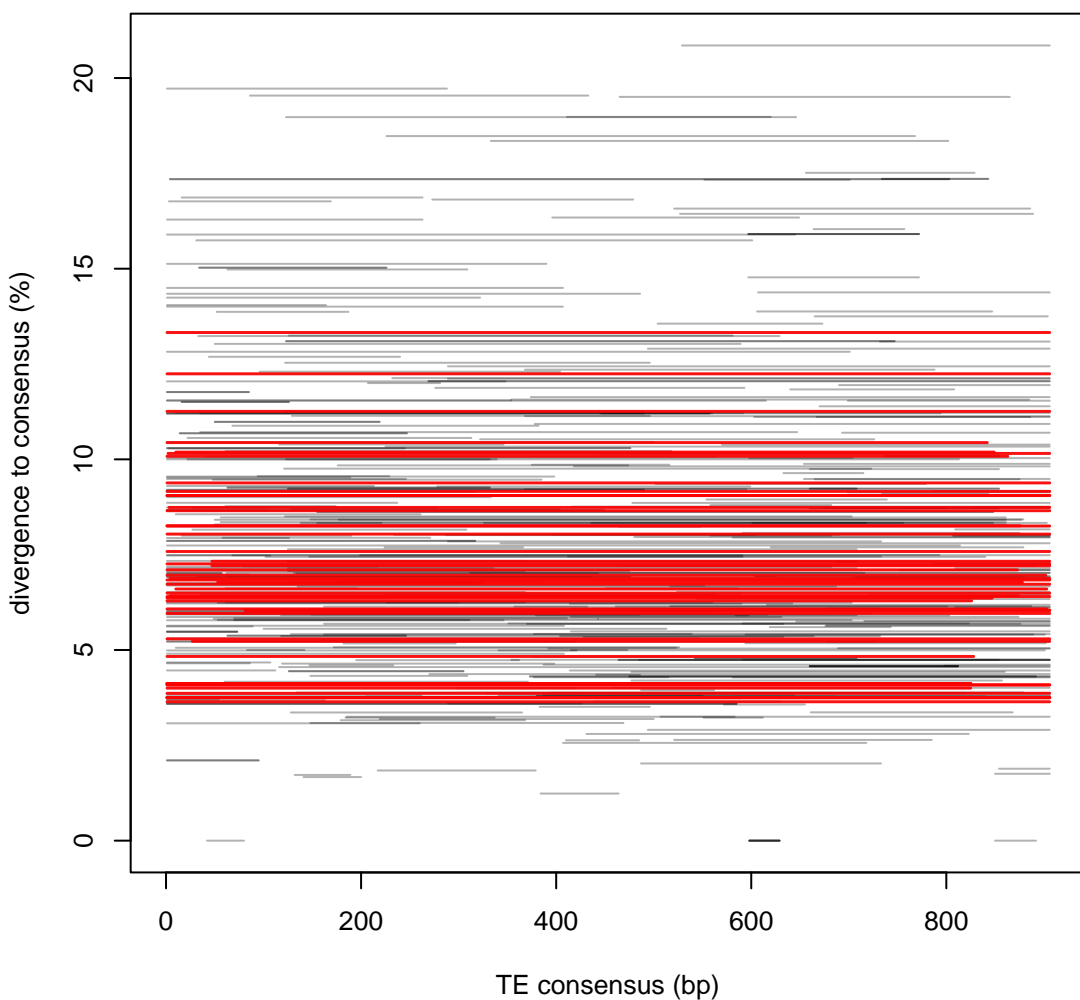
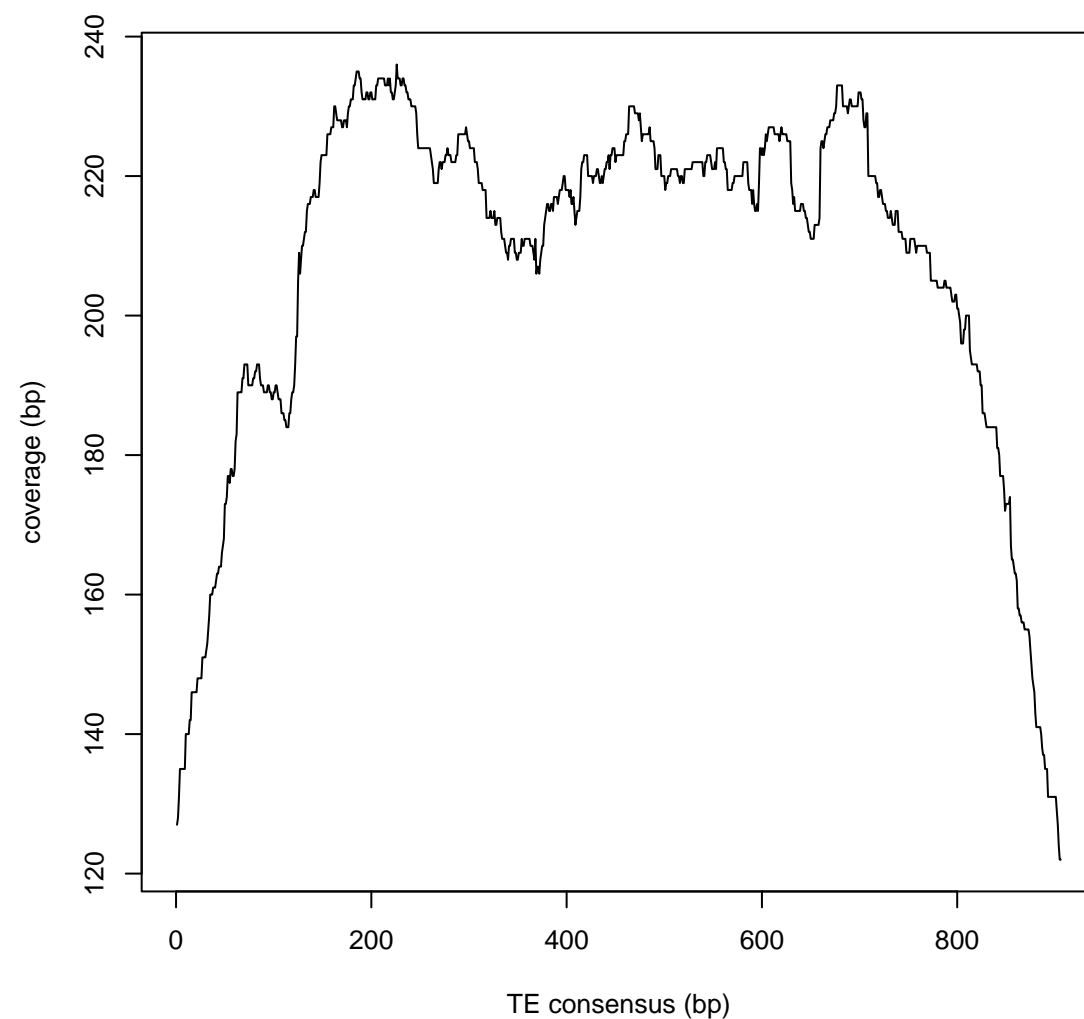


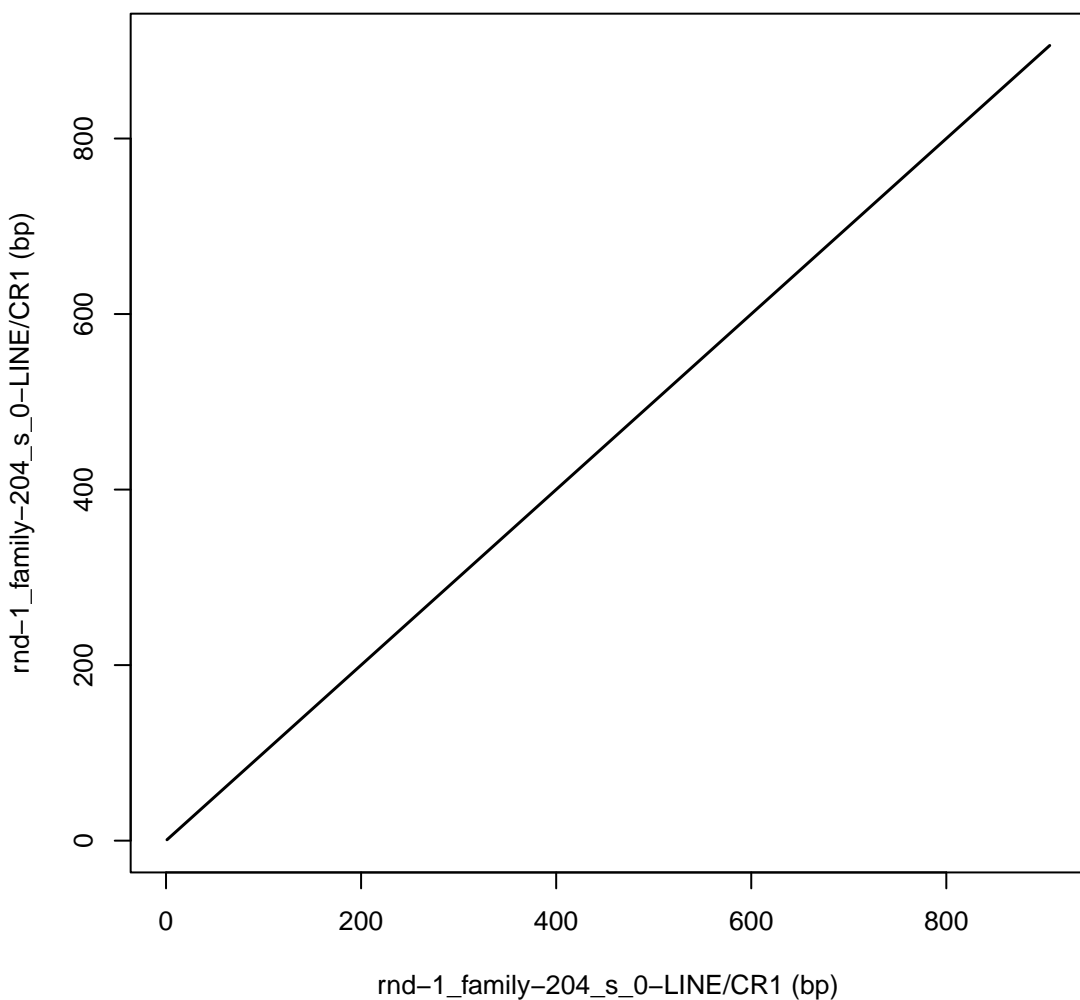
TE: rnd-1\_family-204\_s\_0-LINE/CR1  
consensus size: 906bp; fragments: 480; full length: 51 (>=815.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

