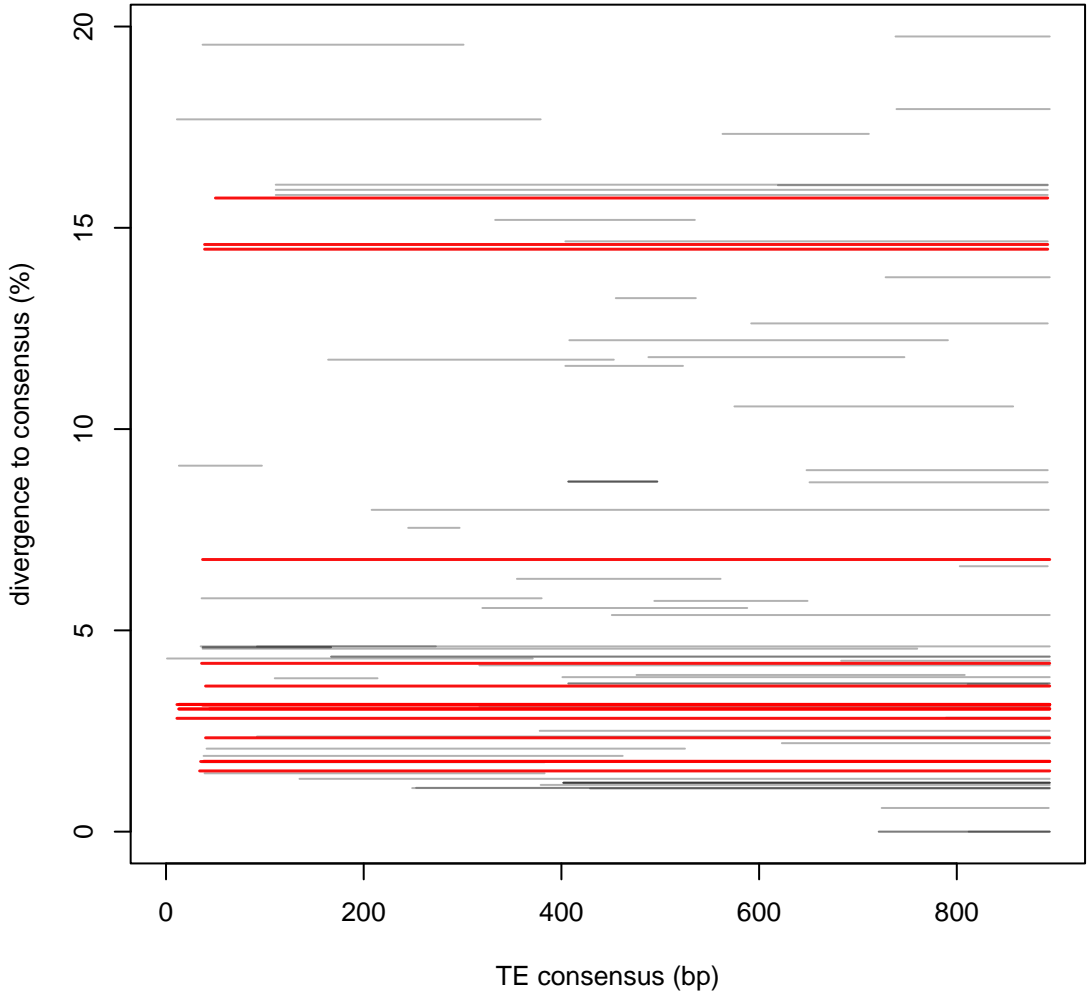
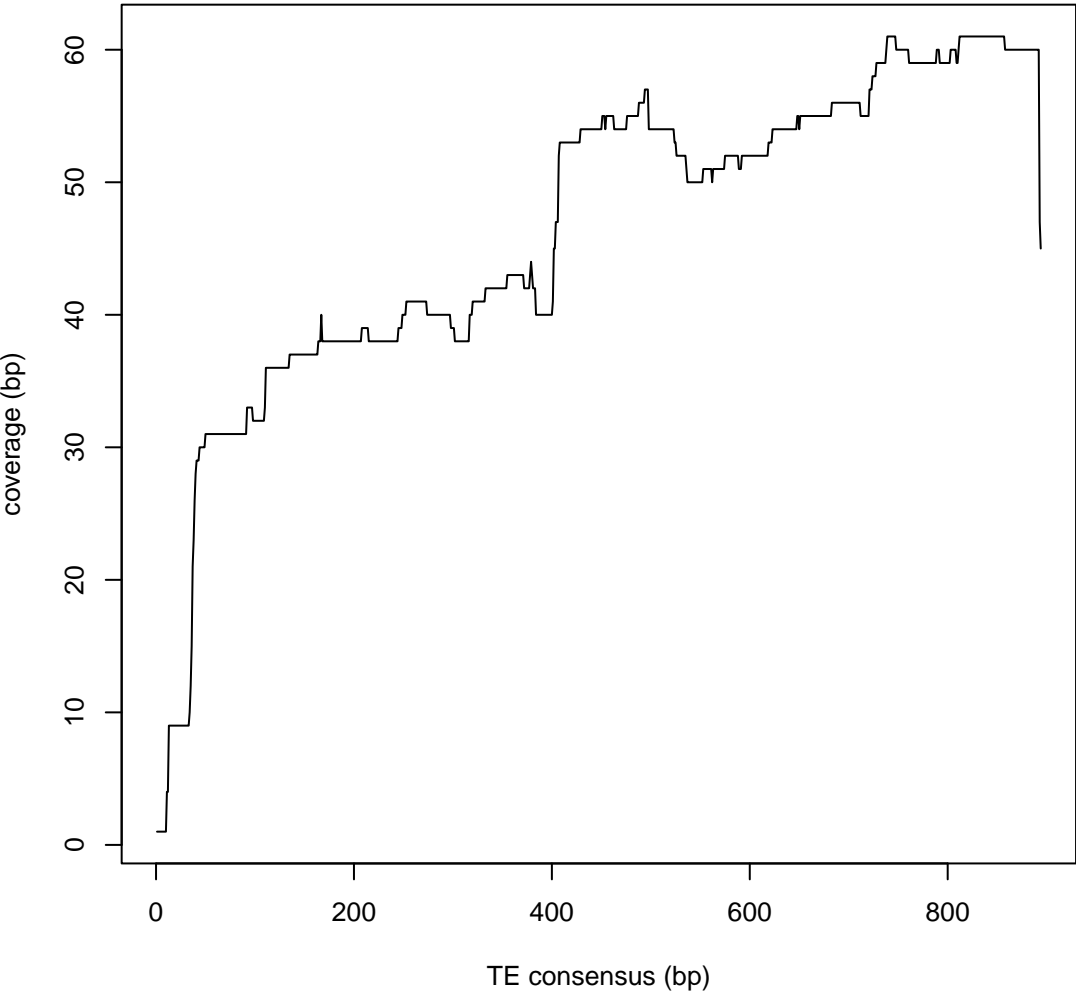


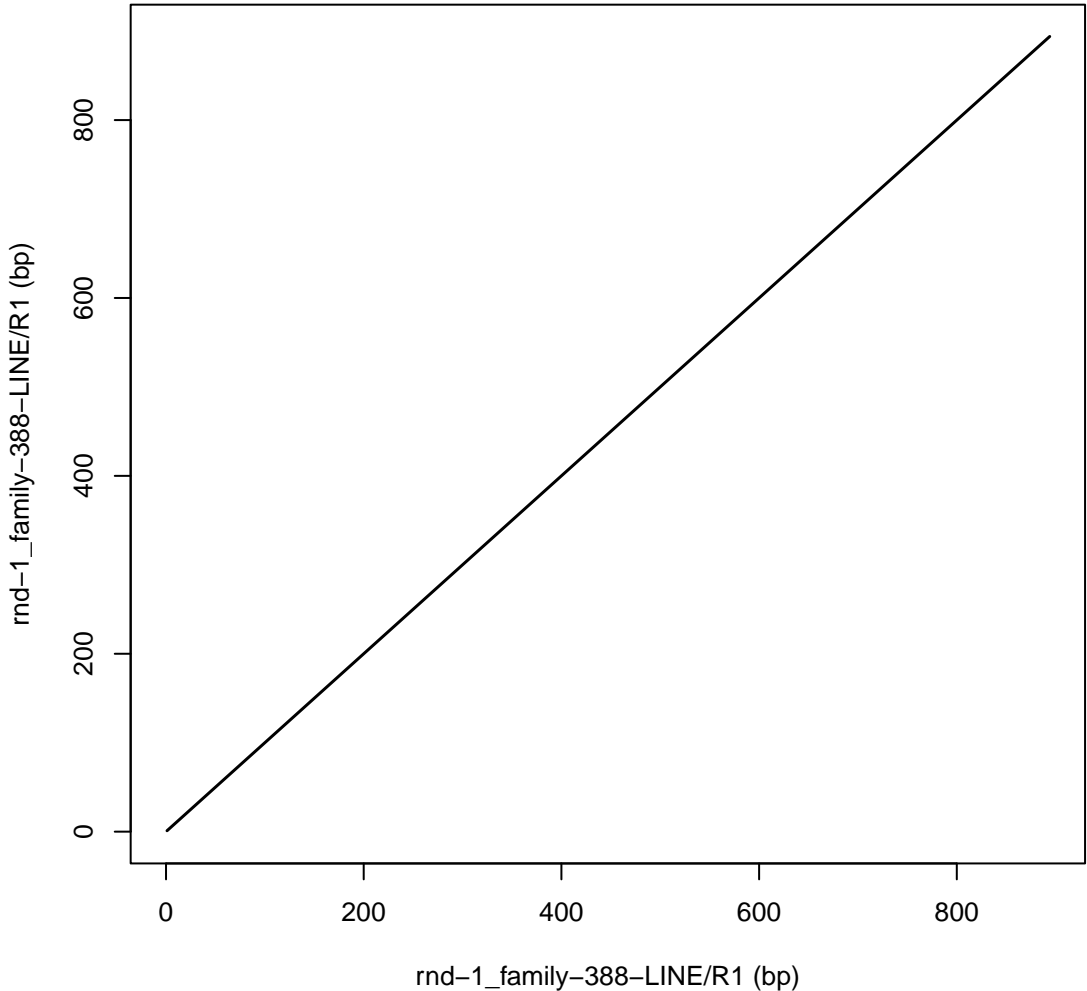
TE: rnd-1_family-388-LINE/R1
consensus size: 894bp; fragments: 90; full length: 18 (>=804.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

