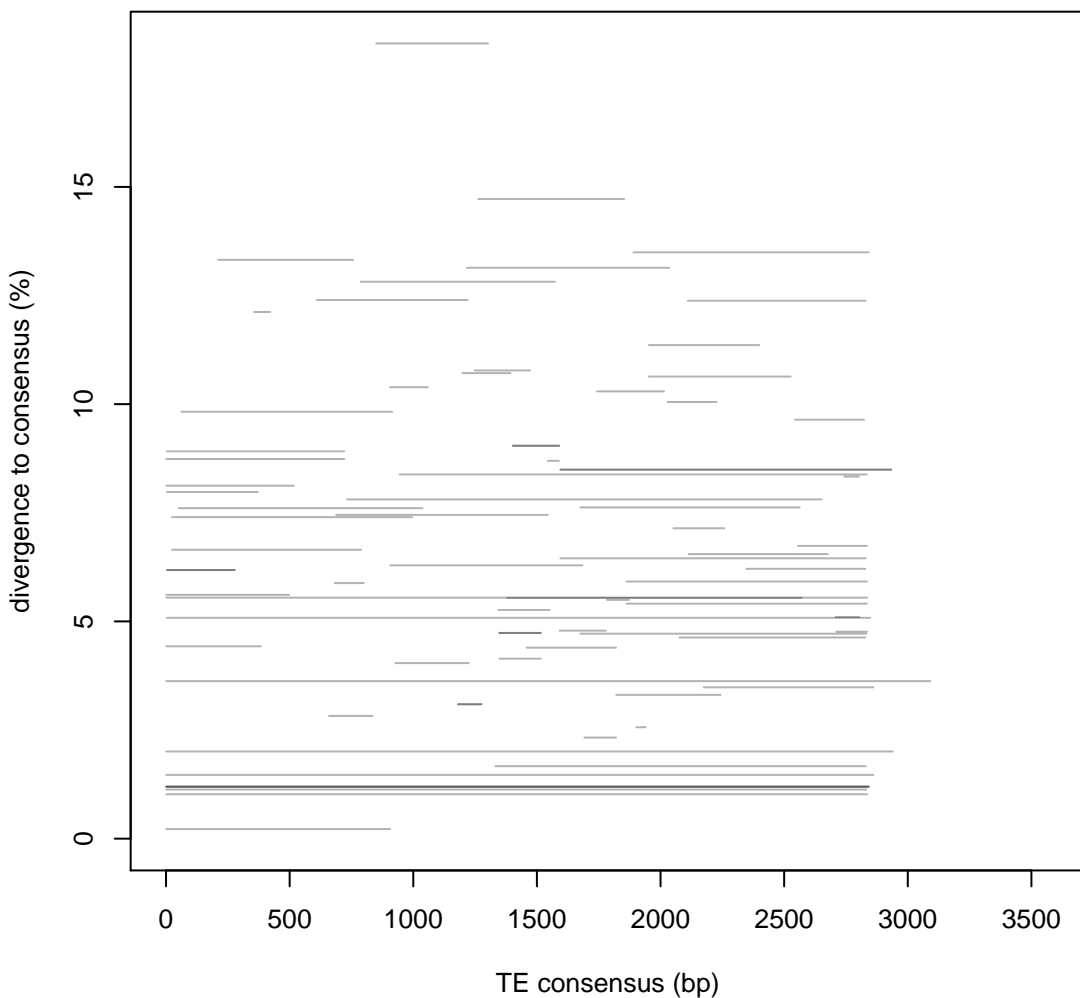
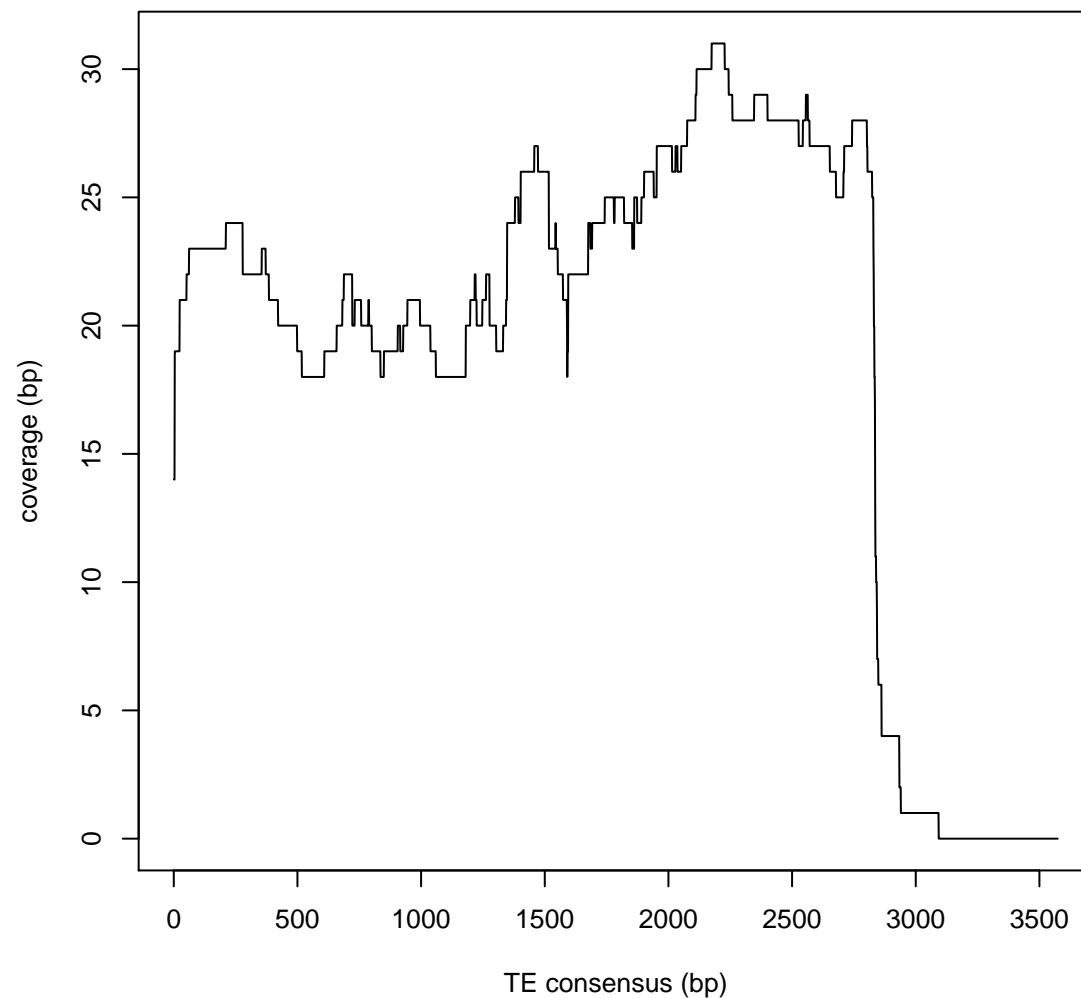


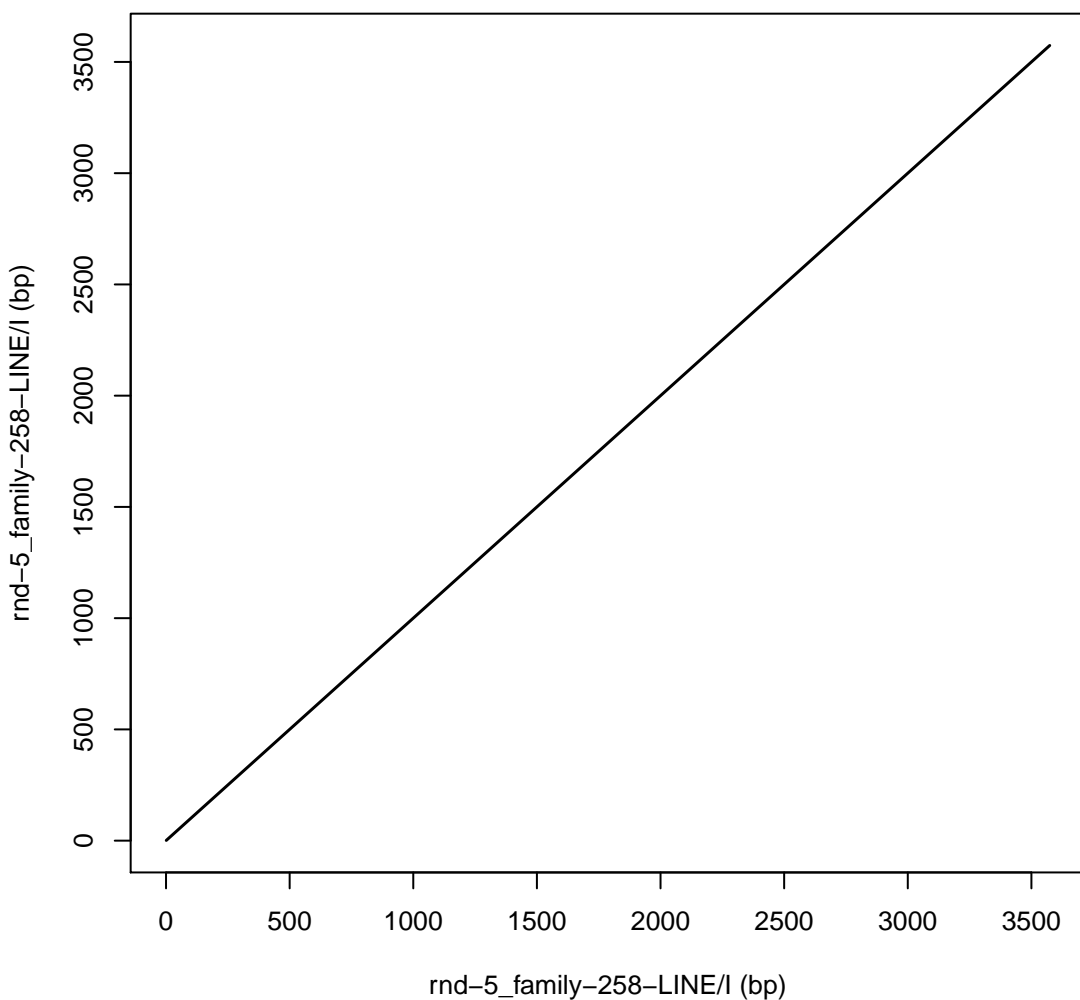
TE: rnd-5\_family-258-LINE/I  
consensus size: 3574bp; fragments: 80; full length: 0 (>=3216.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

