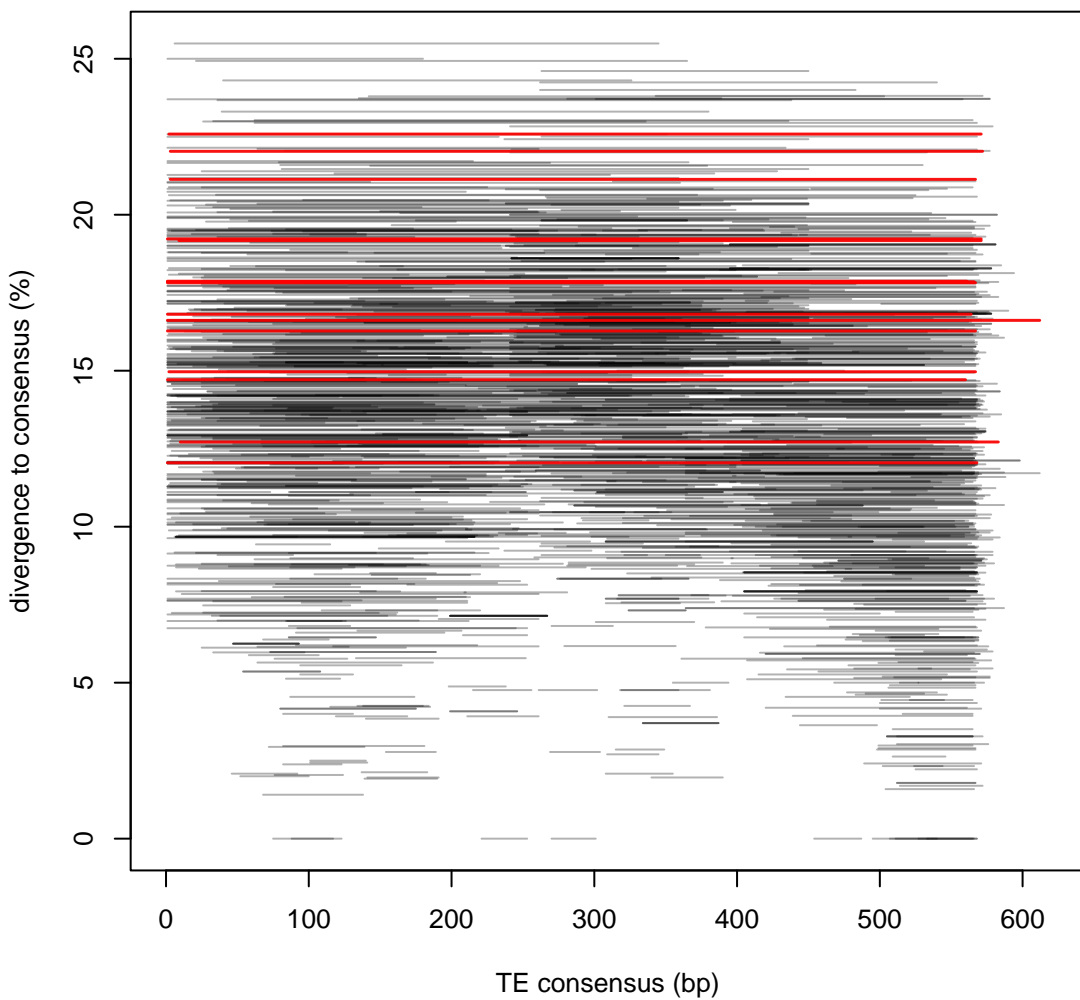
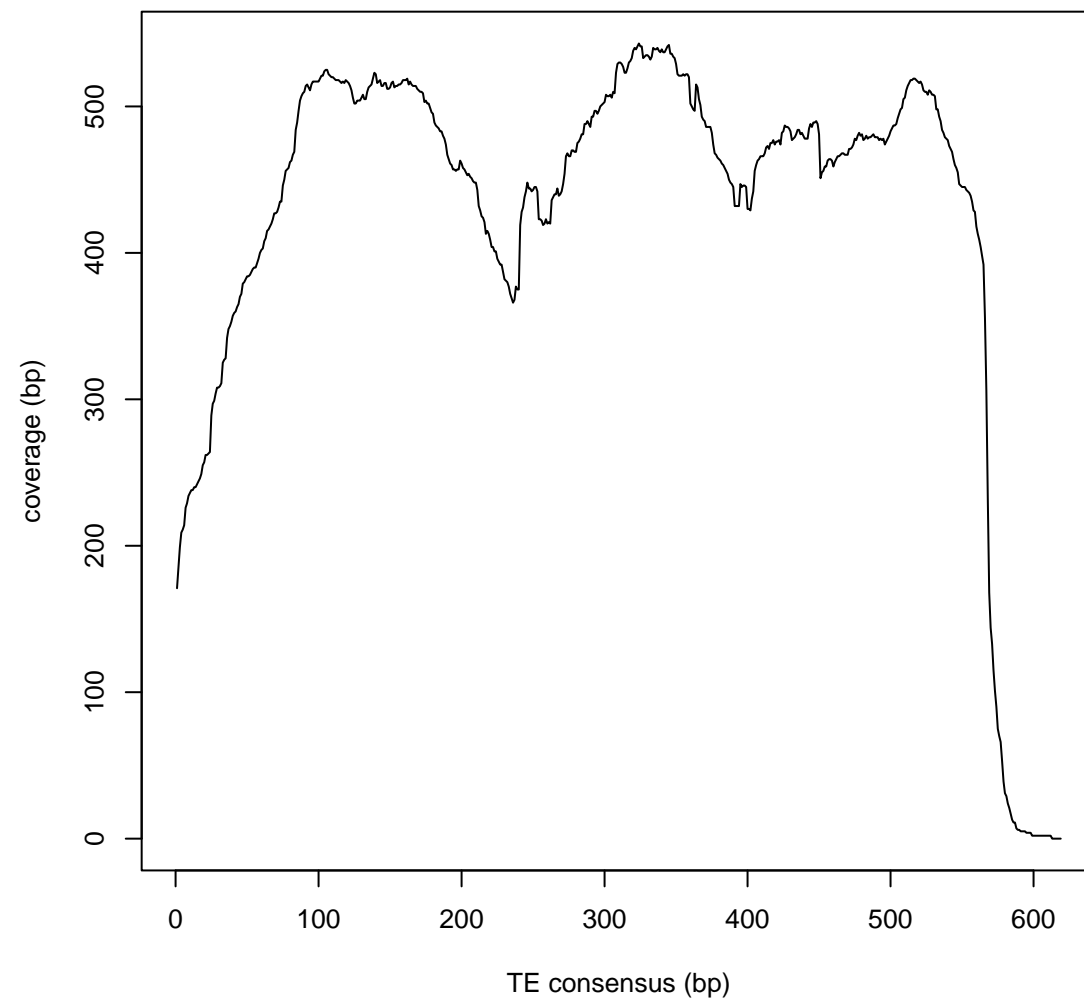


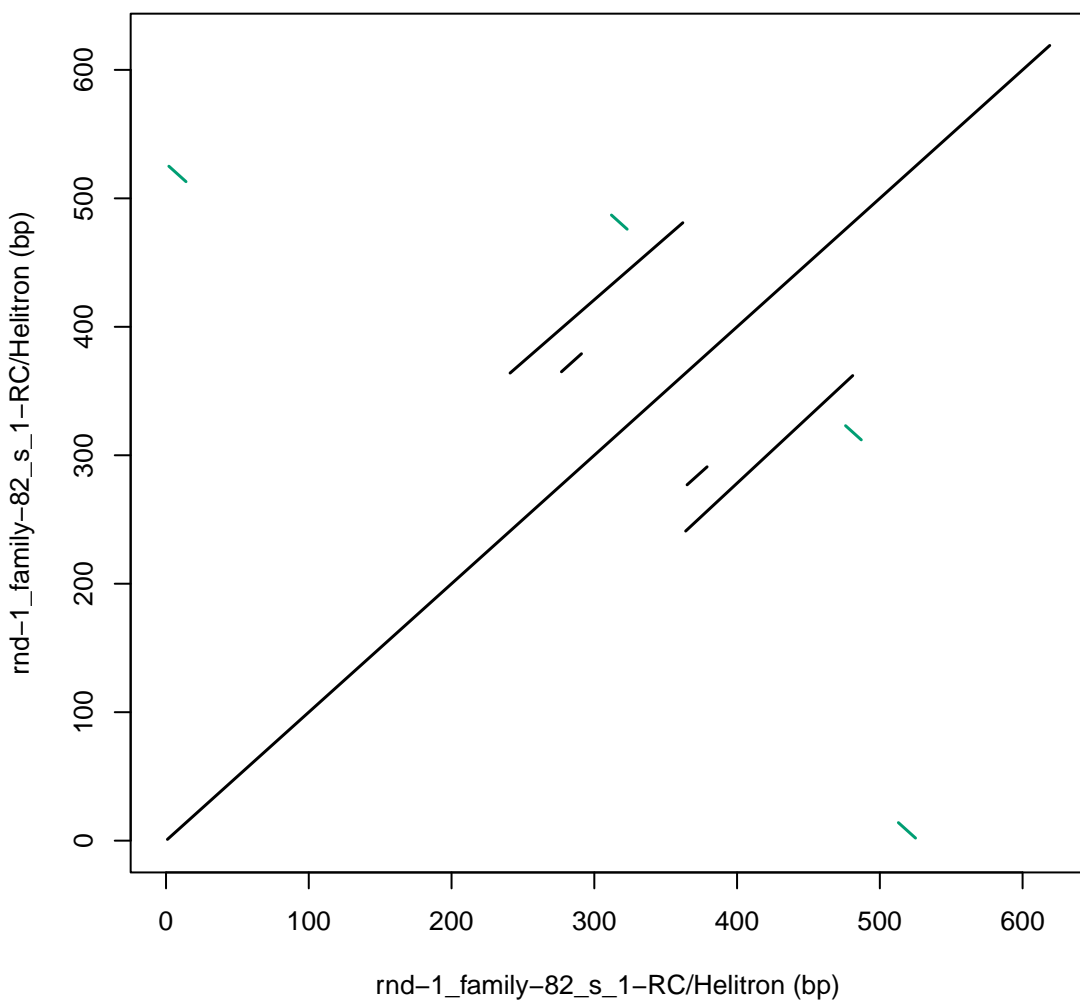
TE: rnd-1\_family-82\_s\_1-RC/Helitron  
consensus size: 619bp; fragments: 1373; full length: 15 (>=557.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

