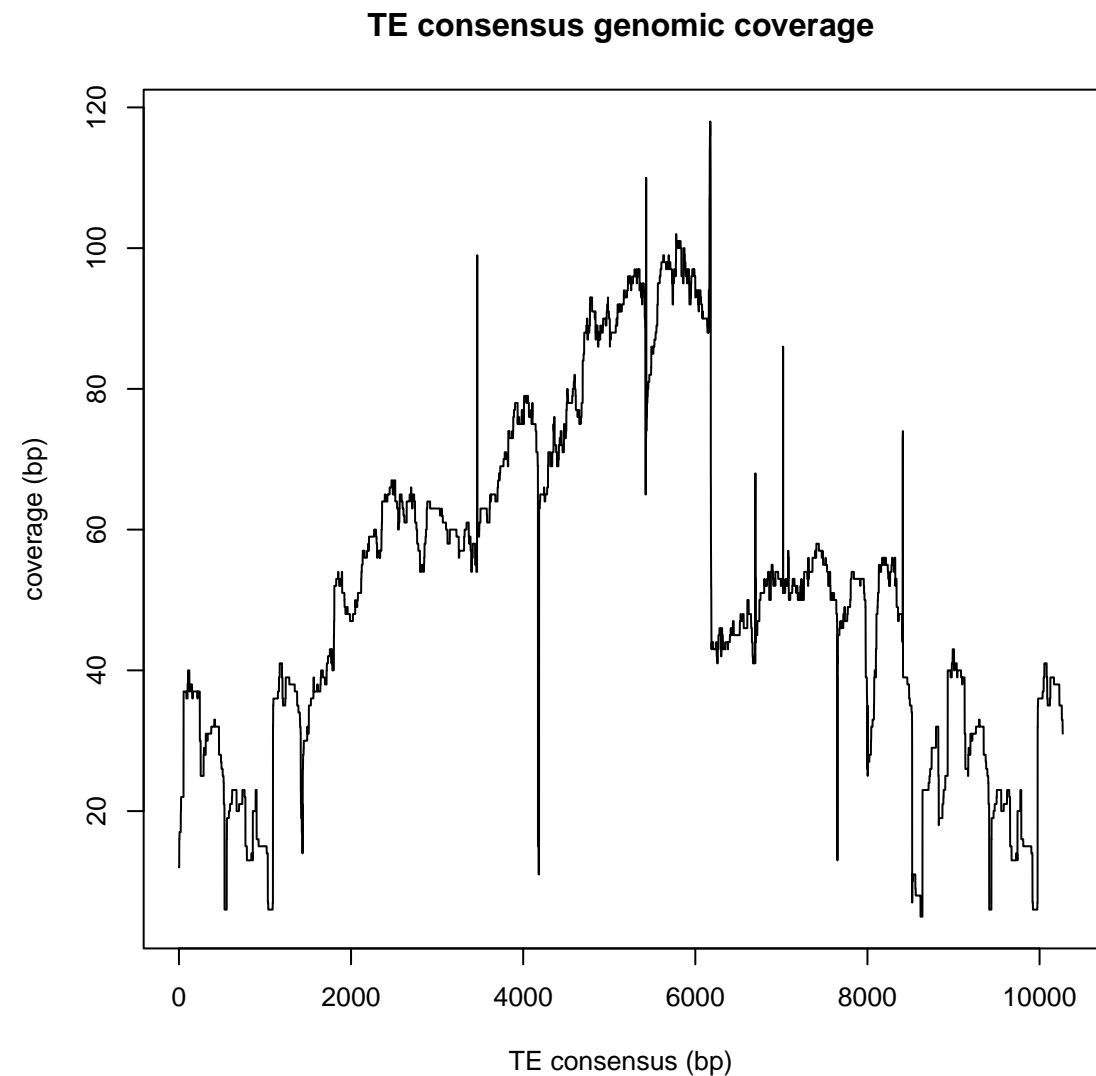
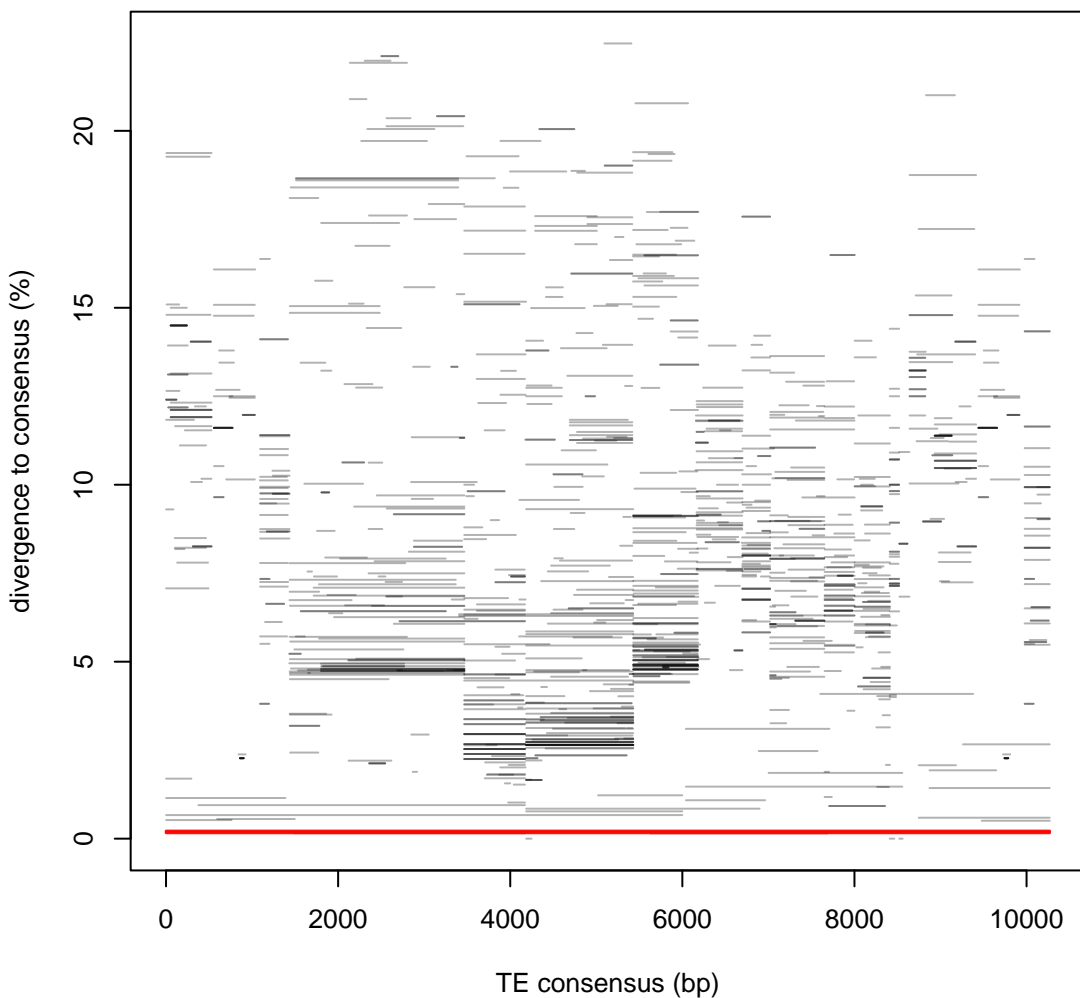
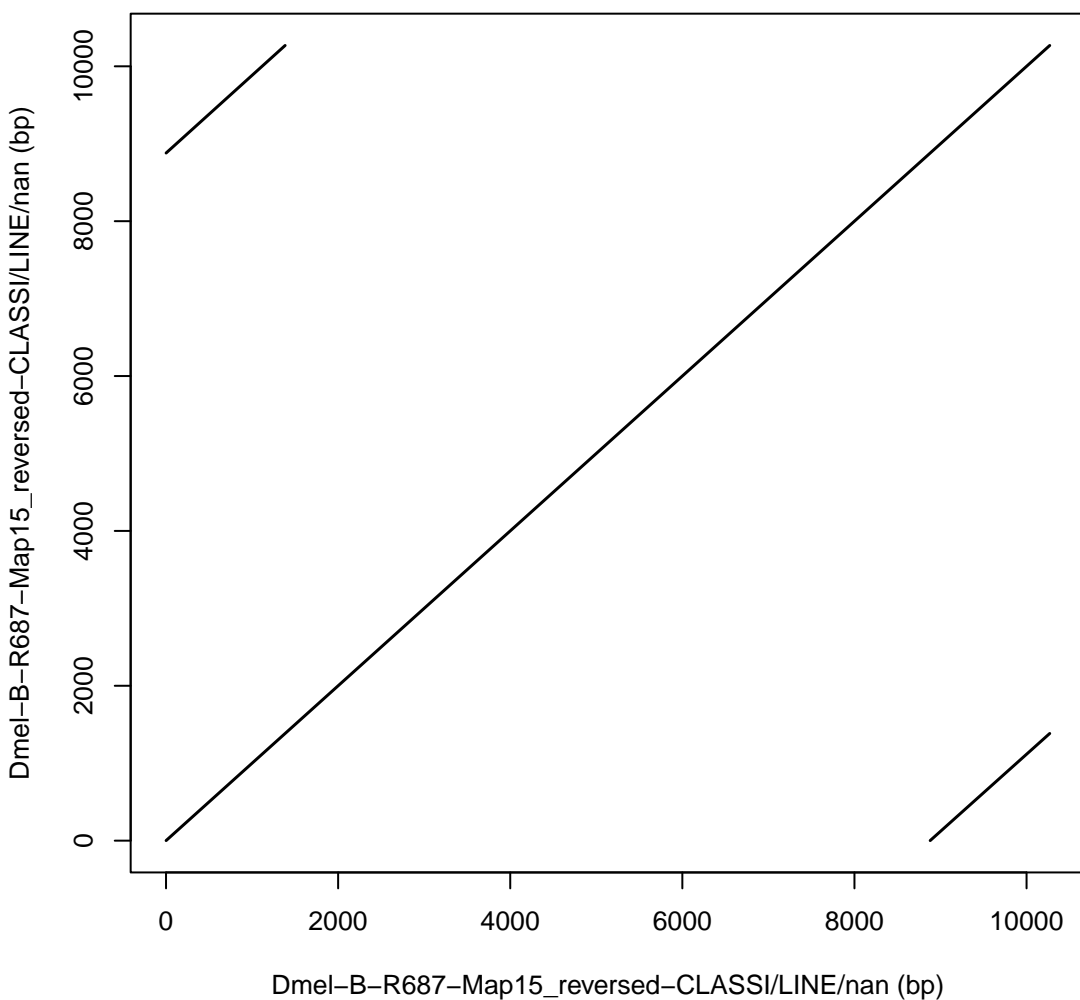


TE: Dmel-B-R687-Map15_reversed-CLASSI/LINE/nan
consensus size: 10269bp; fragments: 1119; full length: 2 (>=9242.1bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

