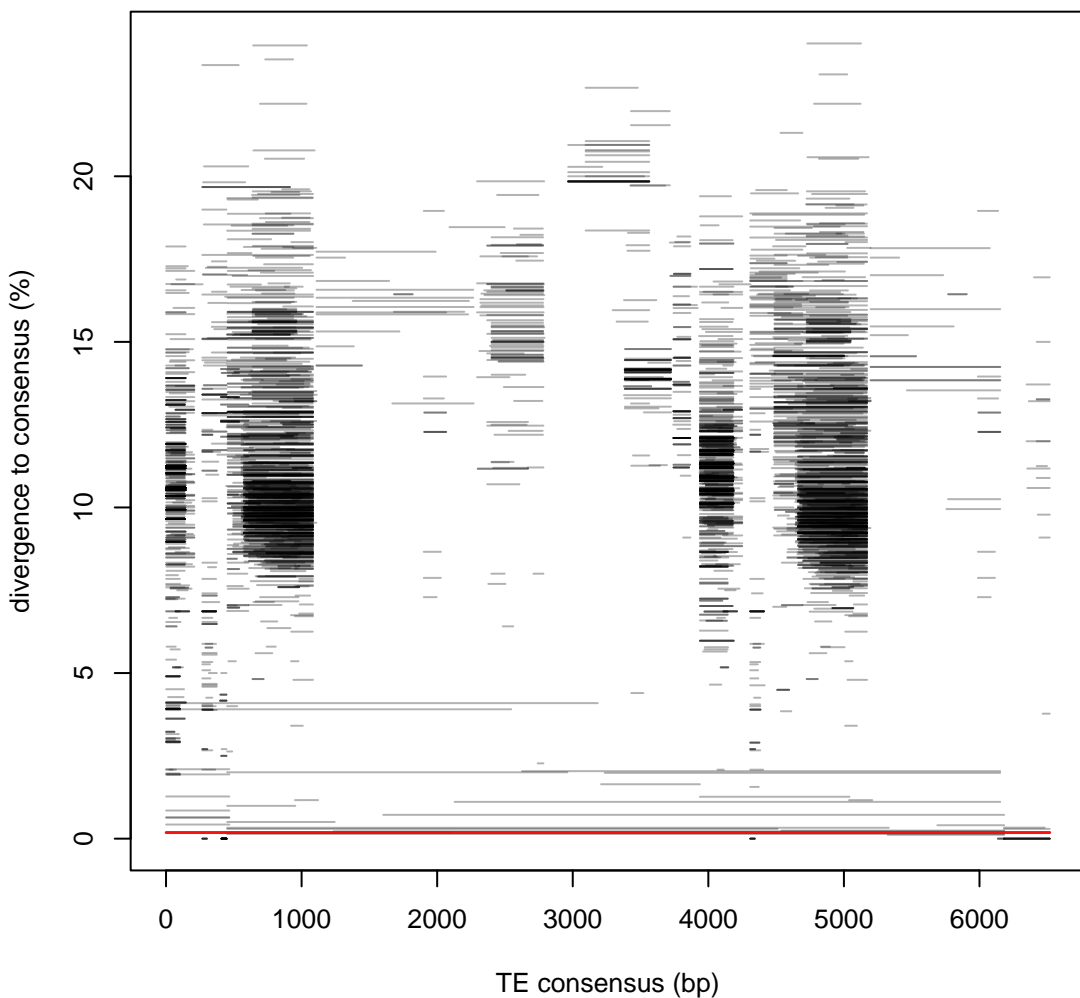
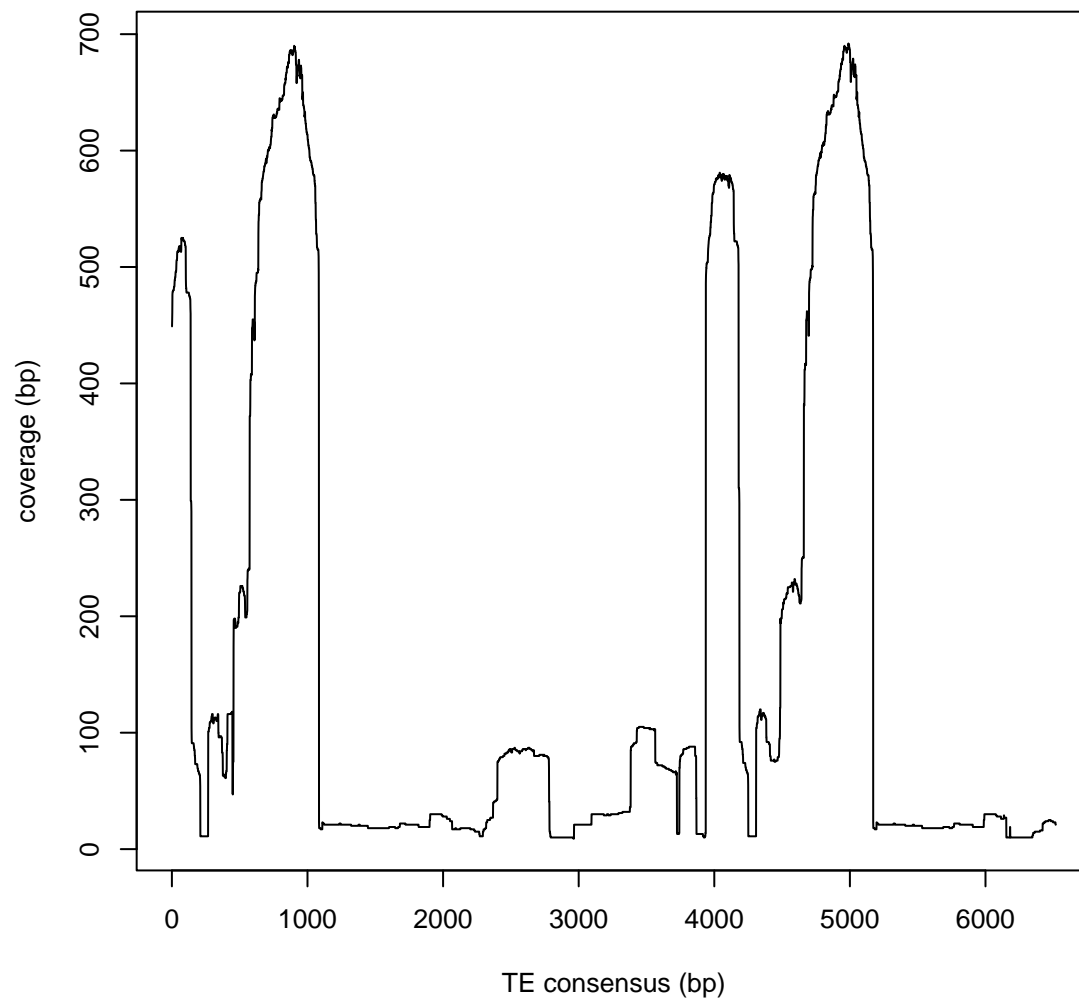


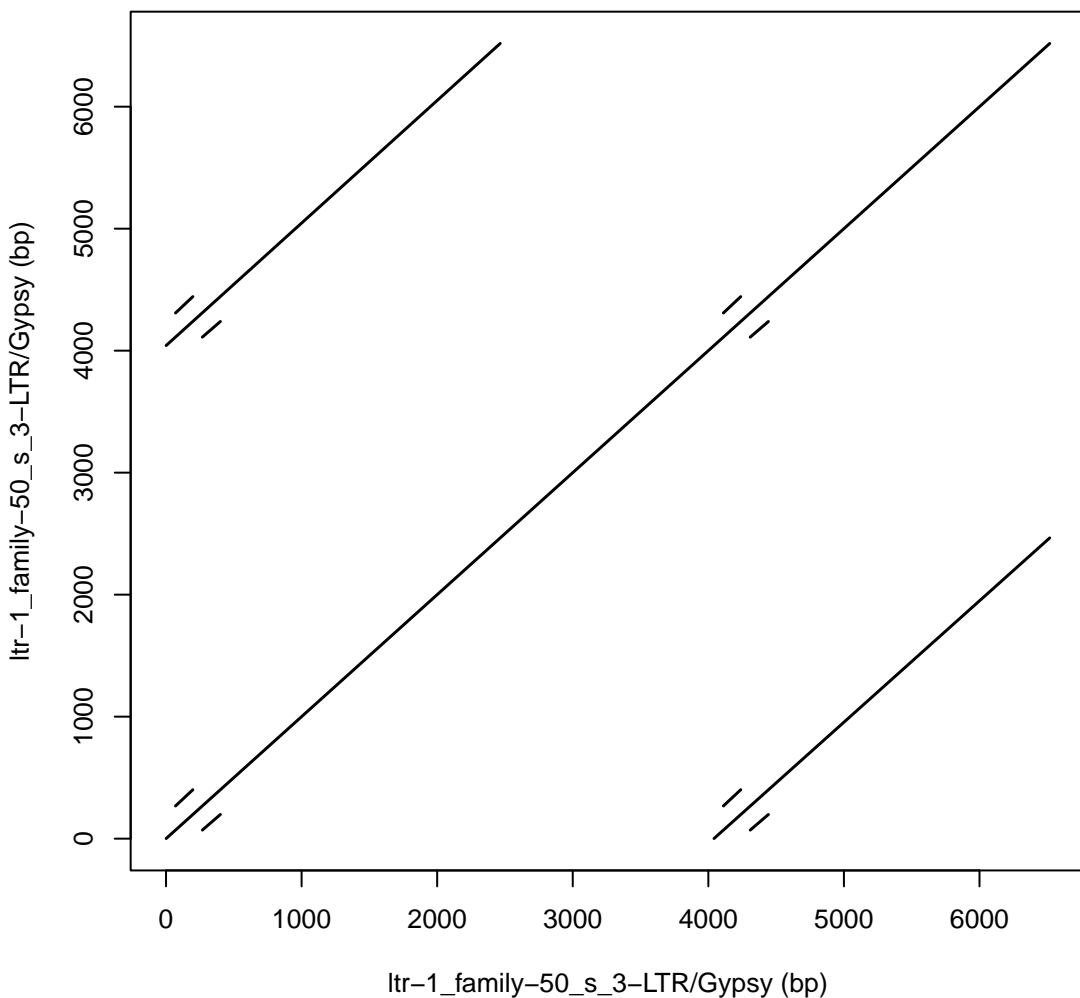
TE: ltr-1\_family-50\_s\_3-LTR/Gypsy  
consensus size: 6518bp; fragments: 3410; full length: 1 (>=5866.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

