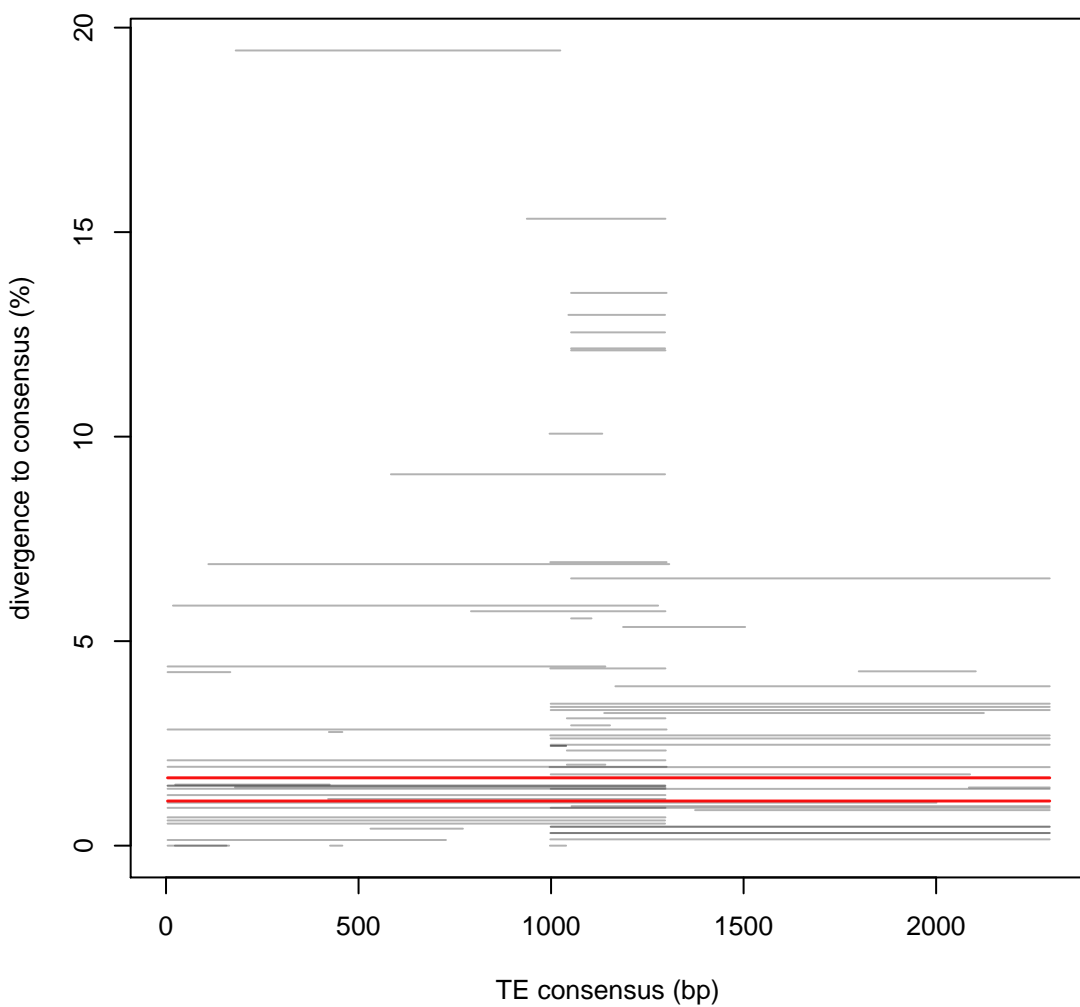
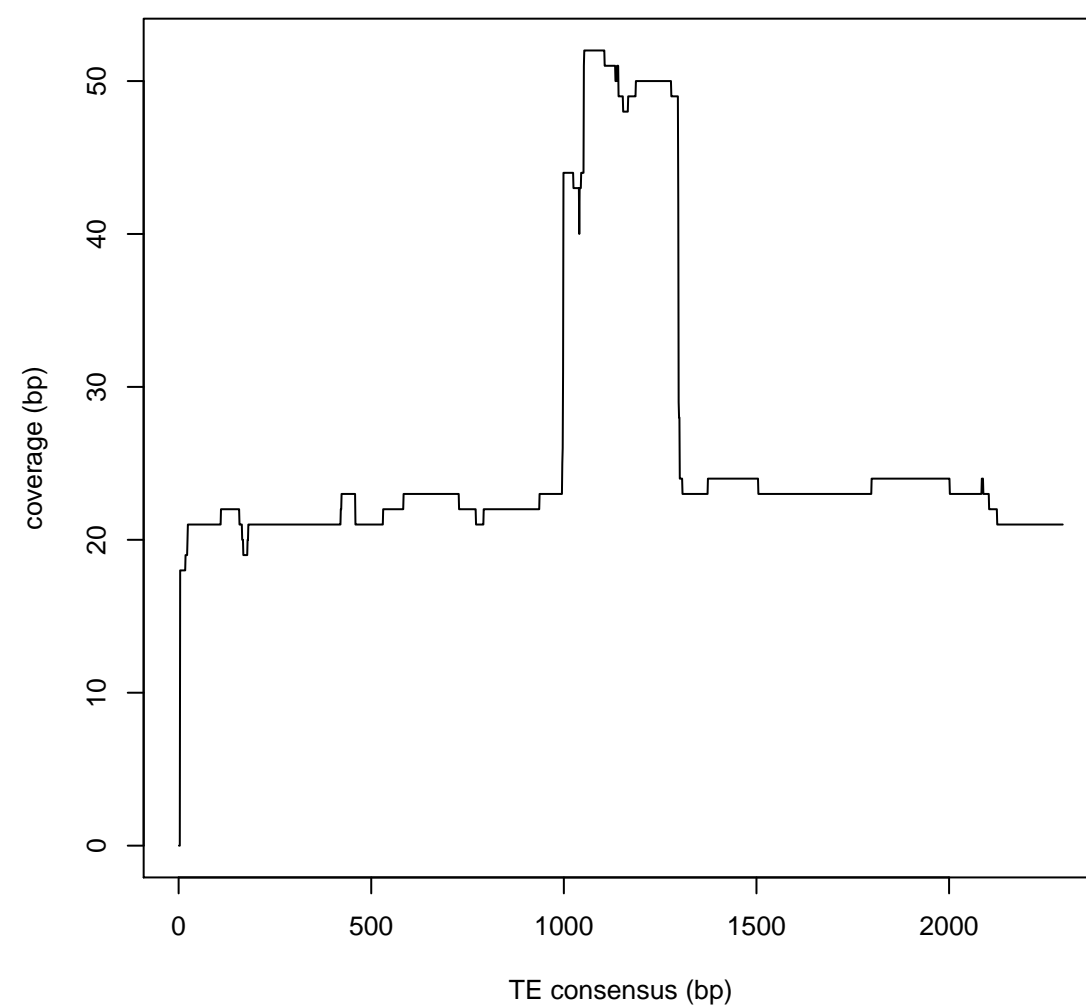


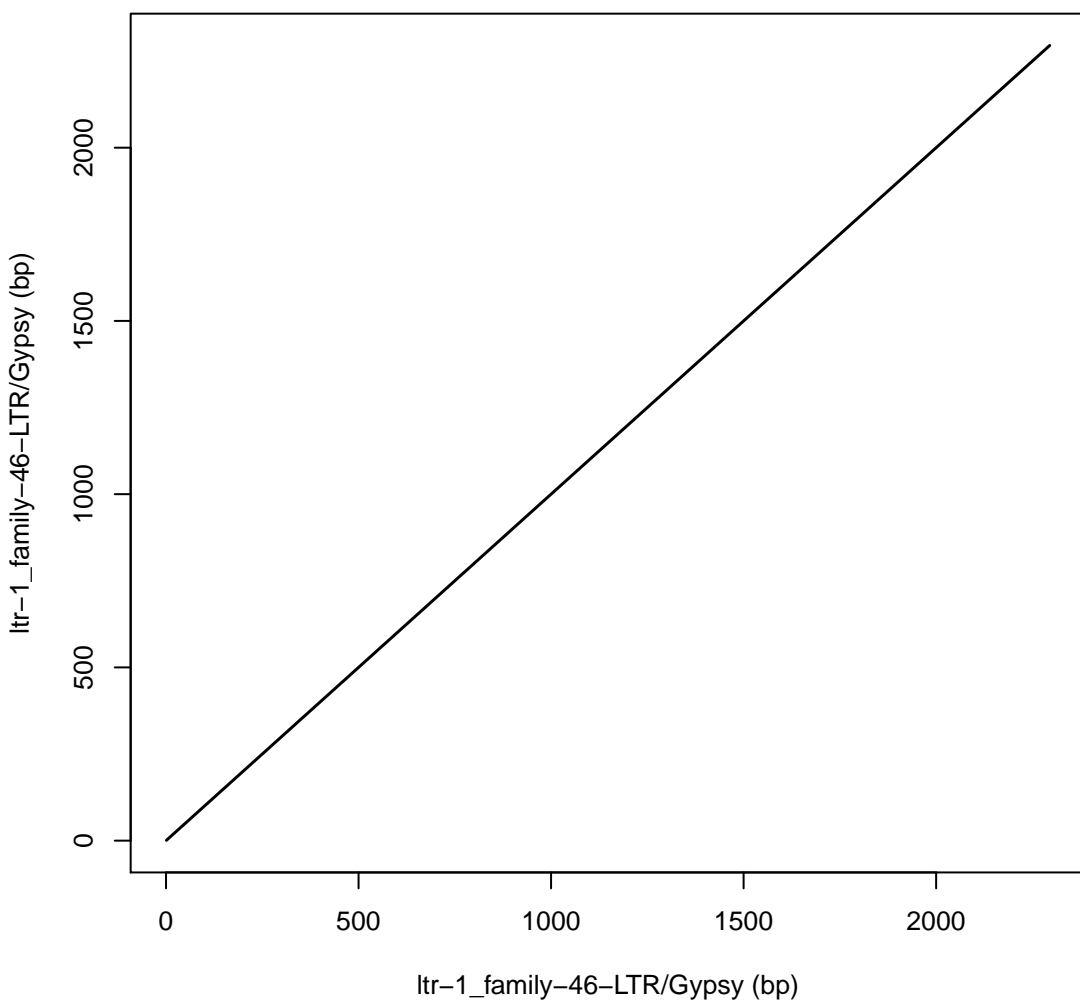
TE: ltr-1\_family-46-LTR/Gypsy  
consensus size: 2295bp; fragments: 70; full length: 2 ( $\geq 2065.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

