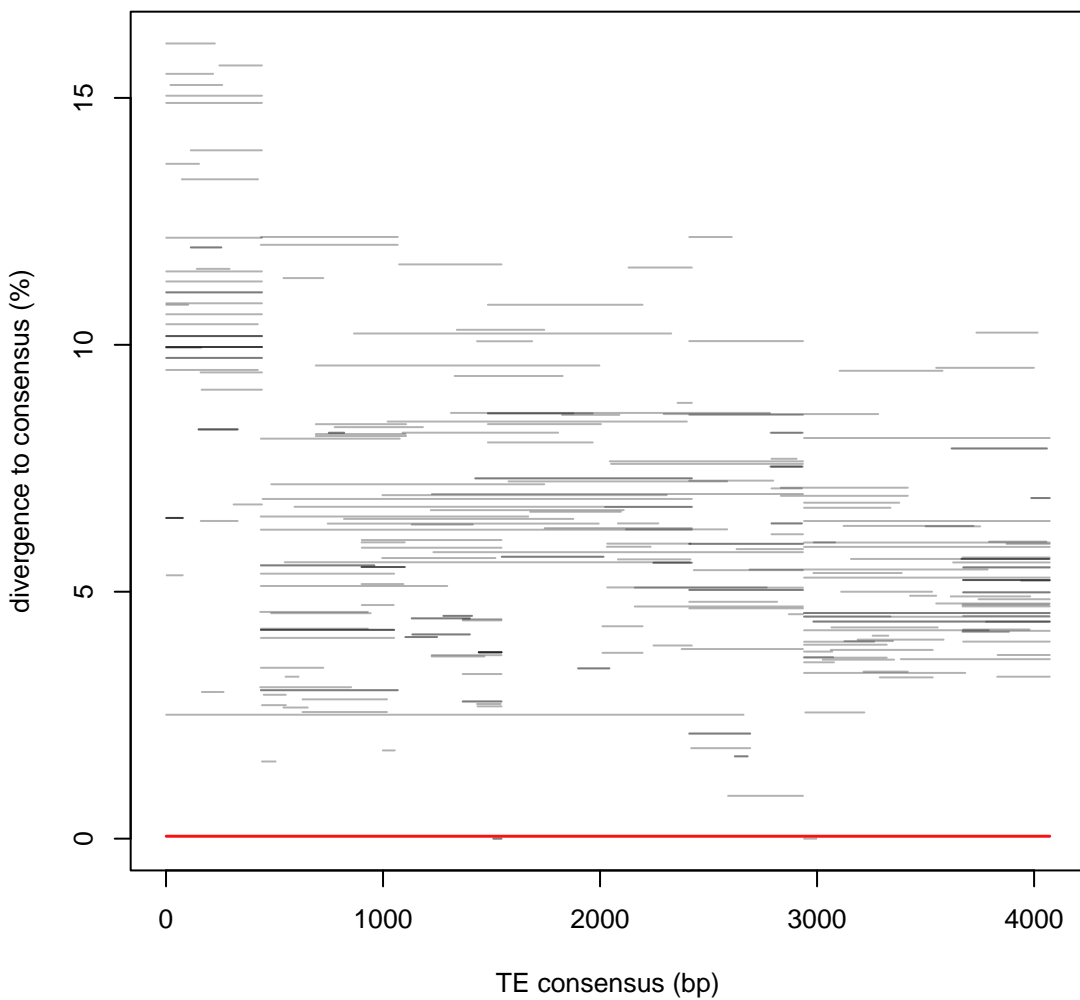
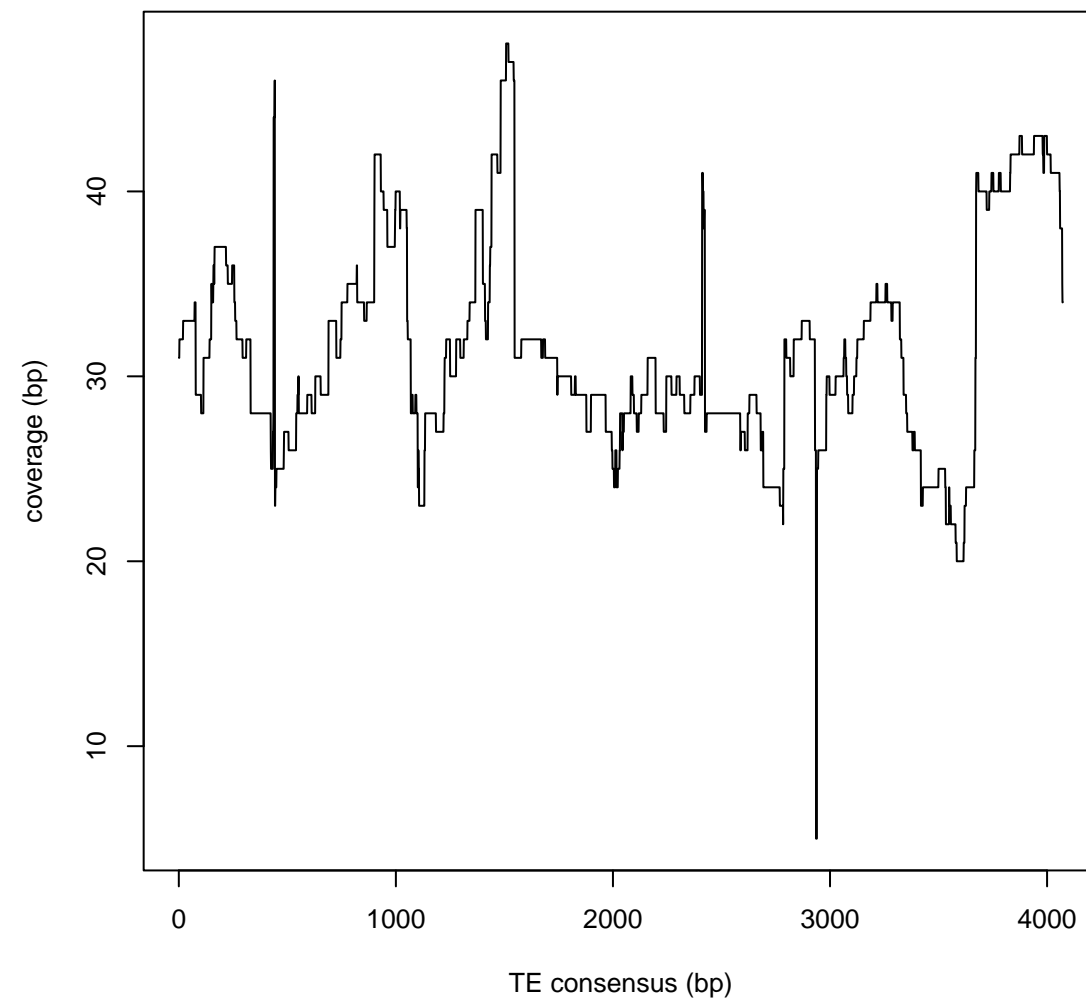


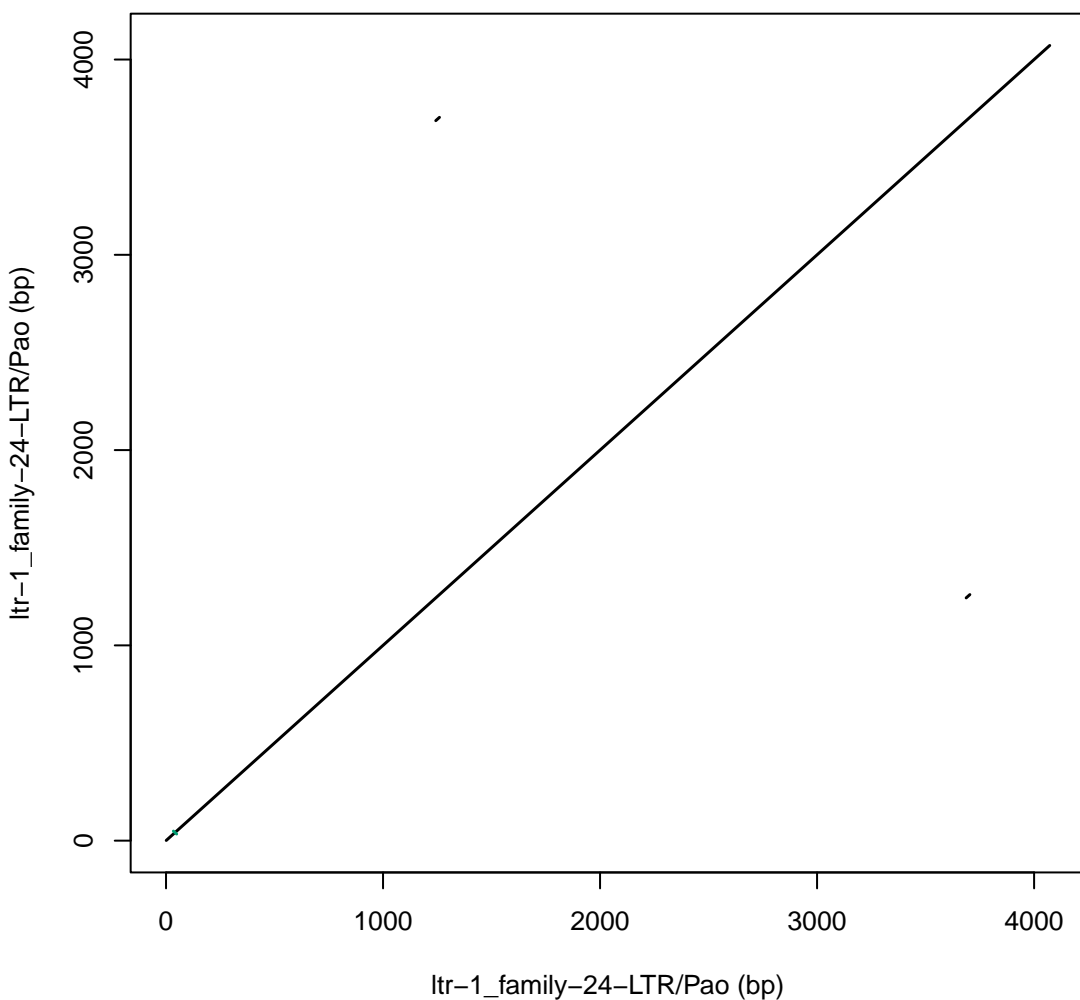
TE: ltr-1_family-24-LTR/Pao
consensus size: 4072bp; fragments: 275; full length: 1 (>=3664.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

