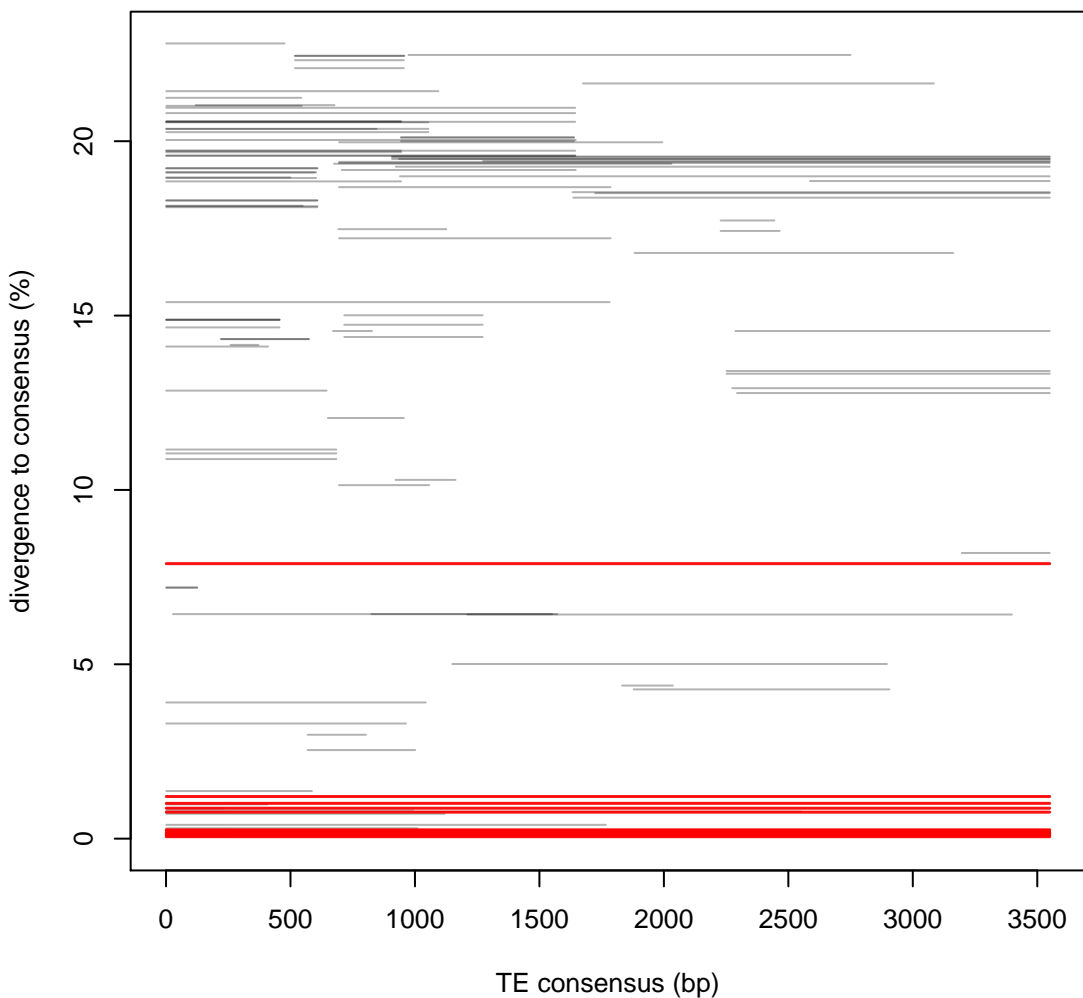
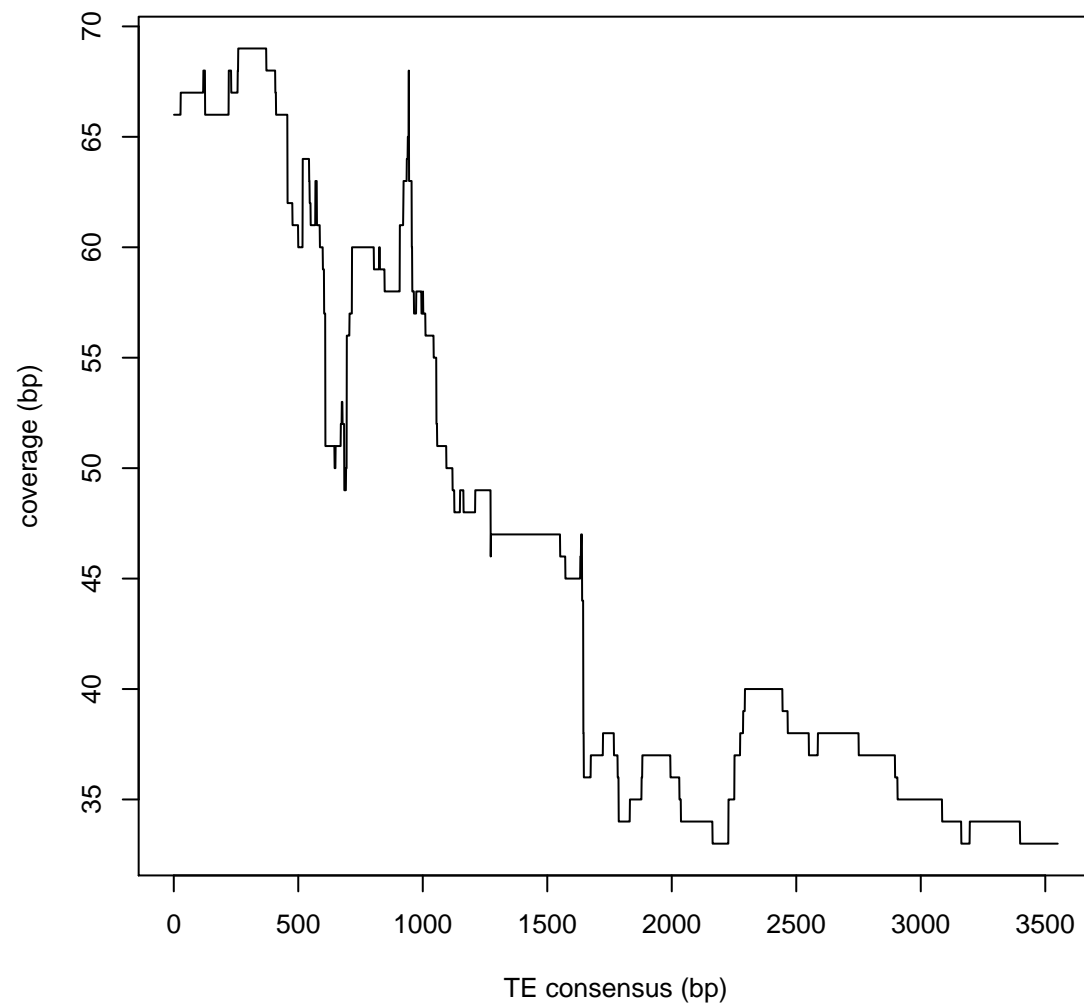


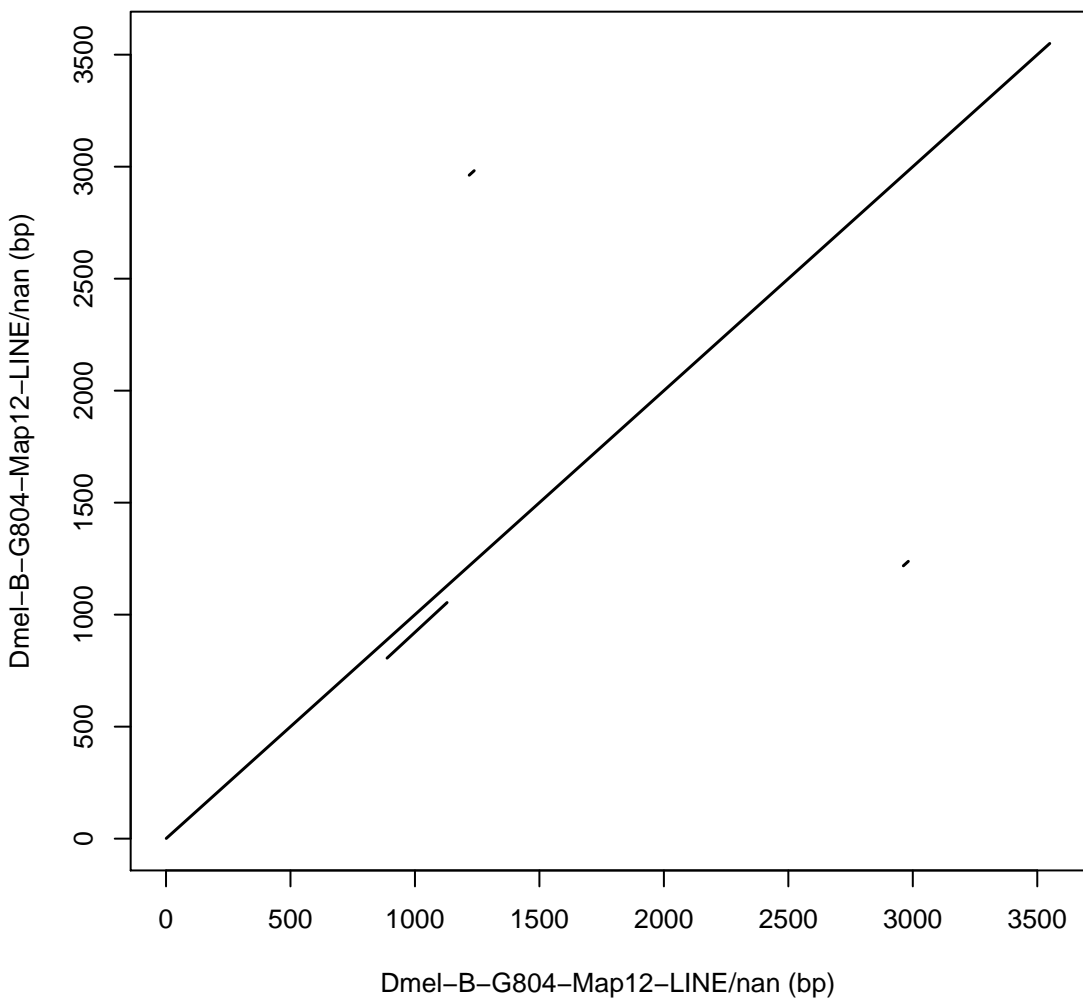
TE: Dmel-B-G804-Map12-LINE/nan
consensus size: 3550bp; fragments: 124; full length: 14 (>=3195bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

