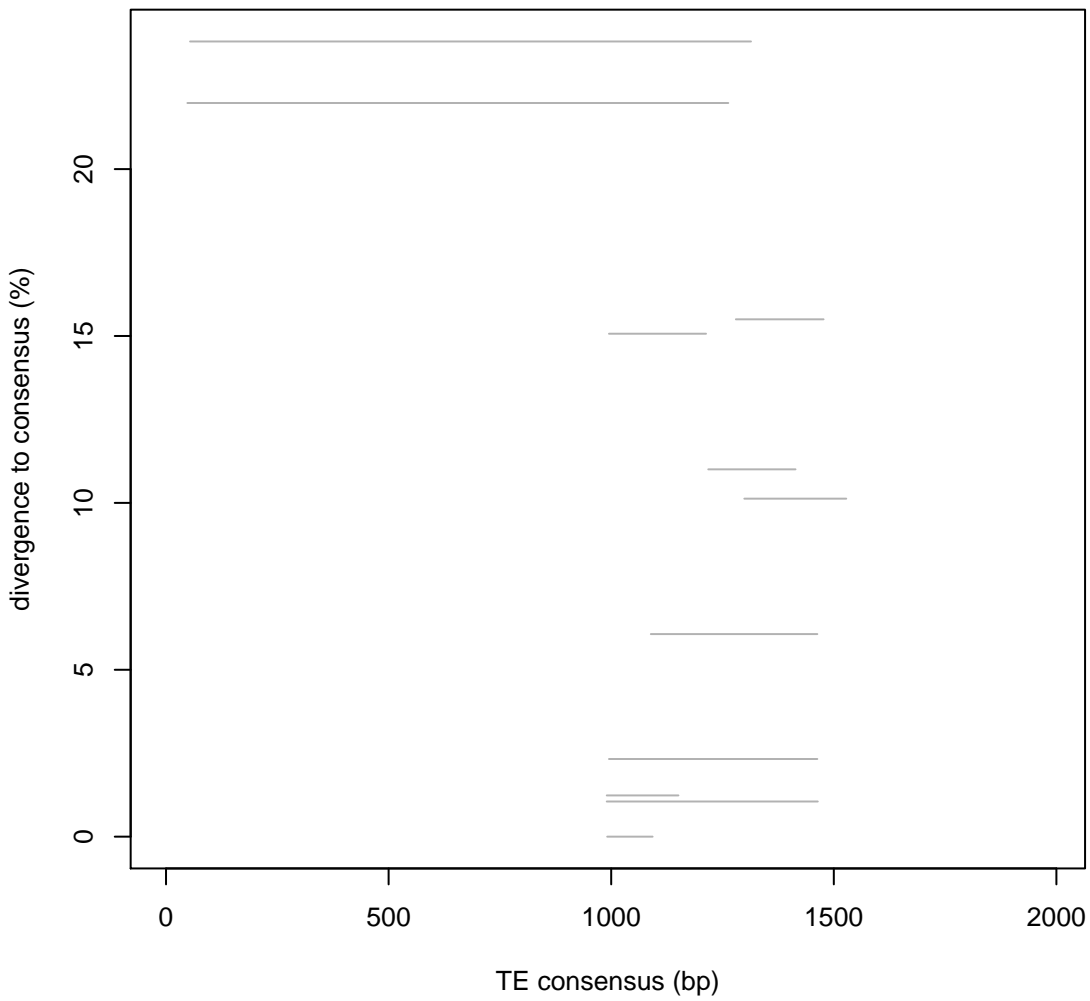
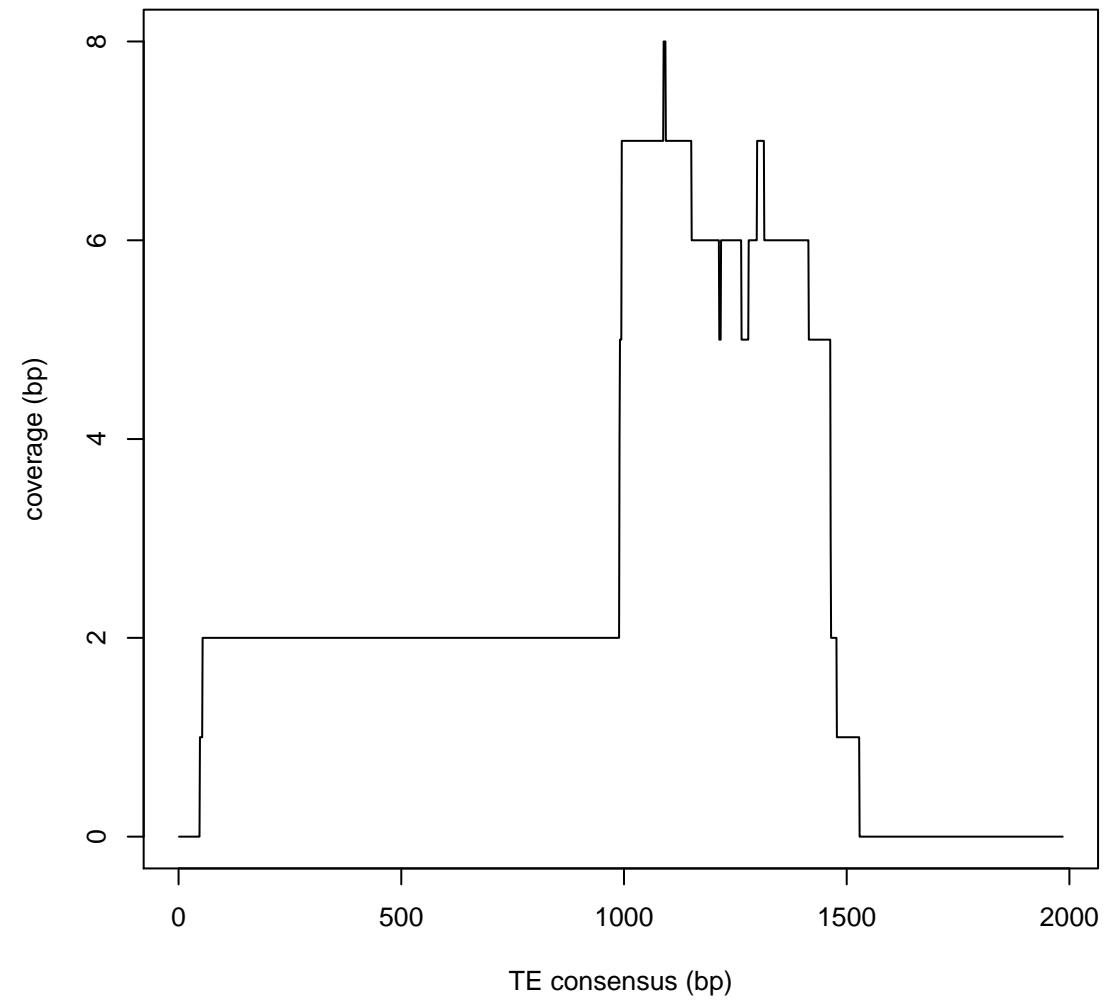


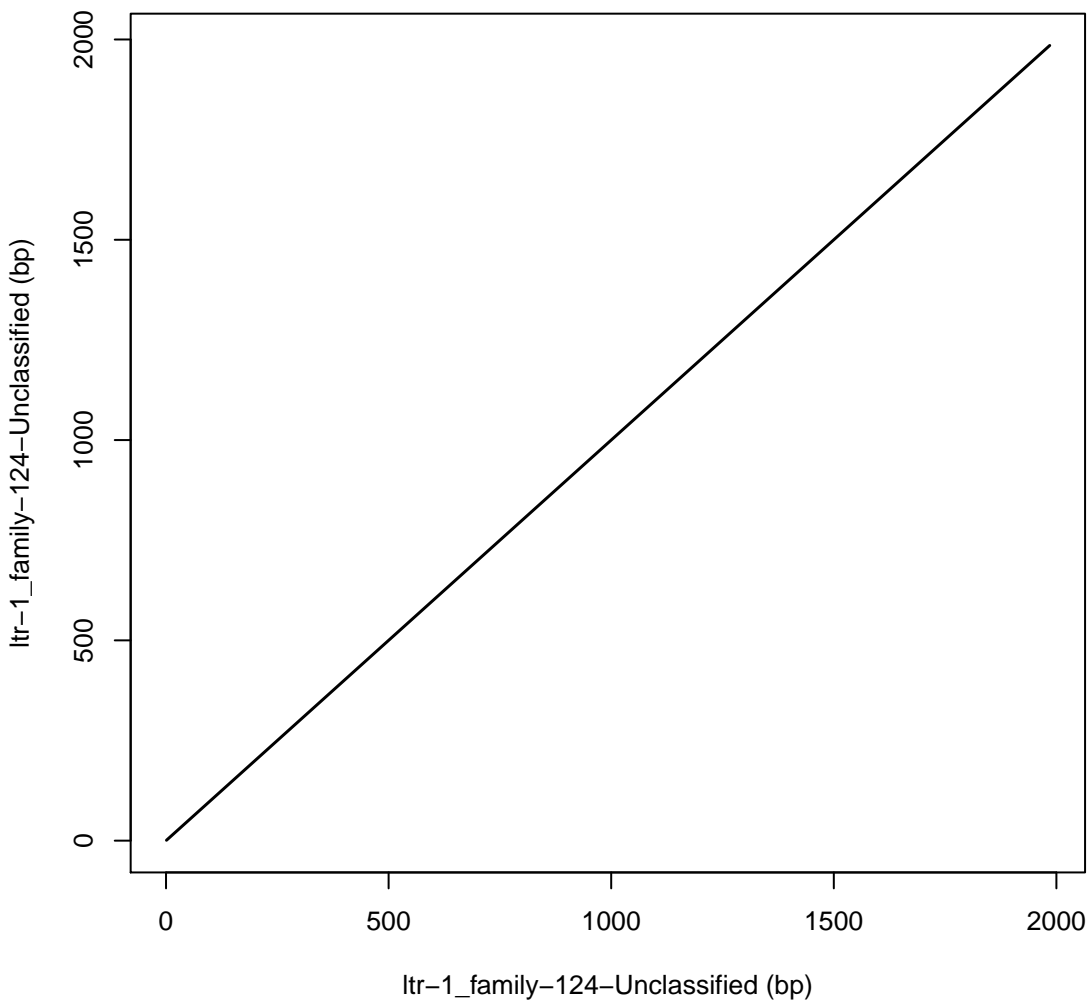
TE: ltr-1_family-124-Unclassified
consensus size: 1985bp; fragments: 11; full length: 0 (>=1786.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

