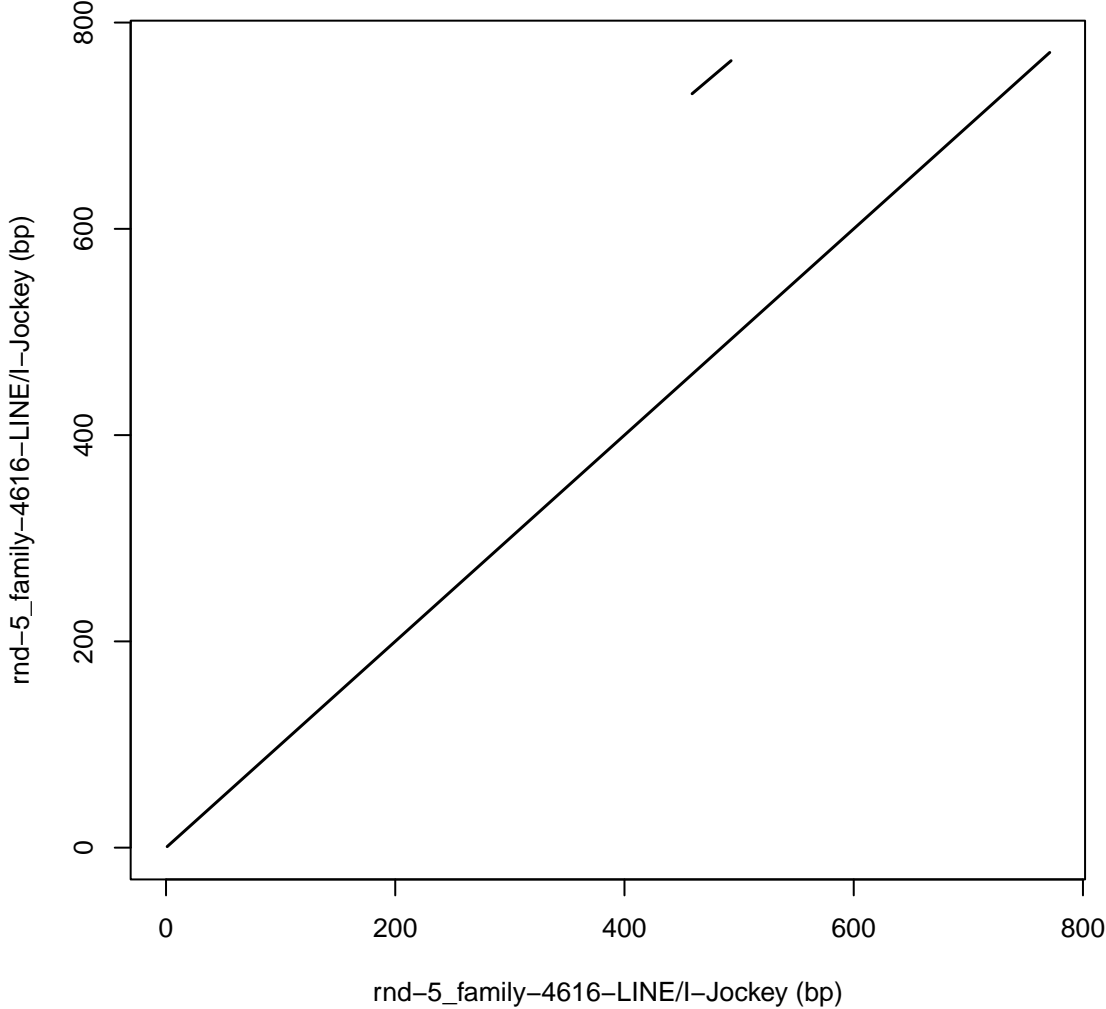
TE: rnd-5_family-4616-LINE/l-Jockey
consensus size: 771bp; fragments: 80; full length: 8 (>=693.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

