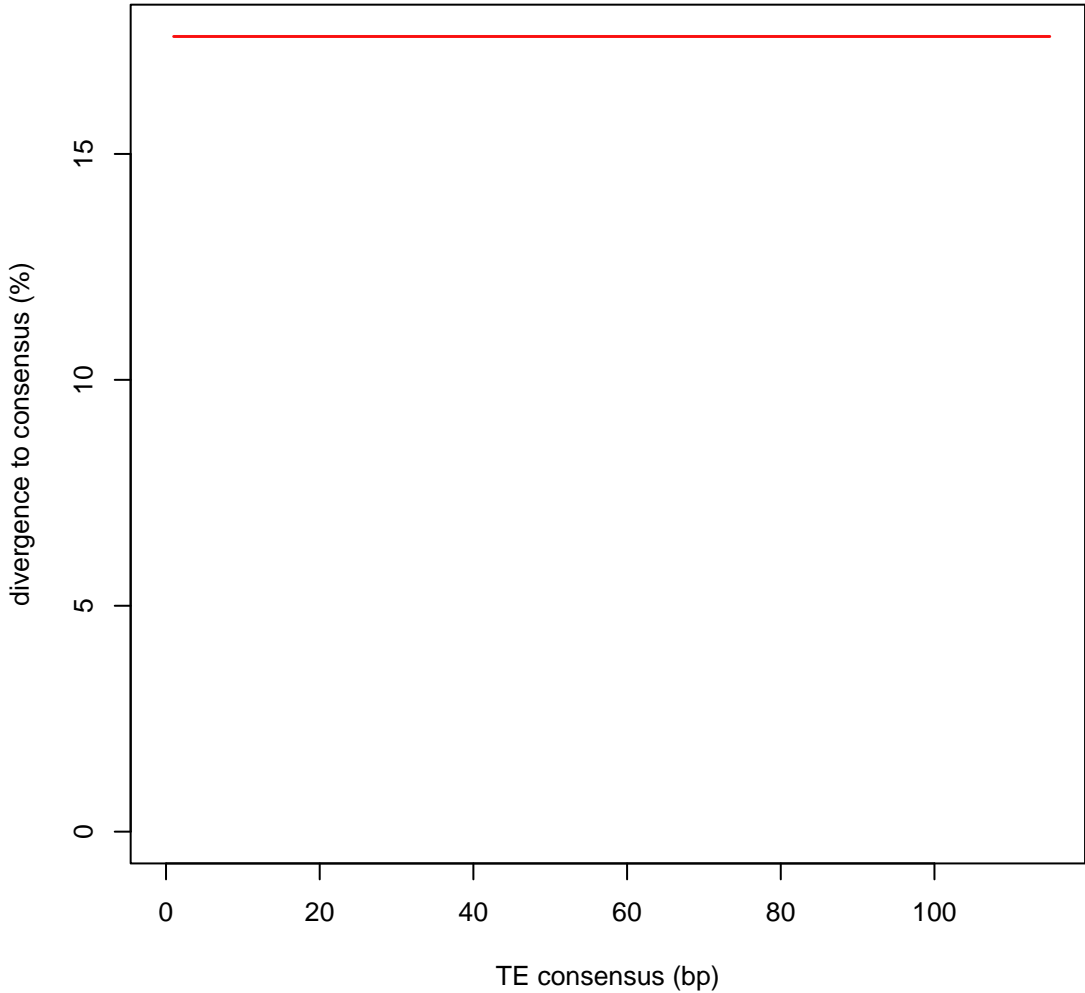
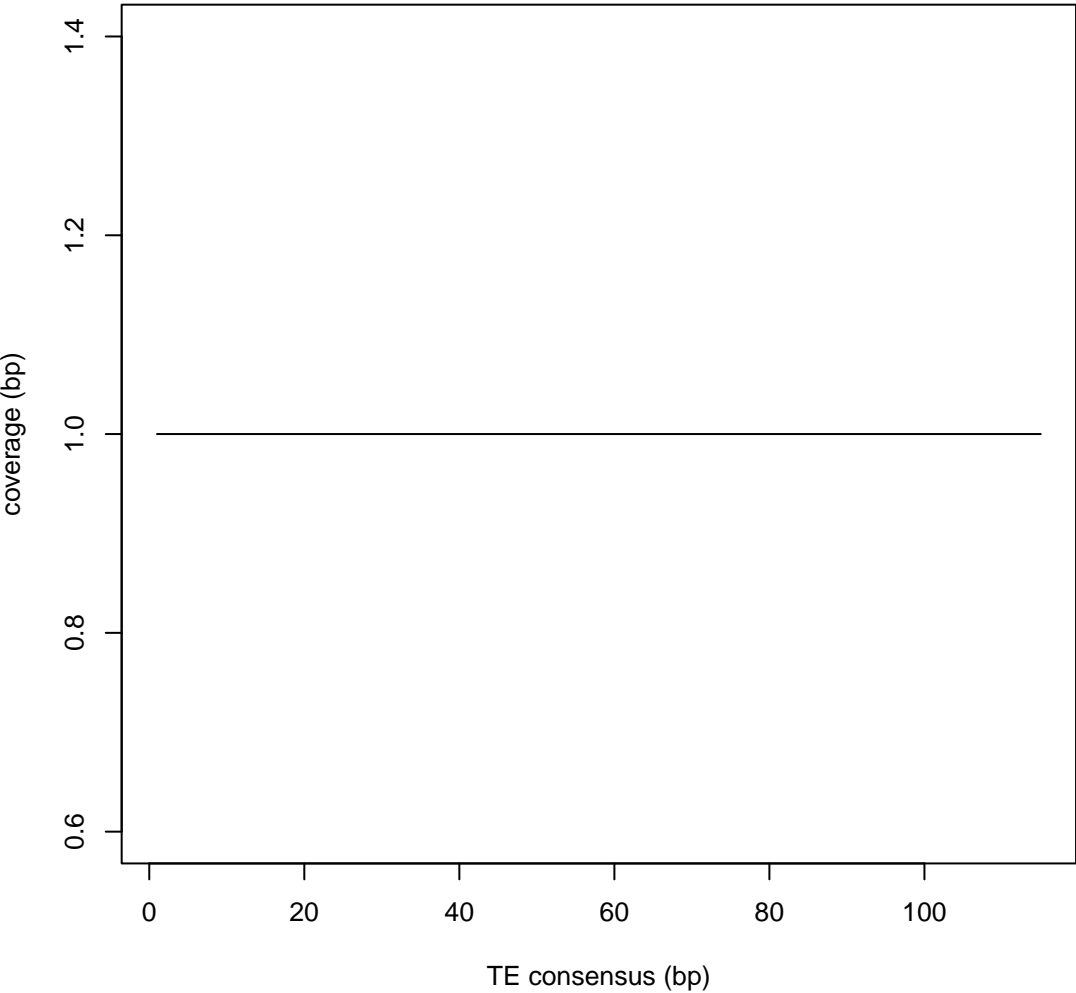


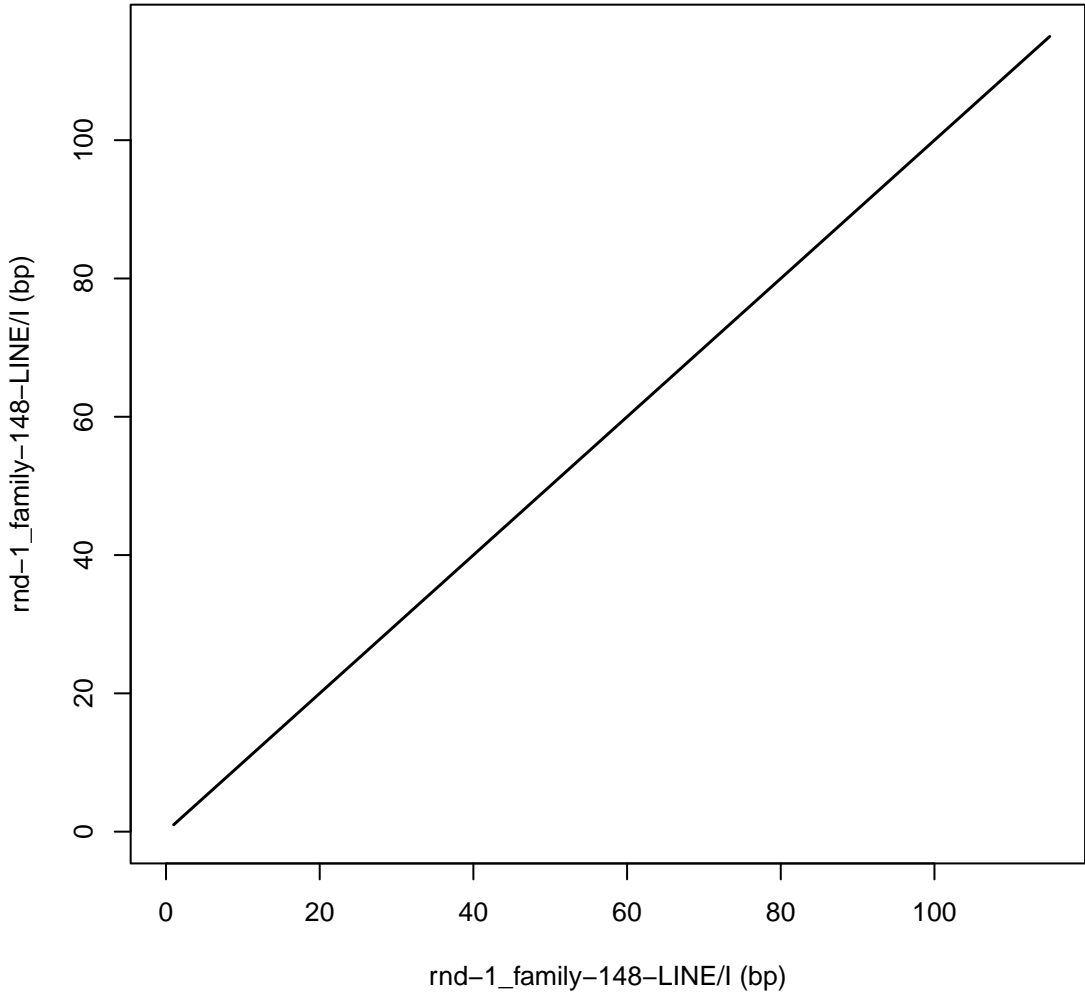
TE: rnd-1\_family-148-LINE/I  
consensus size: 115bp; fragments: 1; full length: 1 (>=103.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

