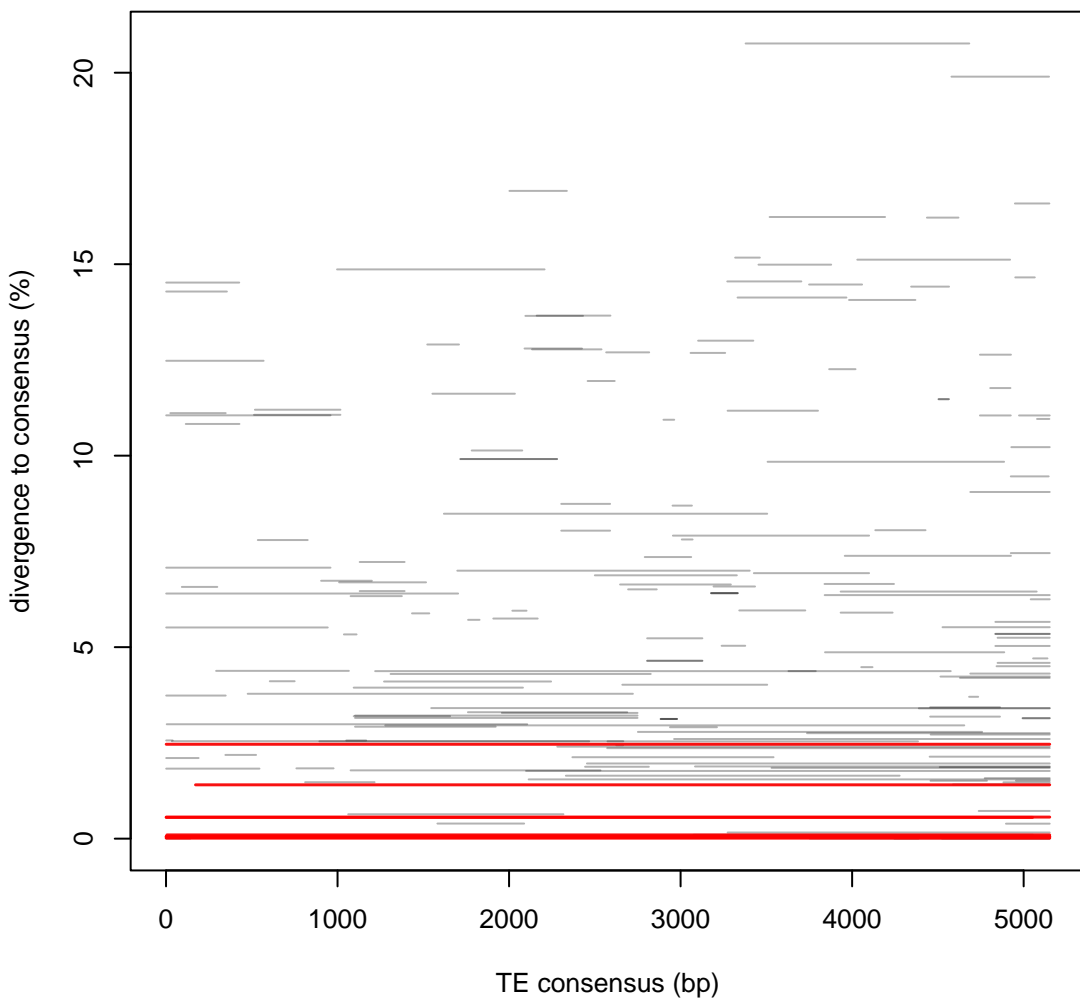
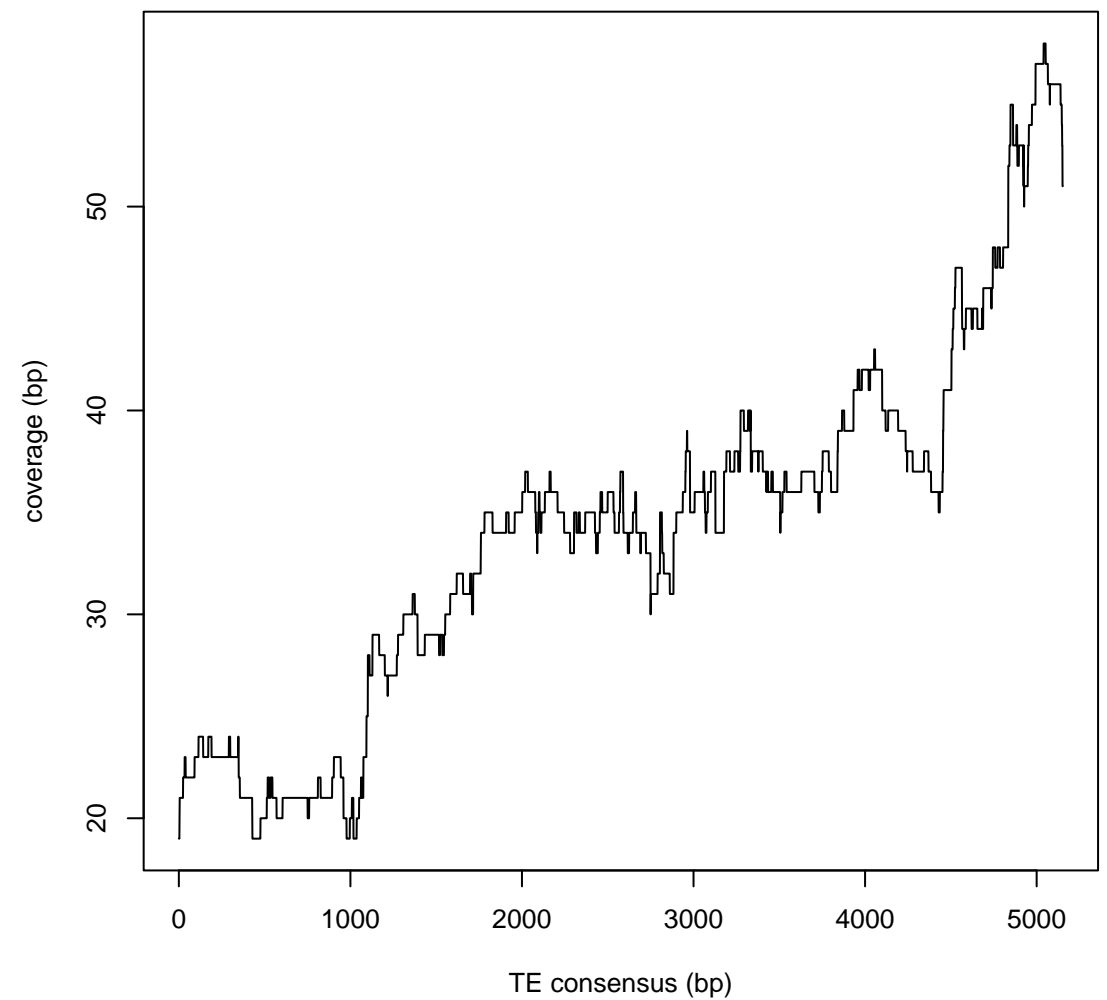


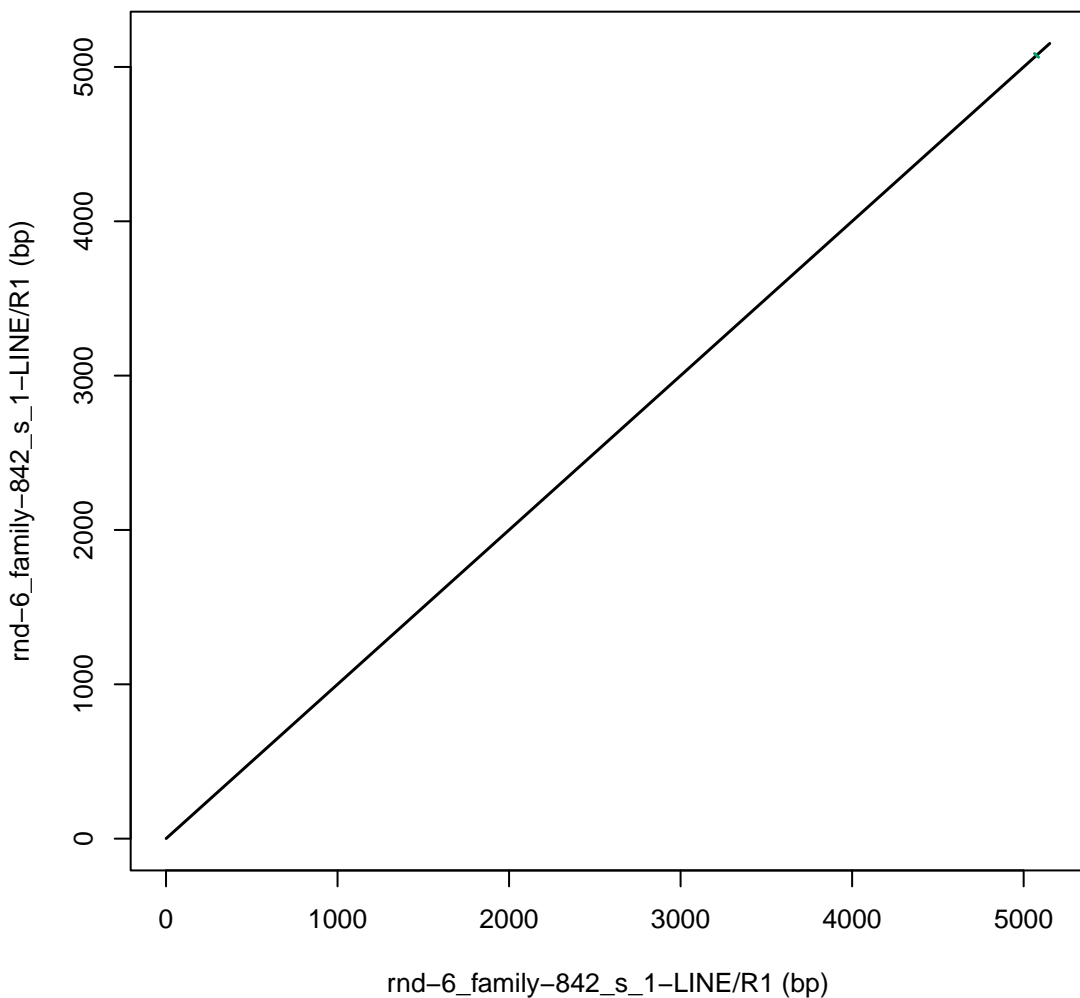
TE: rnd-6\_family-842\_s\_1-LINE/R1  
consensus size: 5152bp; fragments: 191; full length: 9 (>=4636.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

