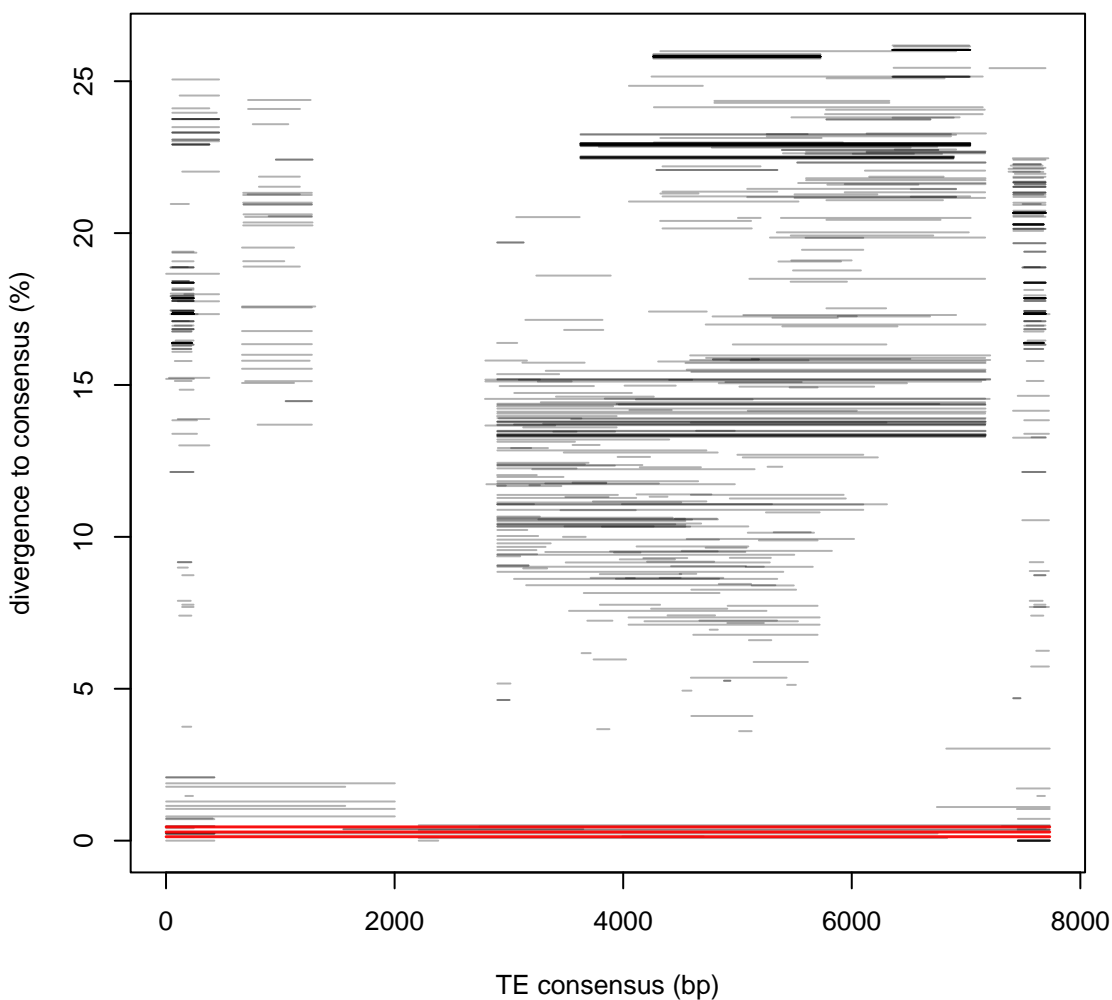
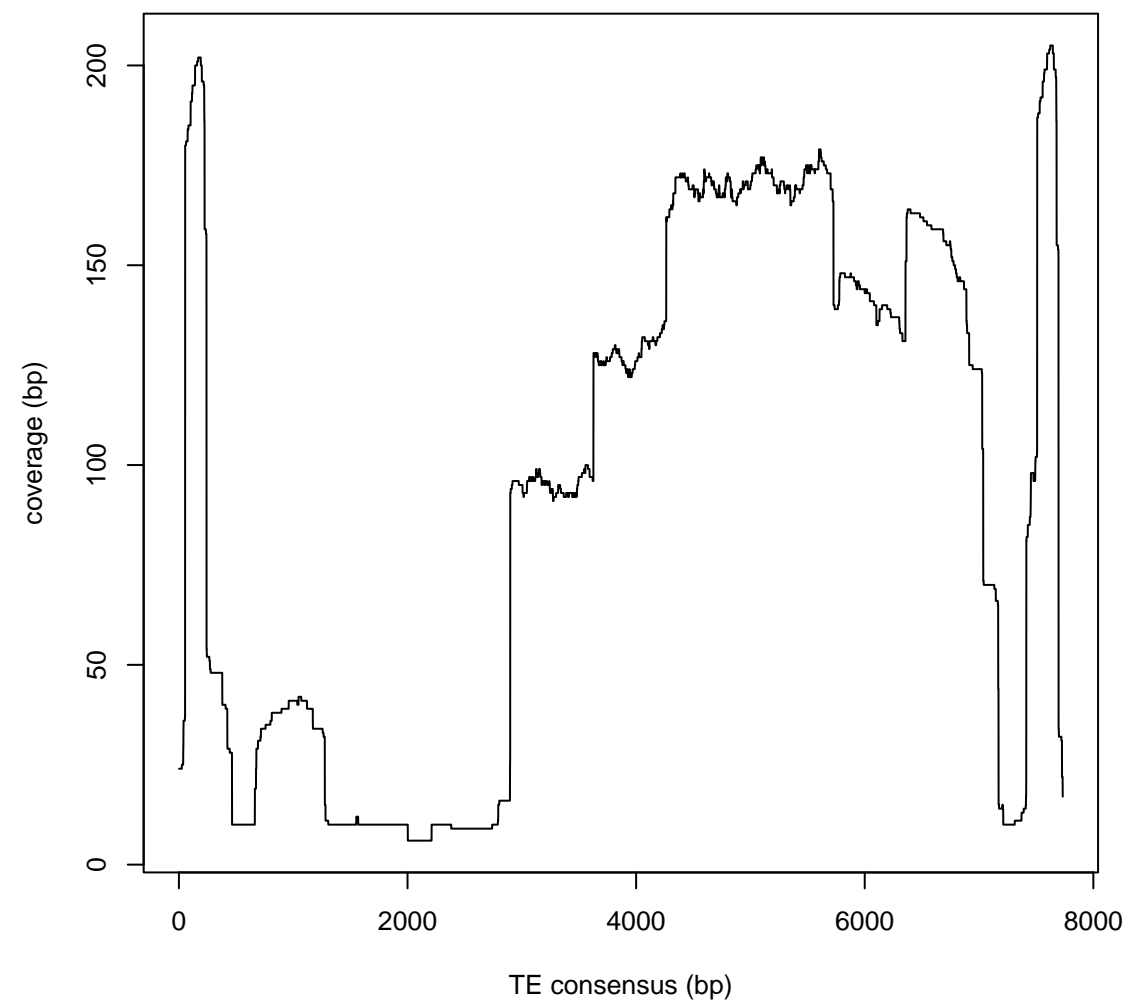


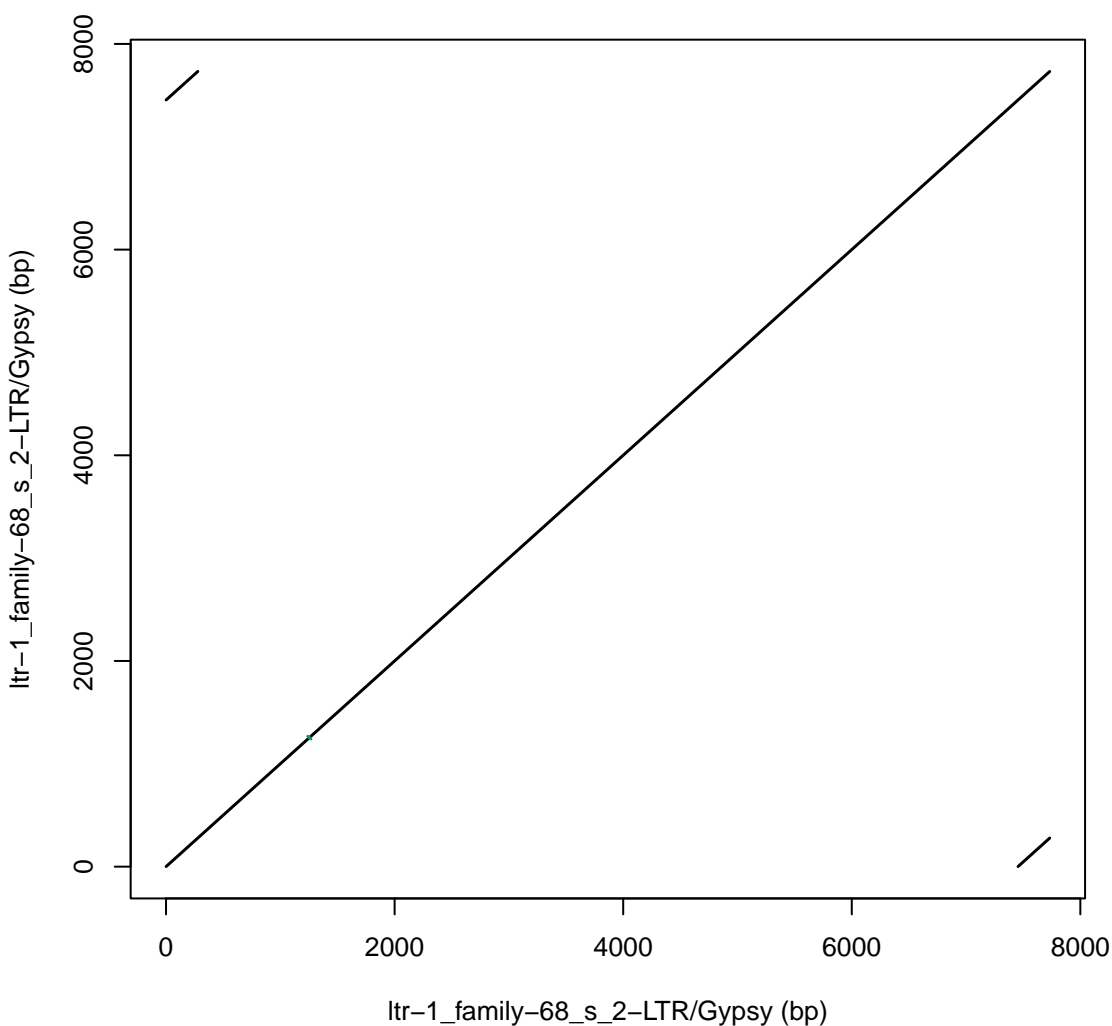
TE: ltr-1\_family-68\_s\_2-LTR/Gypsy  
consensus size: 7732bp; fragments: 818; full length: 3 (>=6958.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

