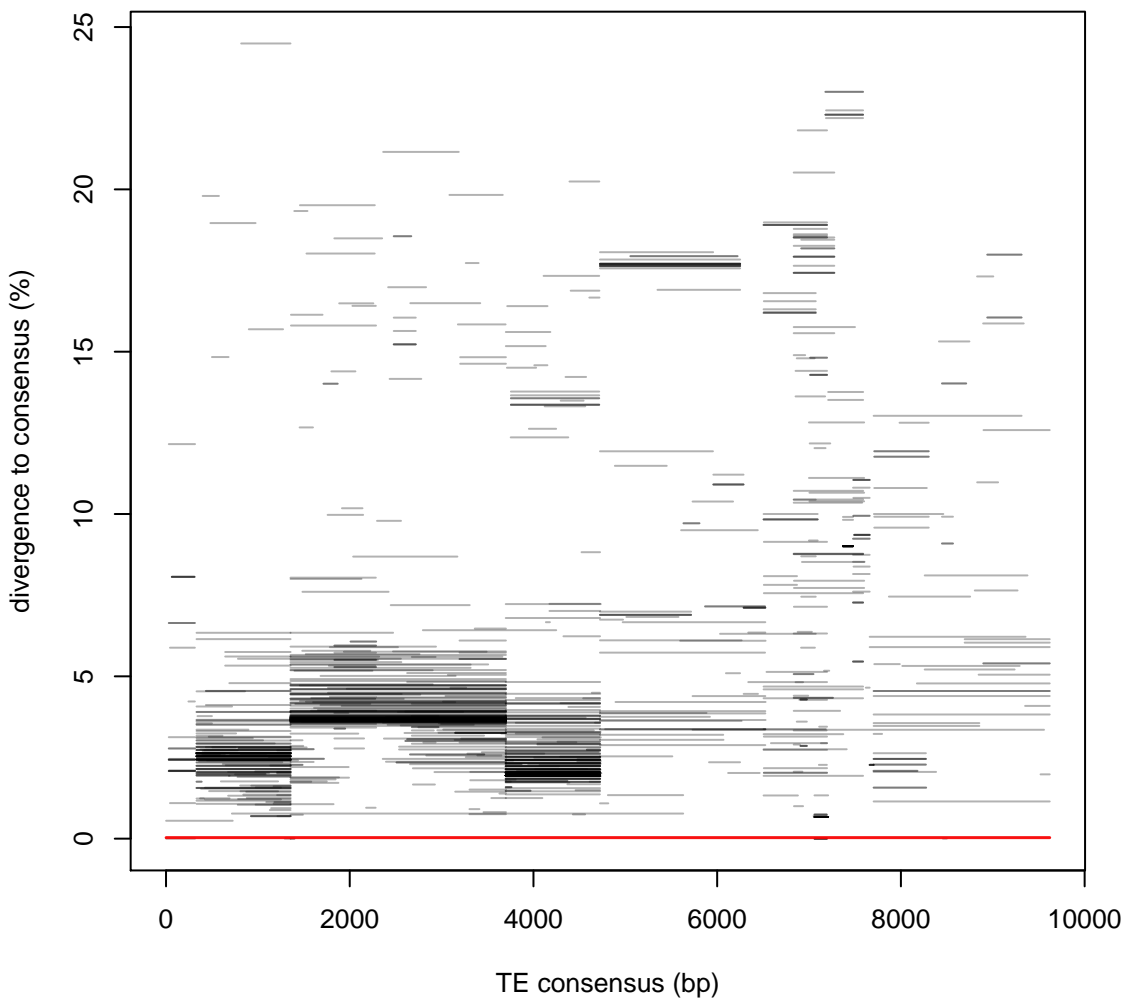
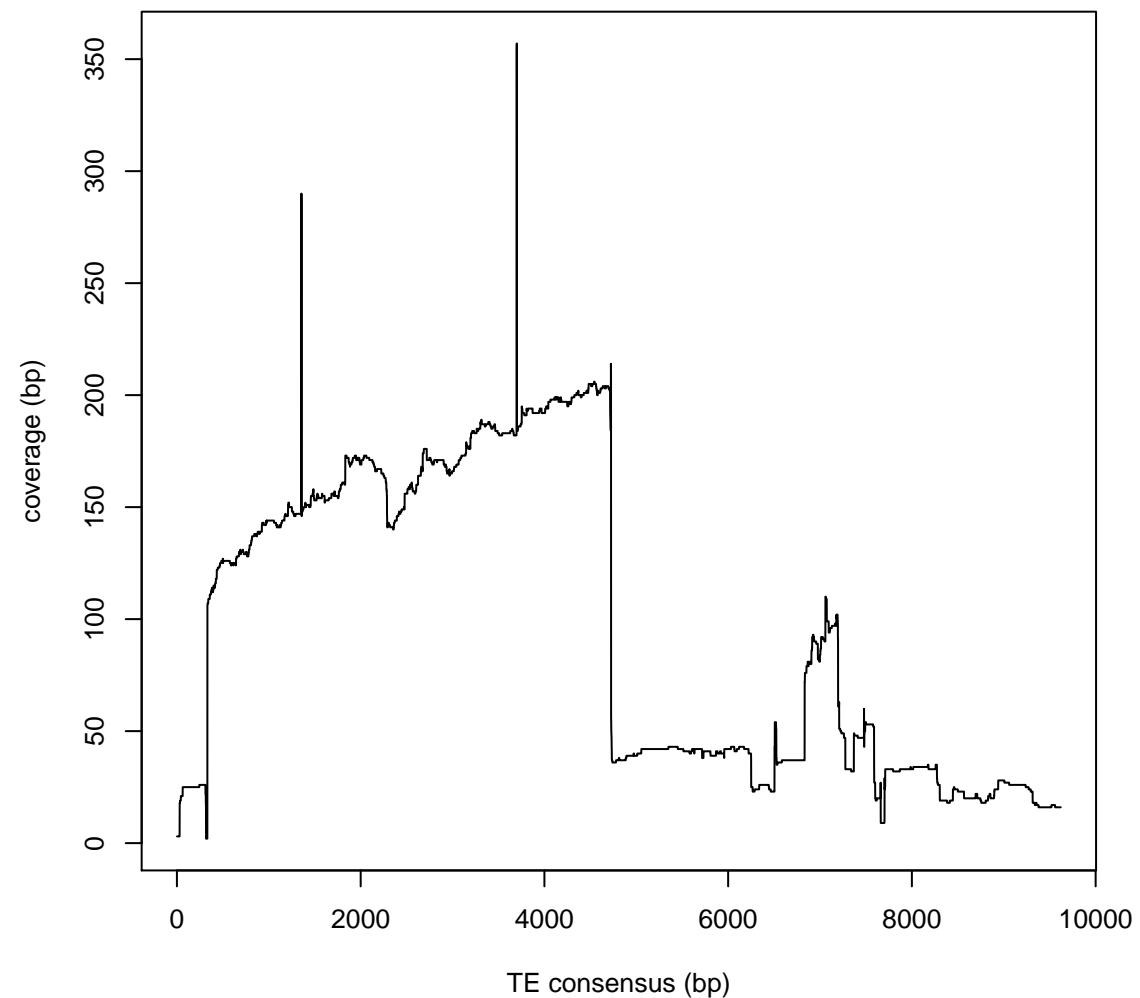


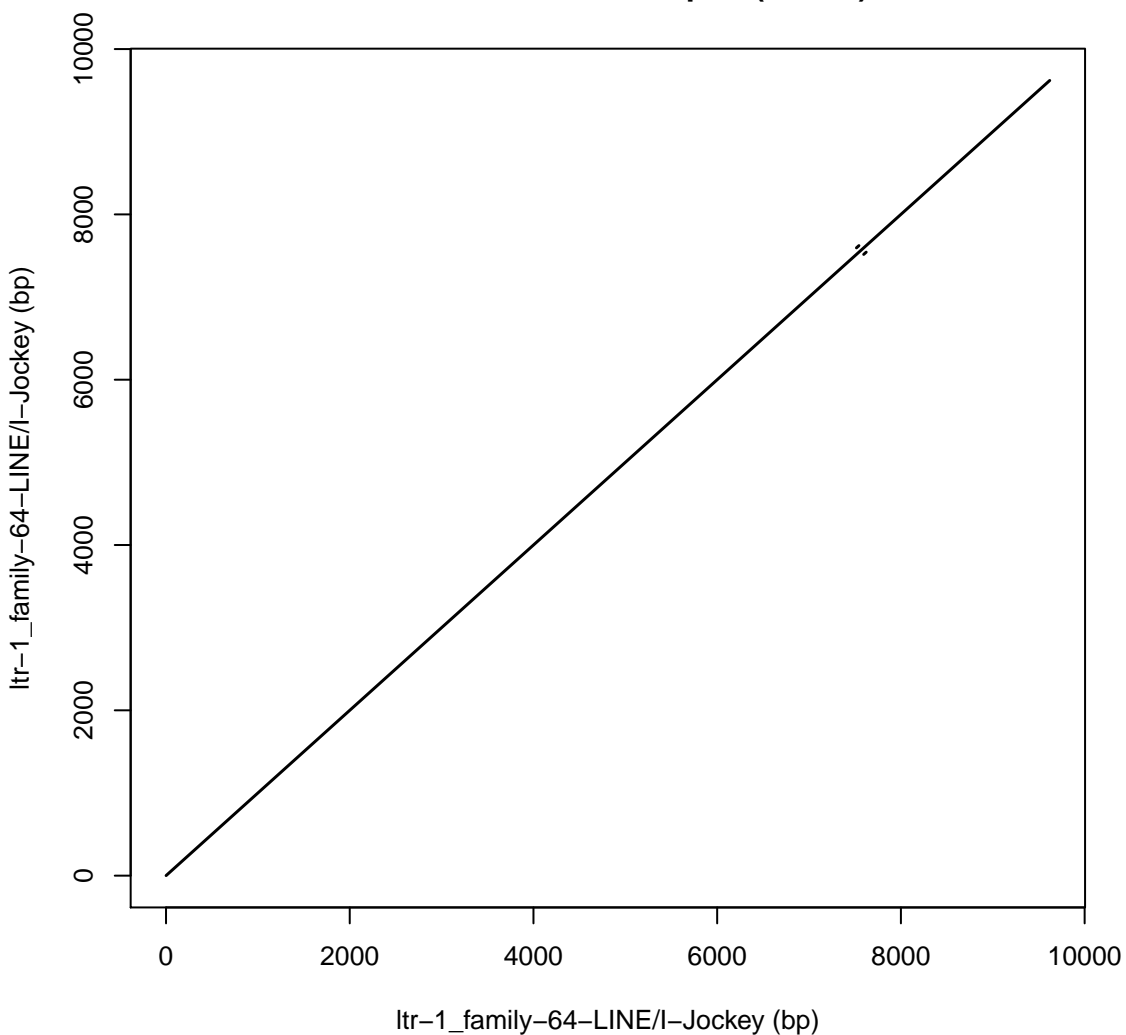
TE: Itr-1_family-64-LINE/I-Jockey
consensus size: 9620bp; fragments: 1044; full length: 1 (>=8658bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

