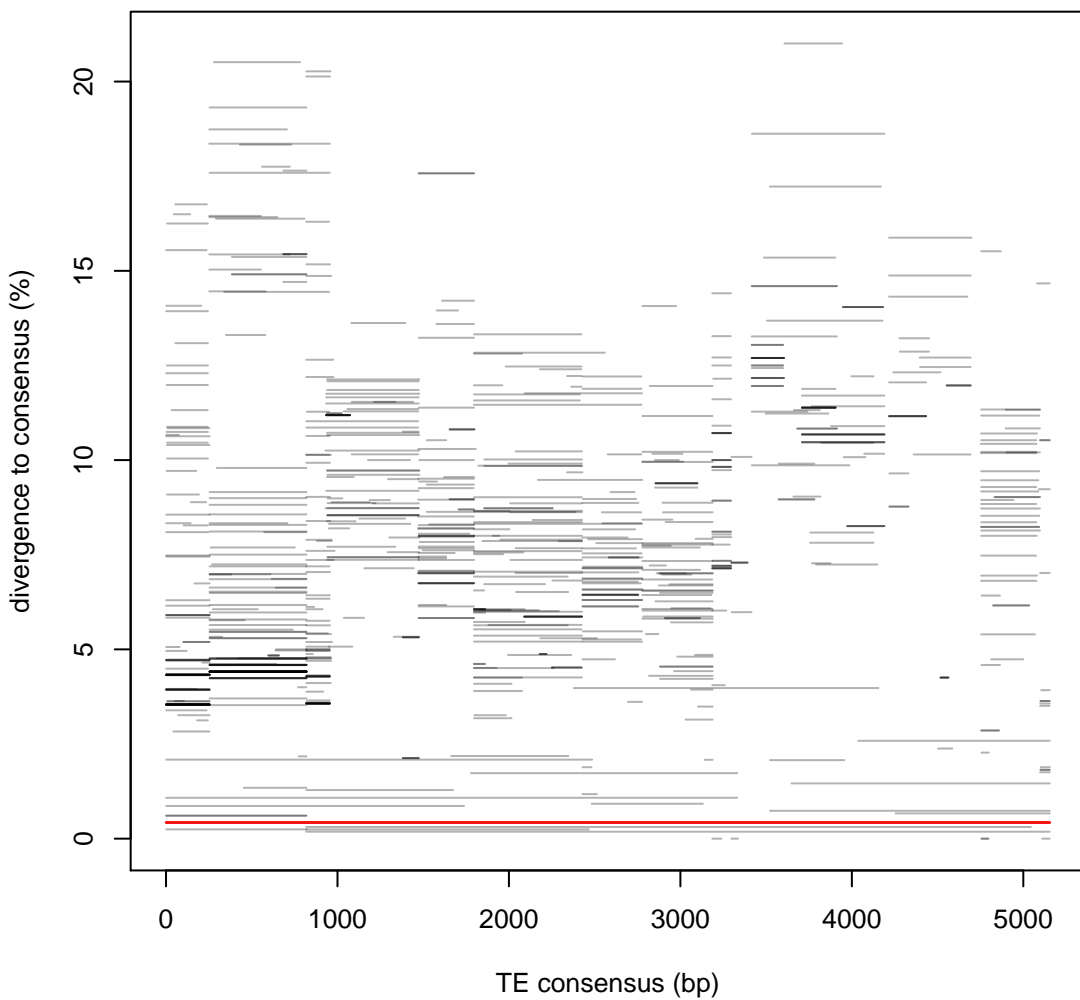
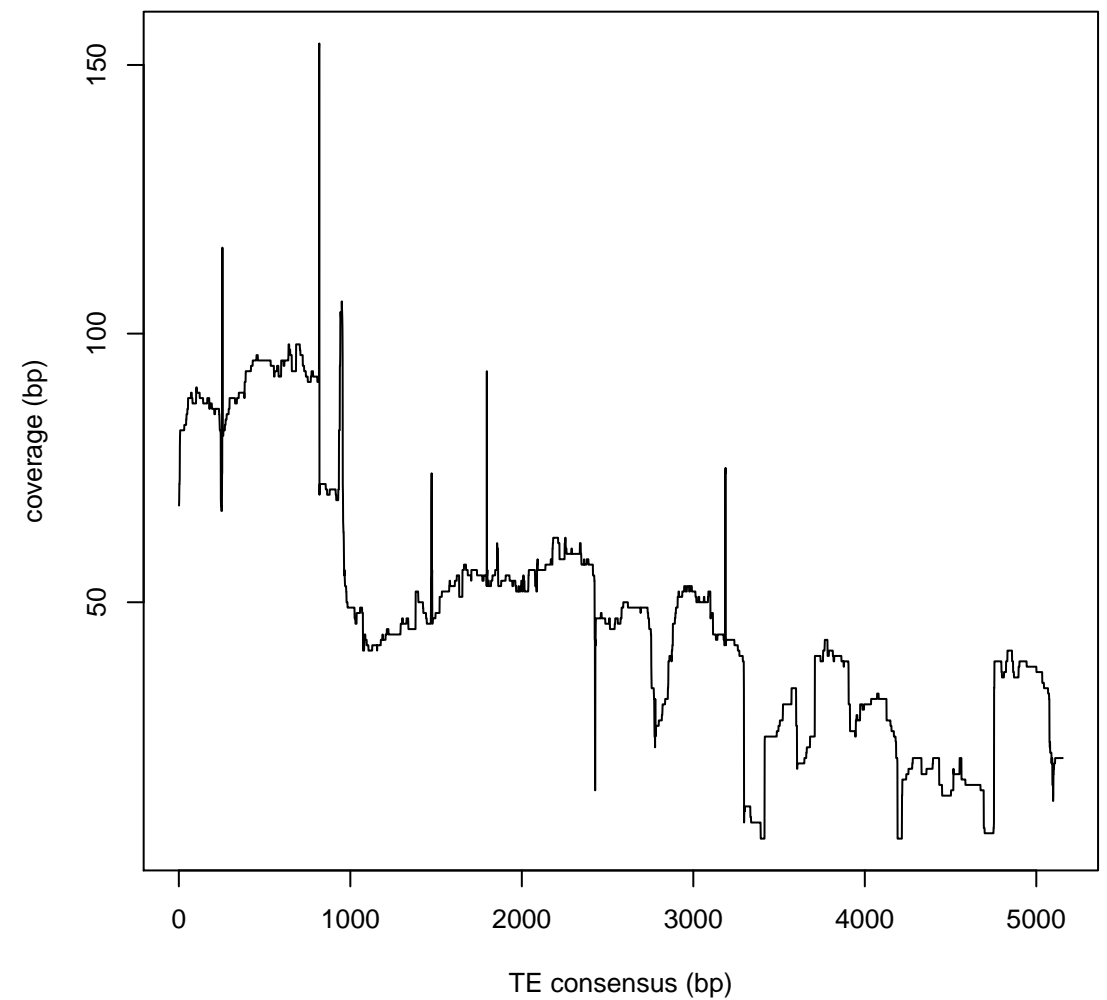


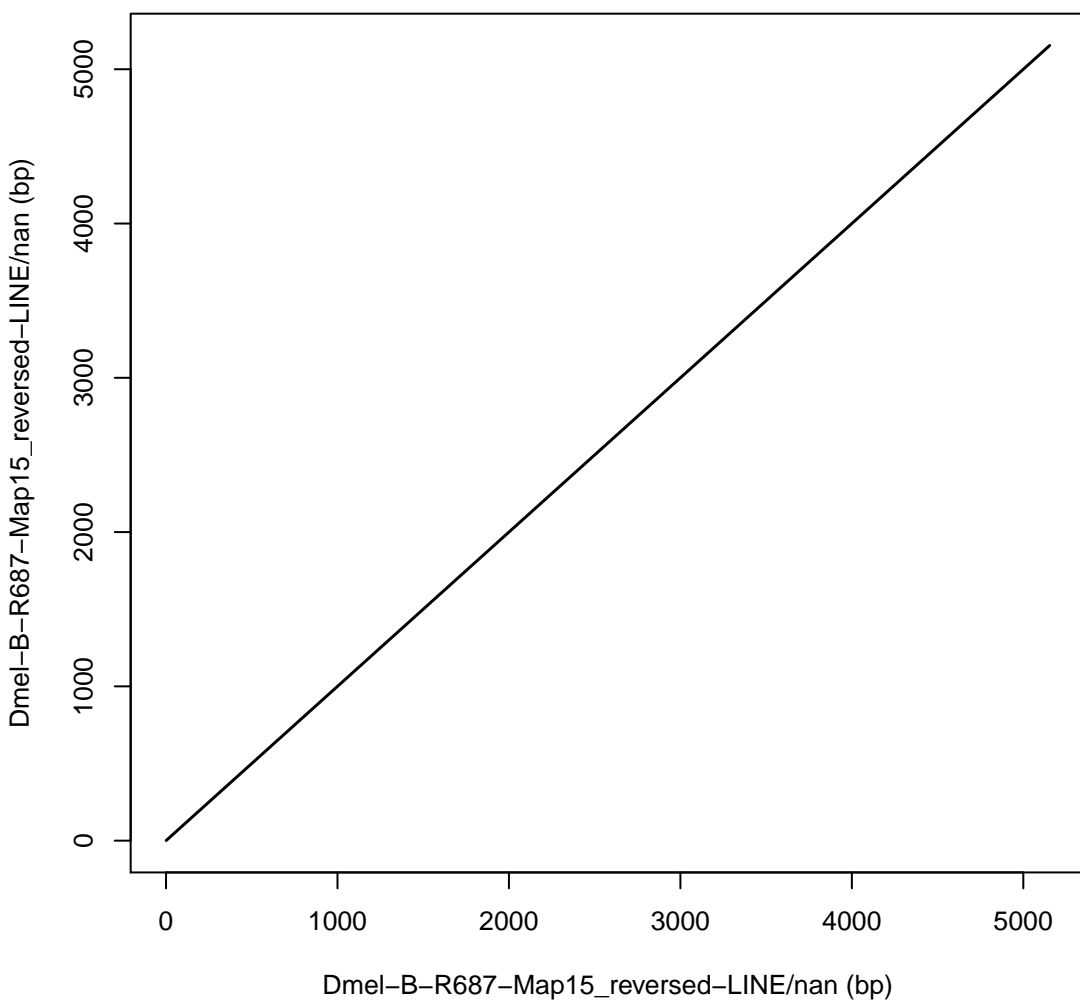
TE: Dmel-B-R687-Map15_reversed-LINE/nan
consensus size: 5154bp; fragments: 768; full length: 1 (≥ 4638.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

