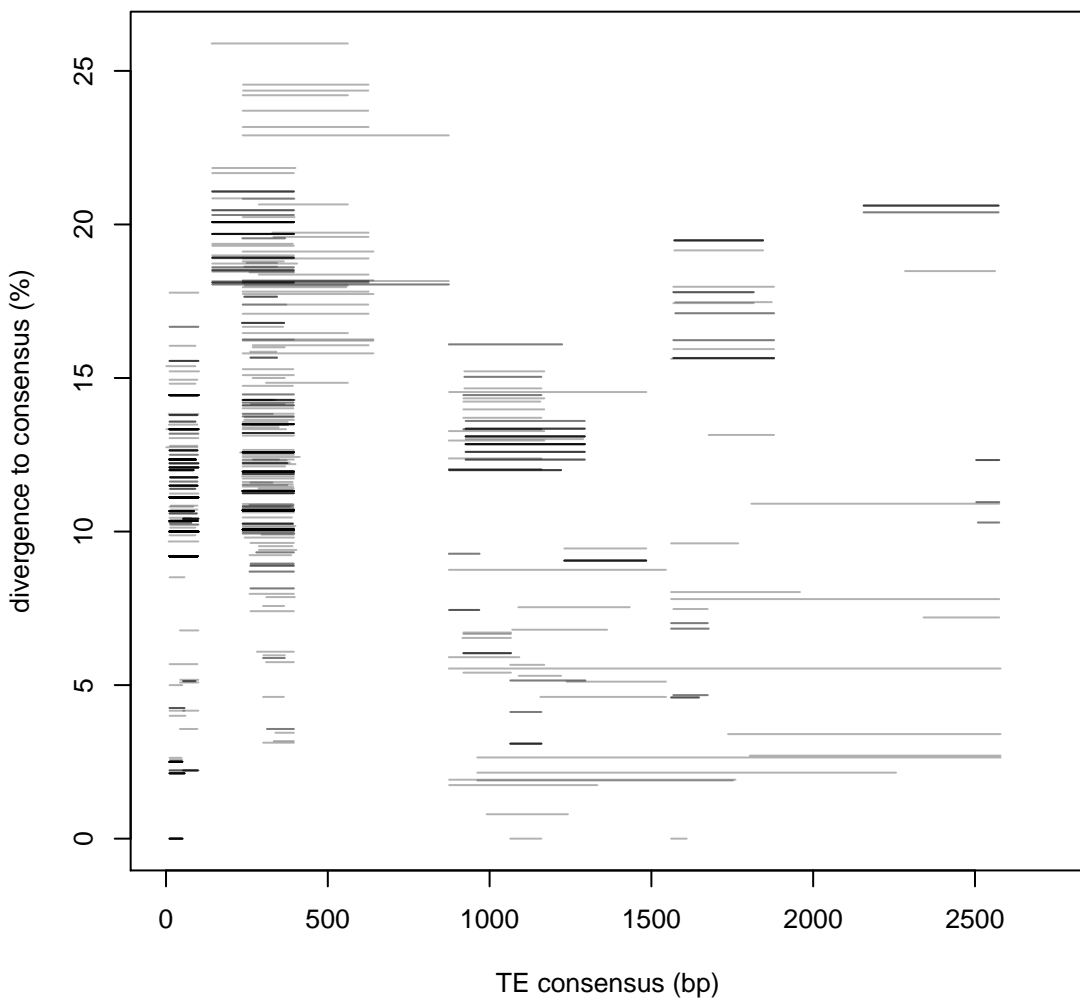
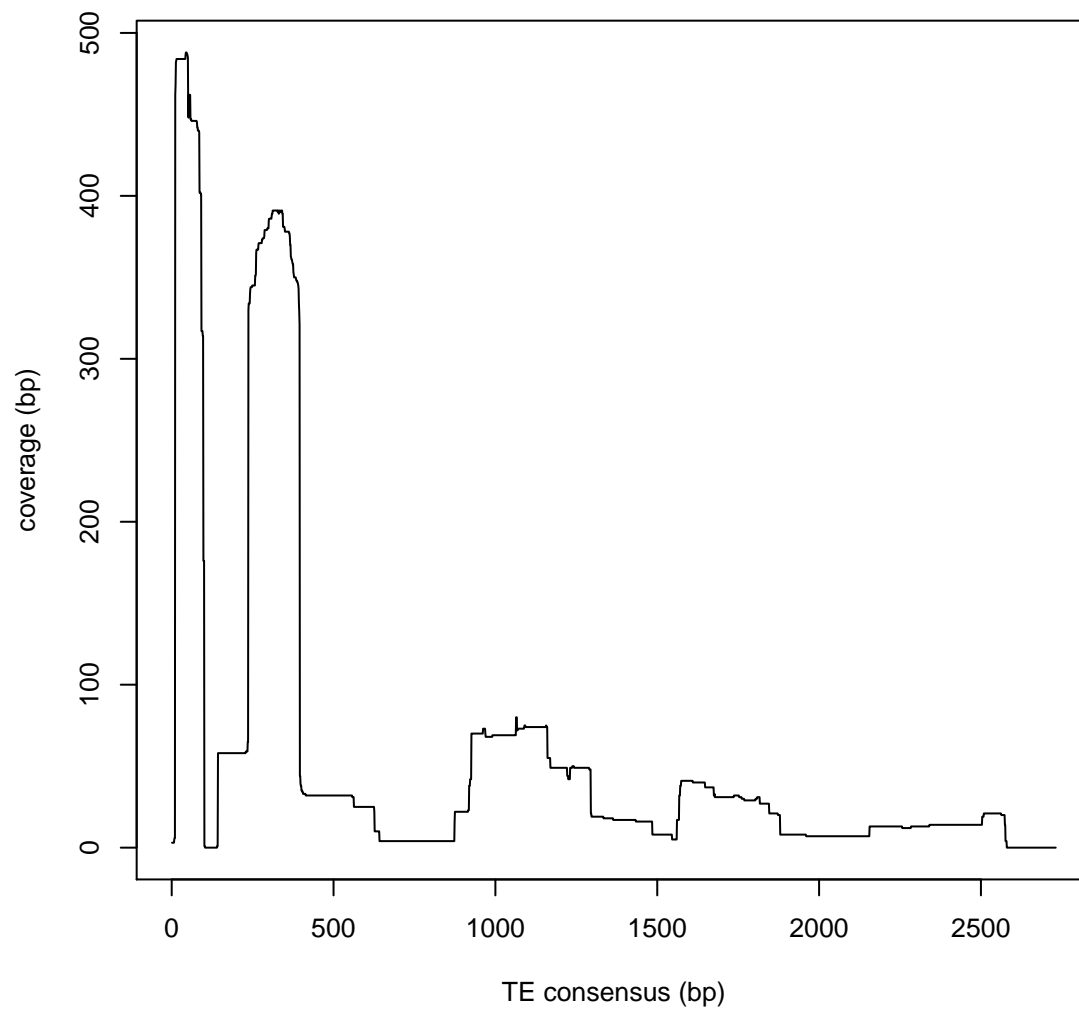


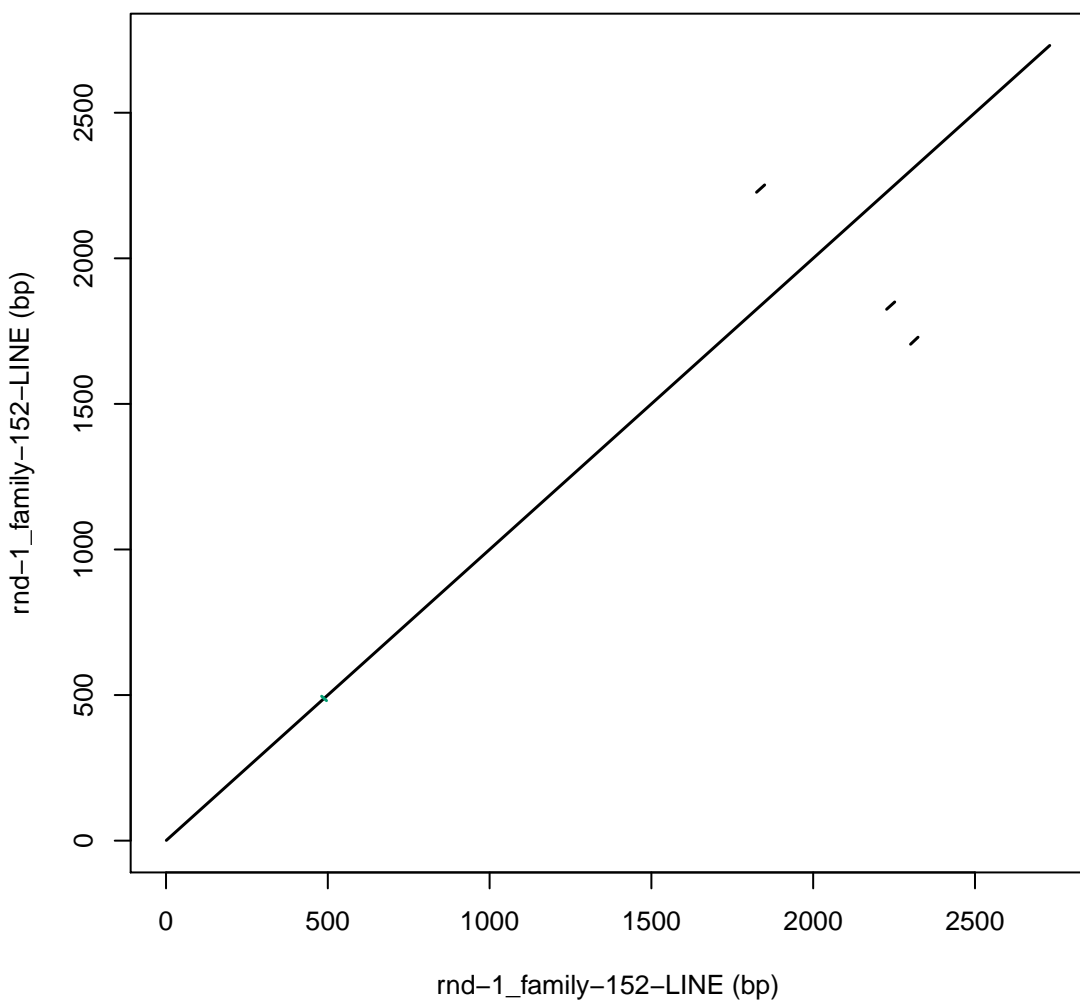
TE: rnd-1\_family-152-LINE  
consensus size: 2731bp; fragments: 1049; full length: 0 (>=2457.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

