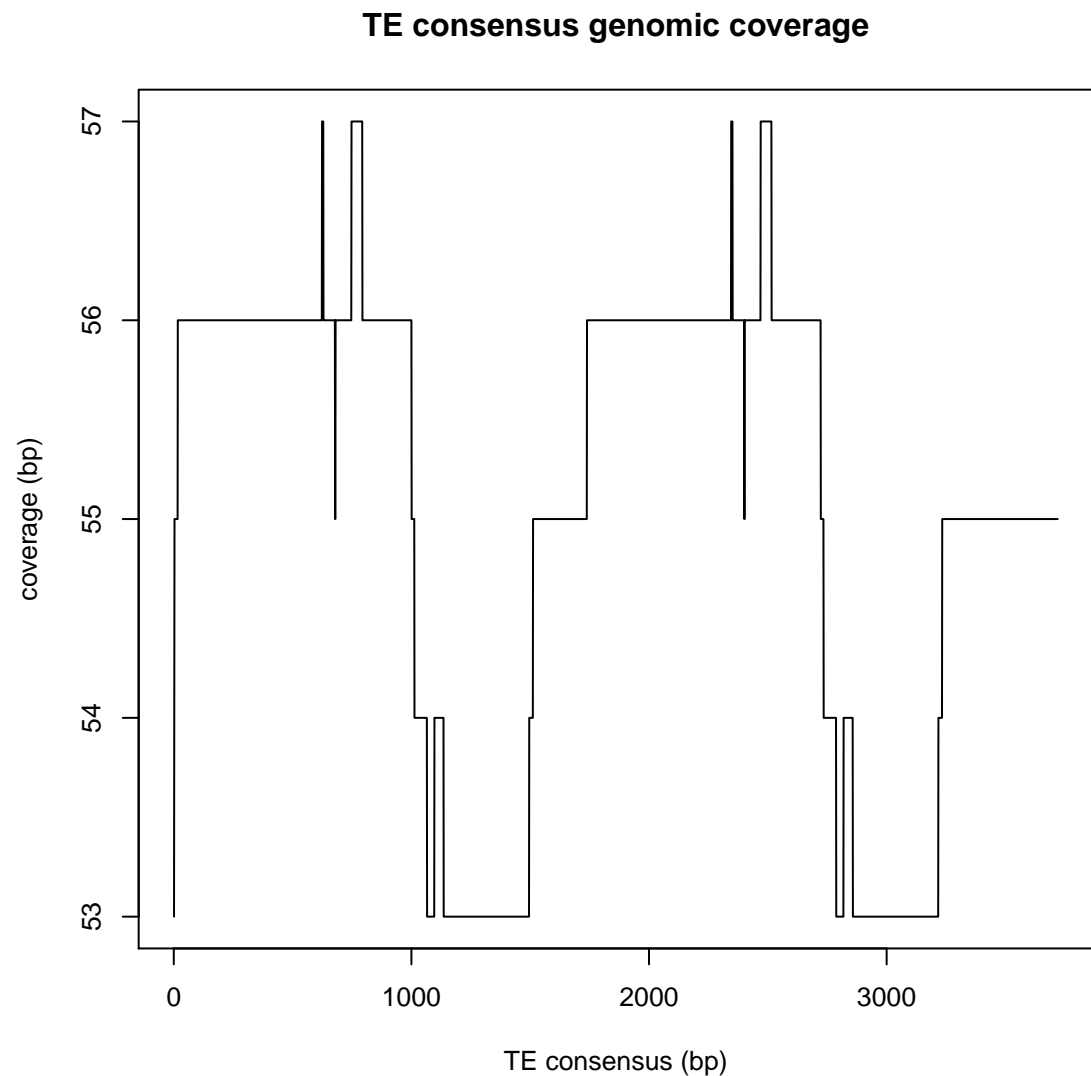
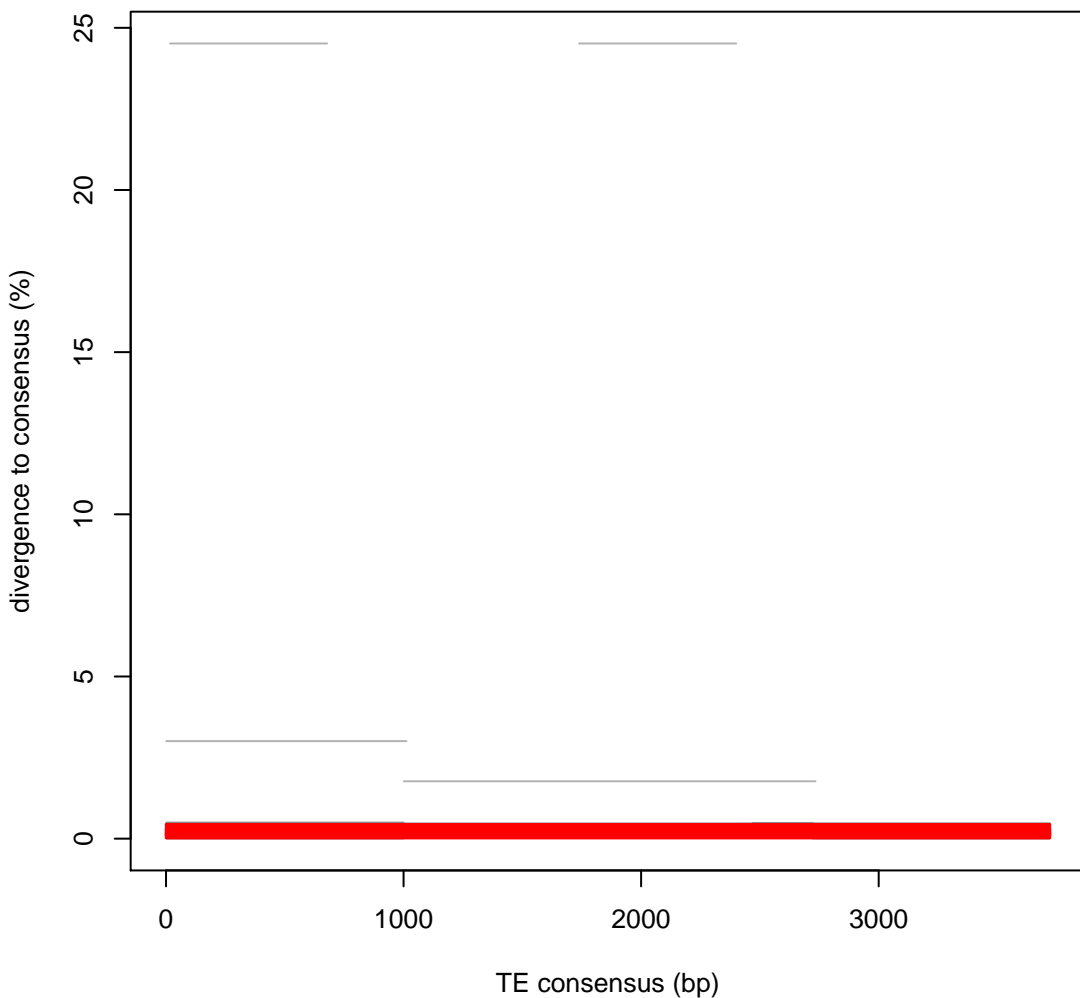
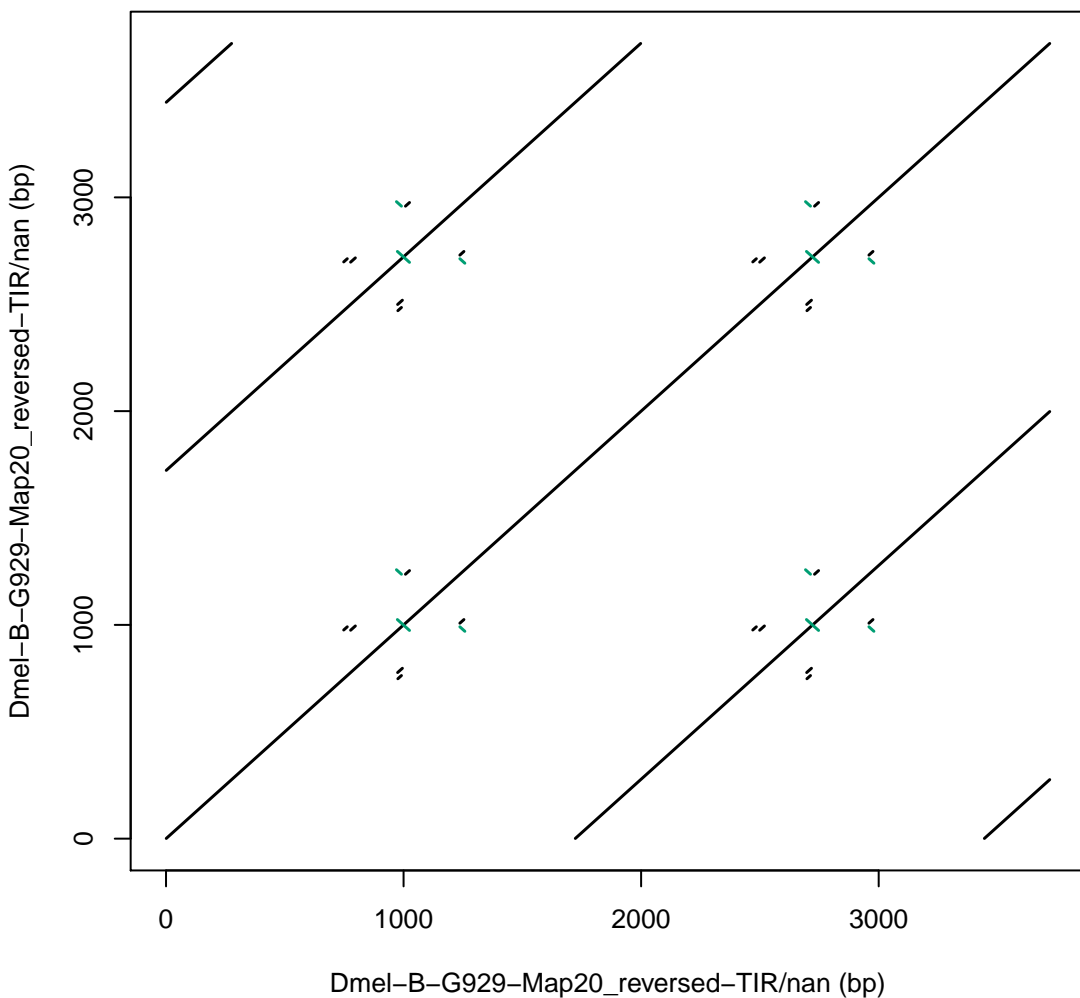


TE: Dmel-B-G929-Map20_reversed-TIR/nan
consensus size: 3720bp; fragments: 81; full length: 38 (>=3348bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

