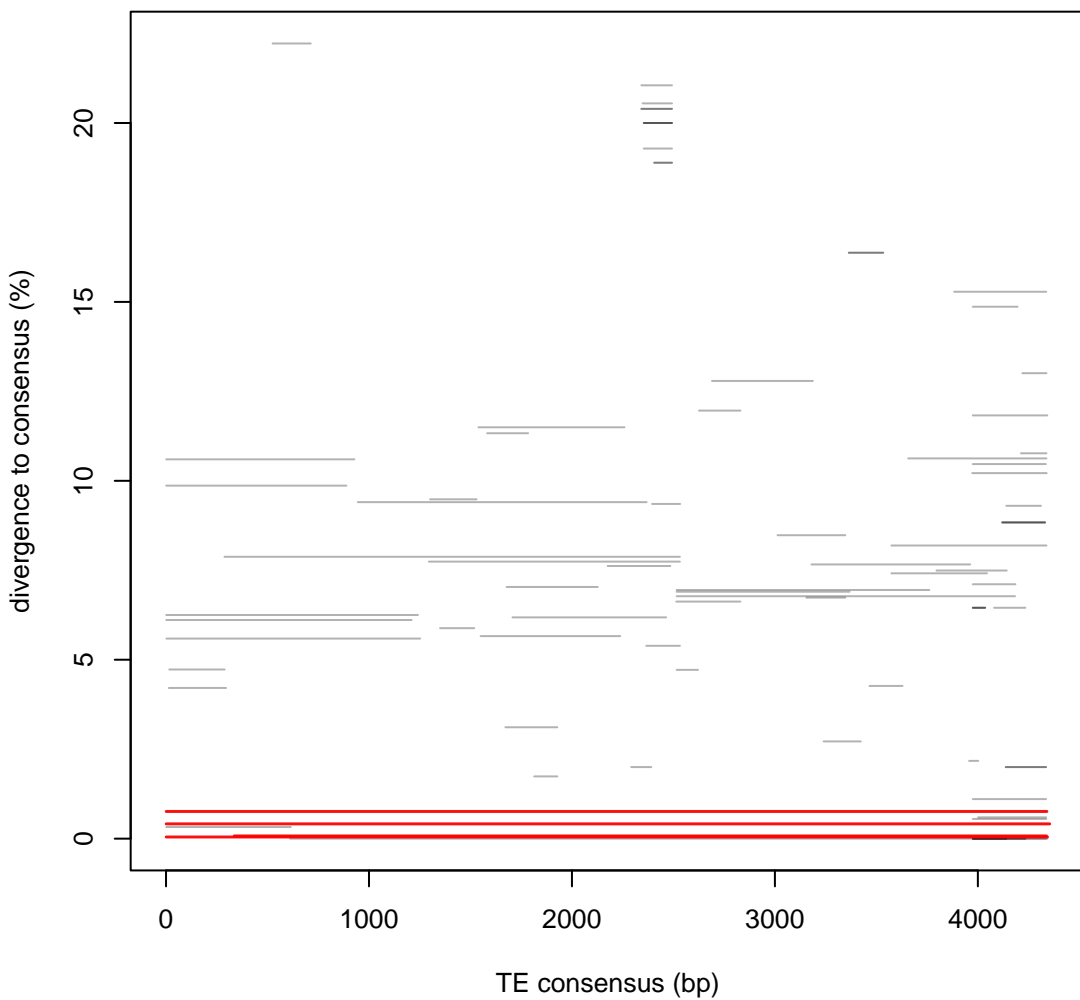
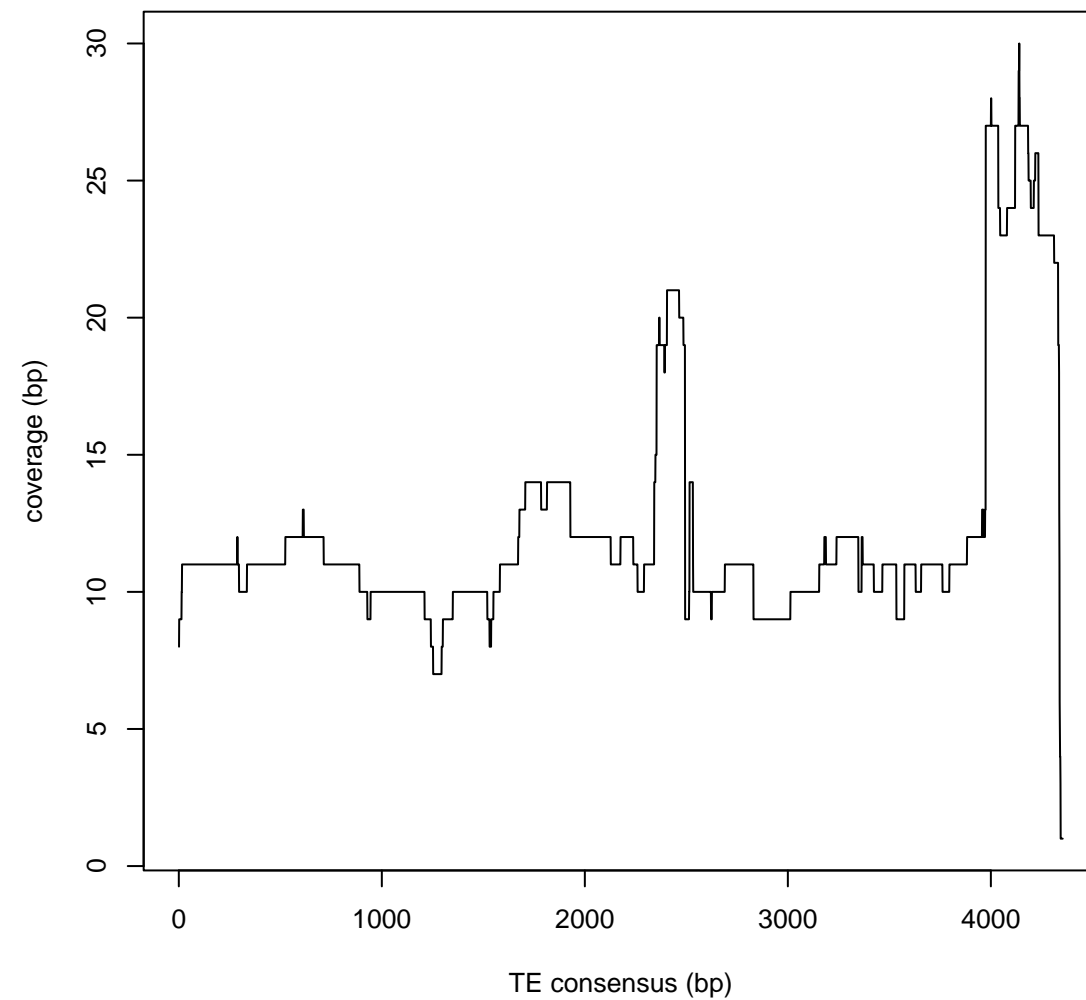


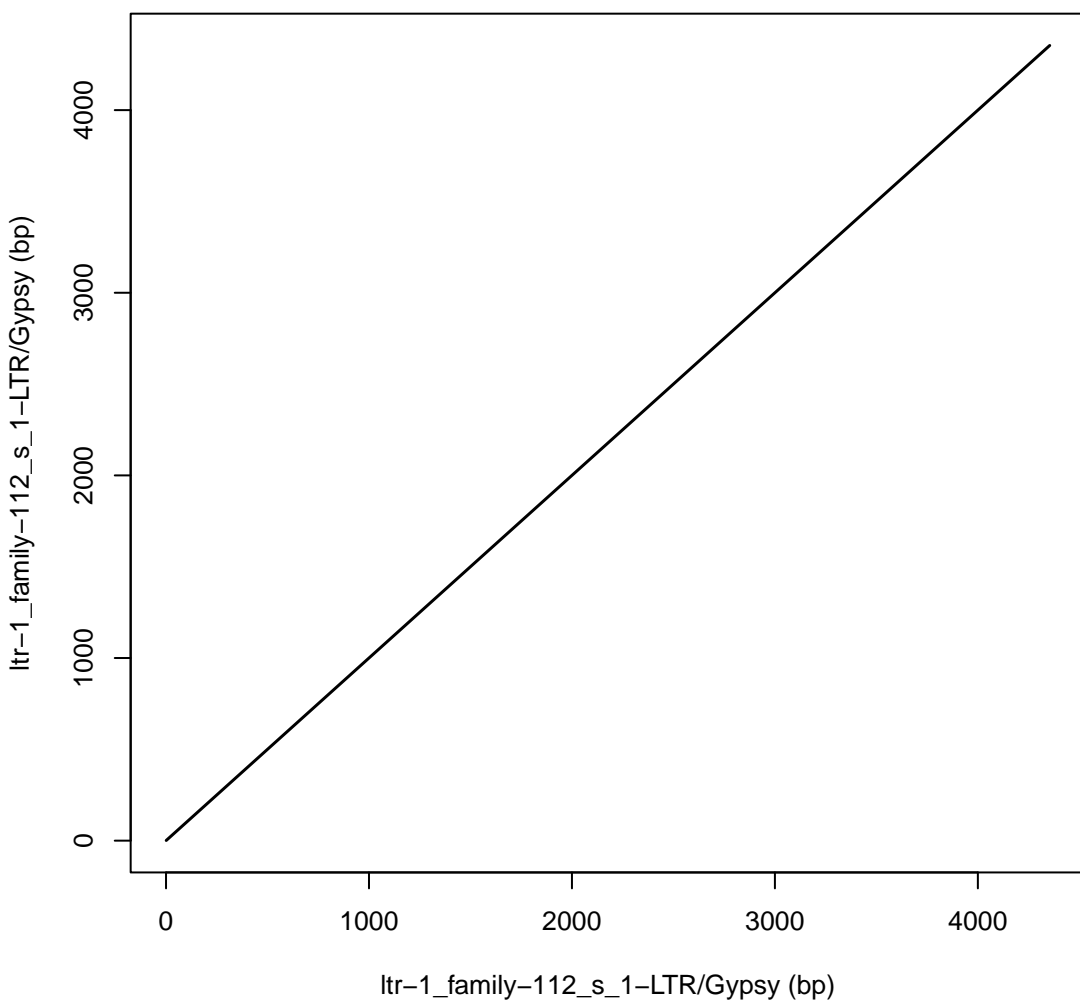
TE: ltr-1\_family-112\_s\_1-LTR/Gypsy  
consensus size: 4354bp; fragments: 85; full length: 4 (>=3918.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

