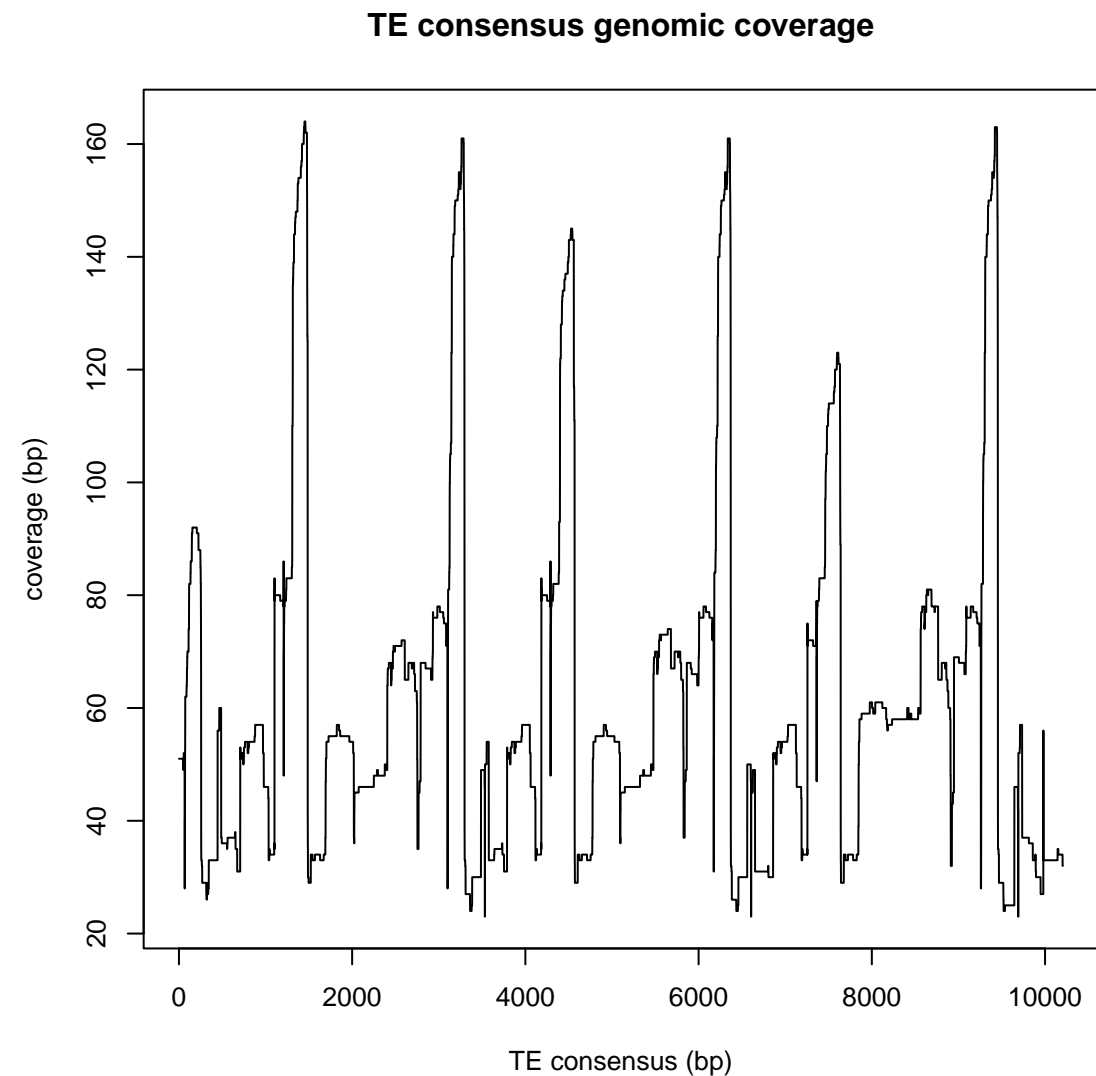
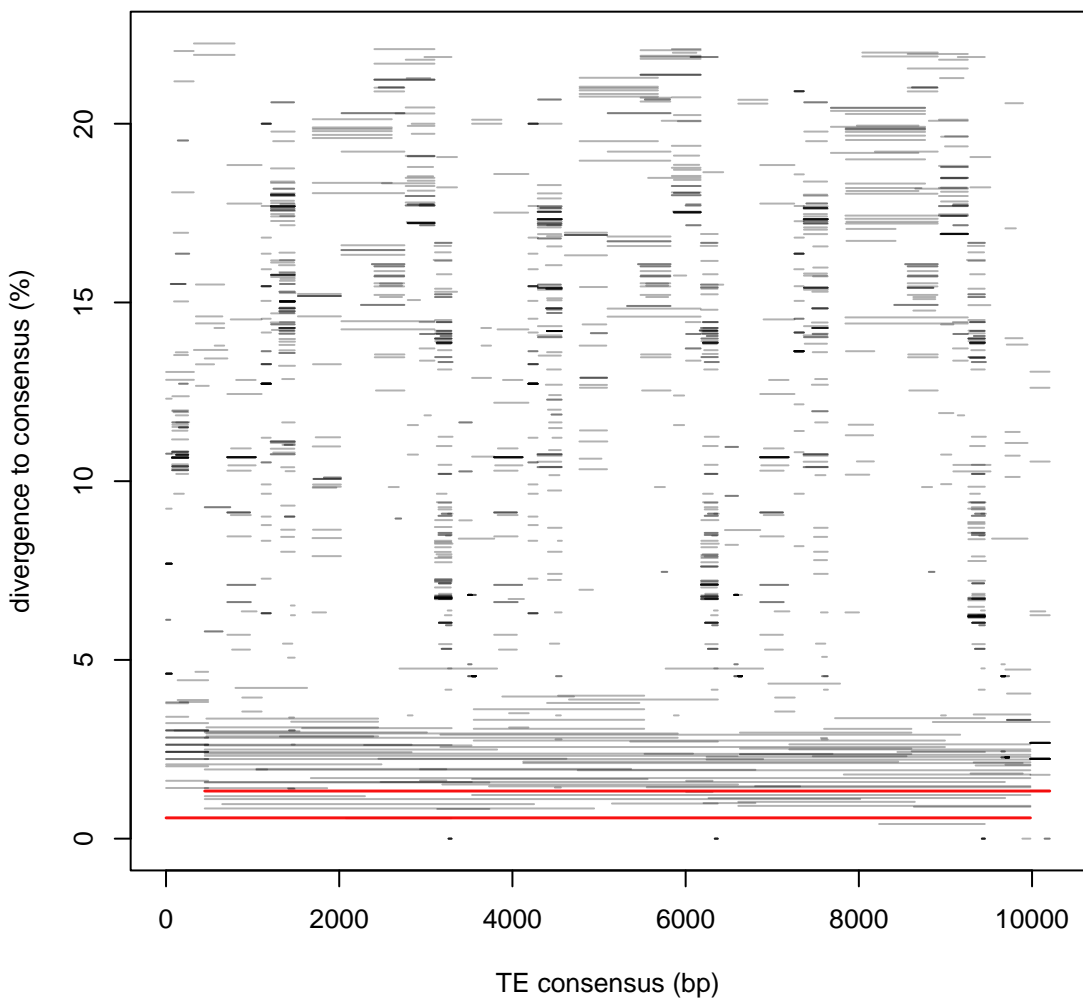
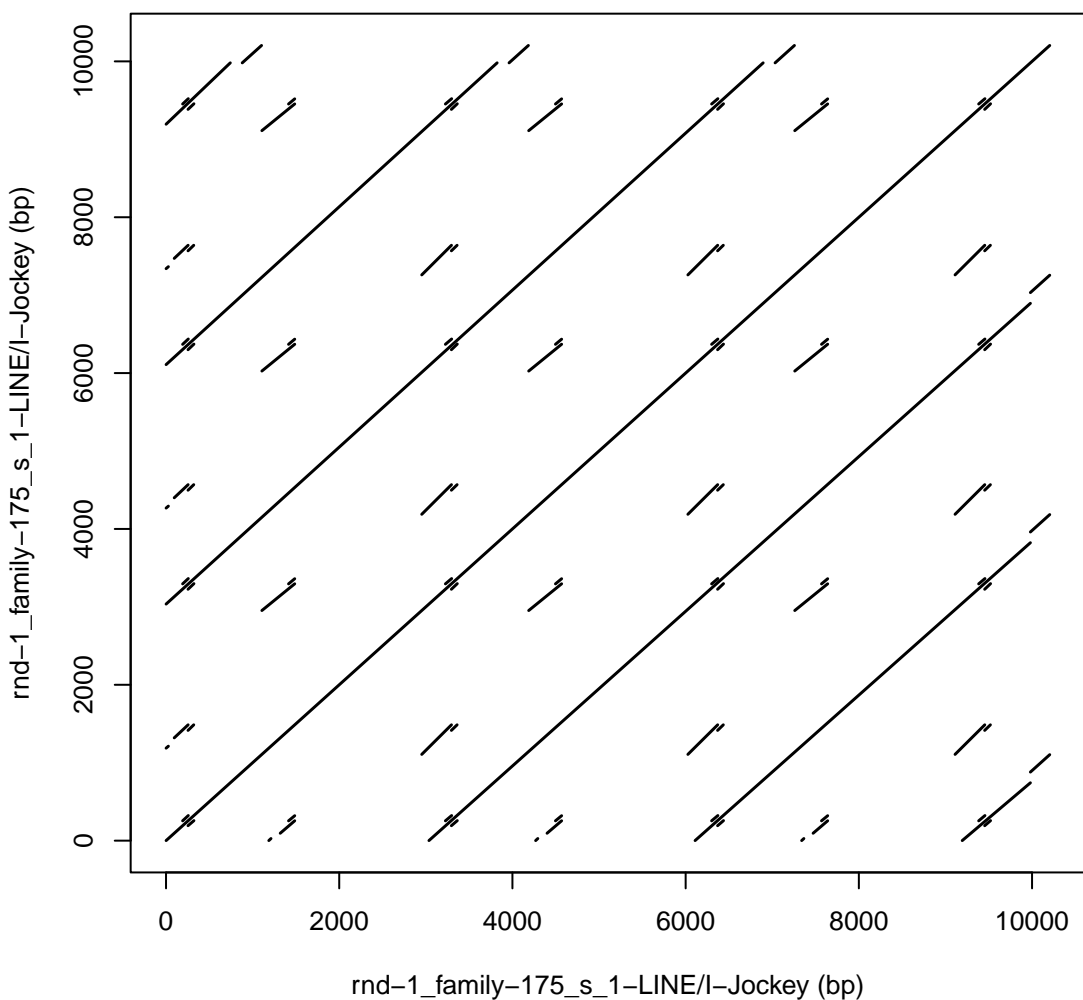


TE: rnd-1_family-175_s_1-LINE/I-Jockey
consensus size: 10204bp; fragments: 1753; full length: 2 (>=9183.6bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

