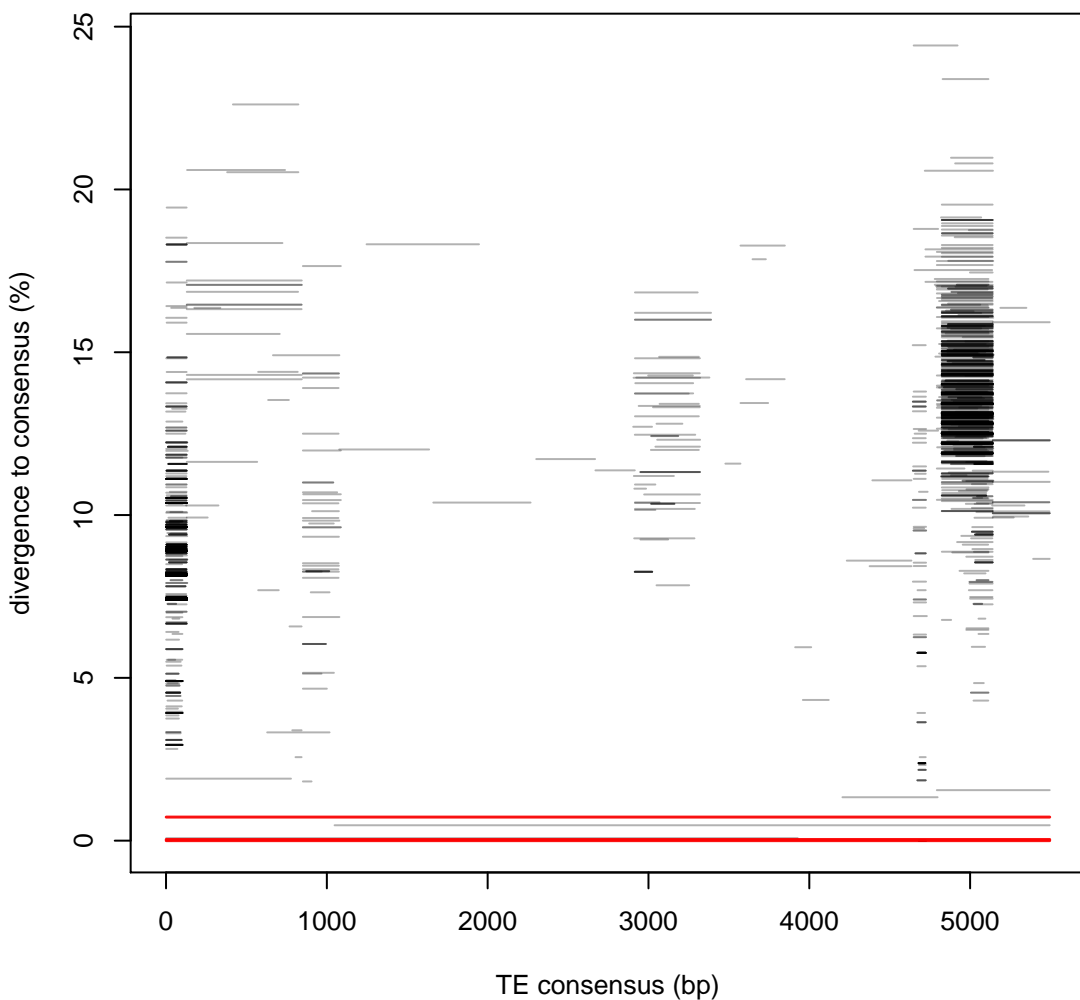
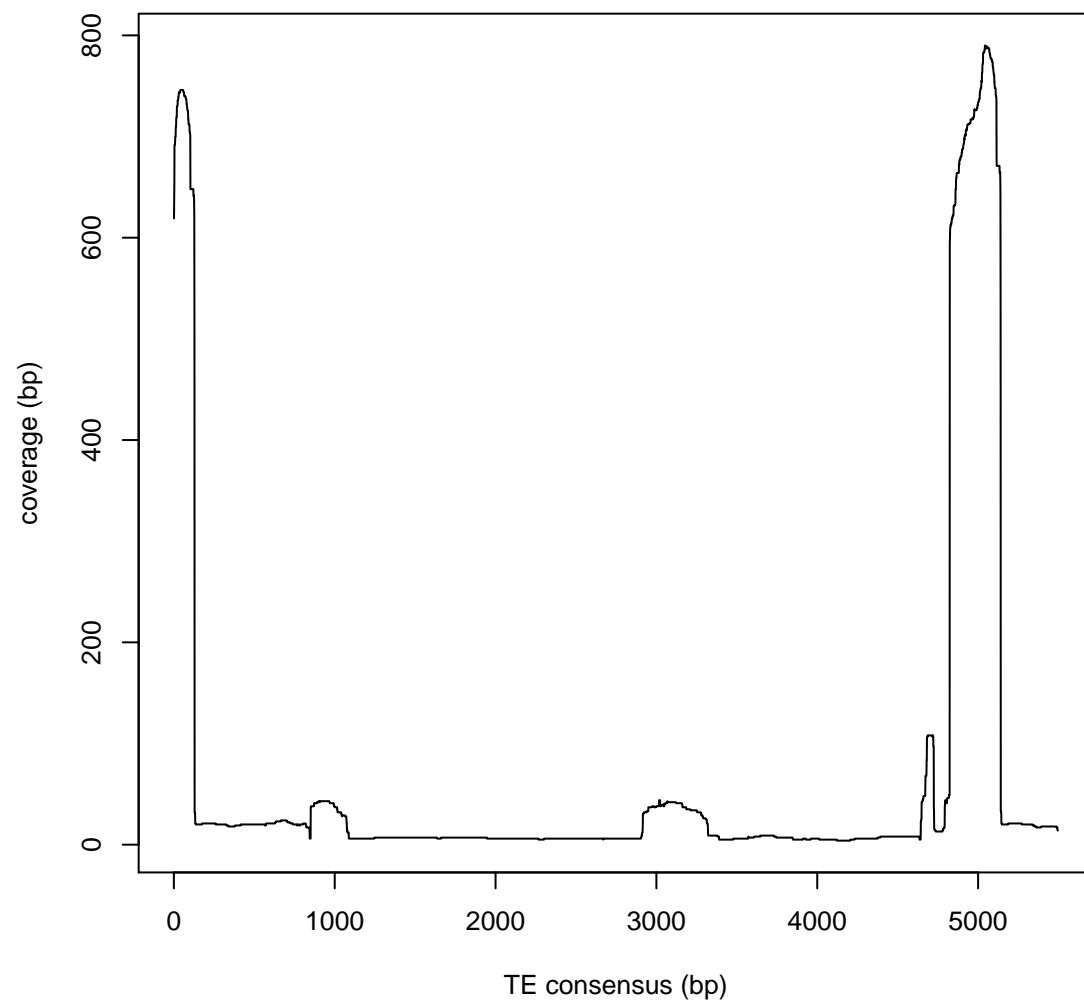


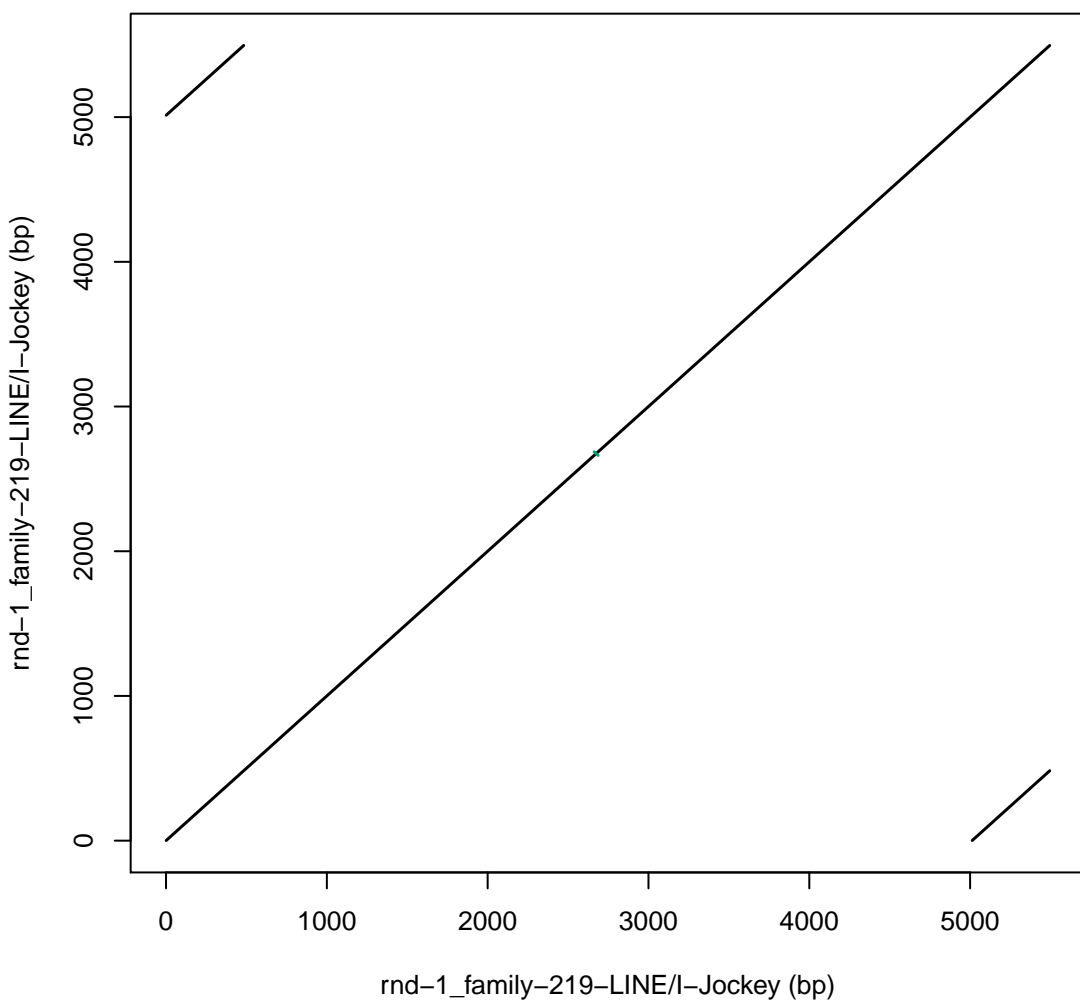
TE: rnd-1\_family-219-LINE/I-Jockey  
consensus size: 5495bp; fragments: 1788; full length: 3 (>=4945.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

