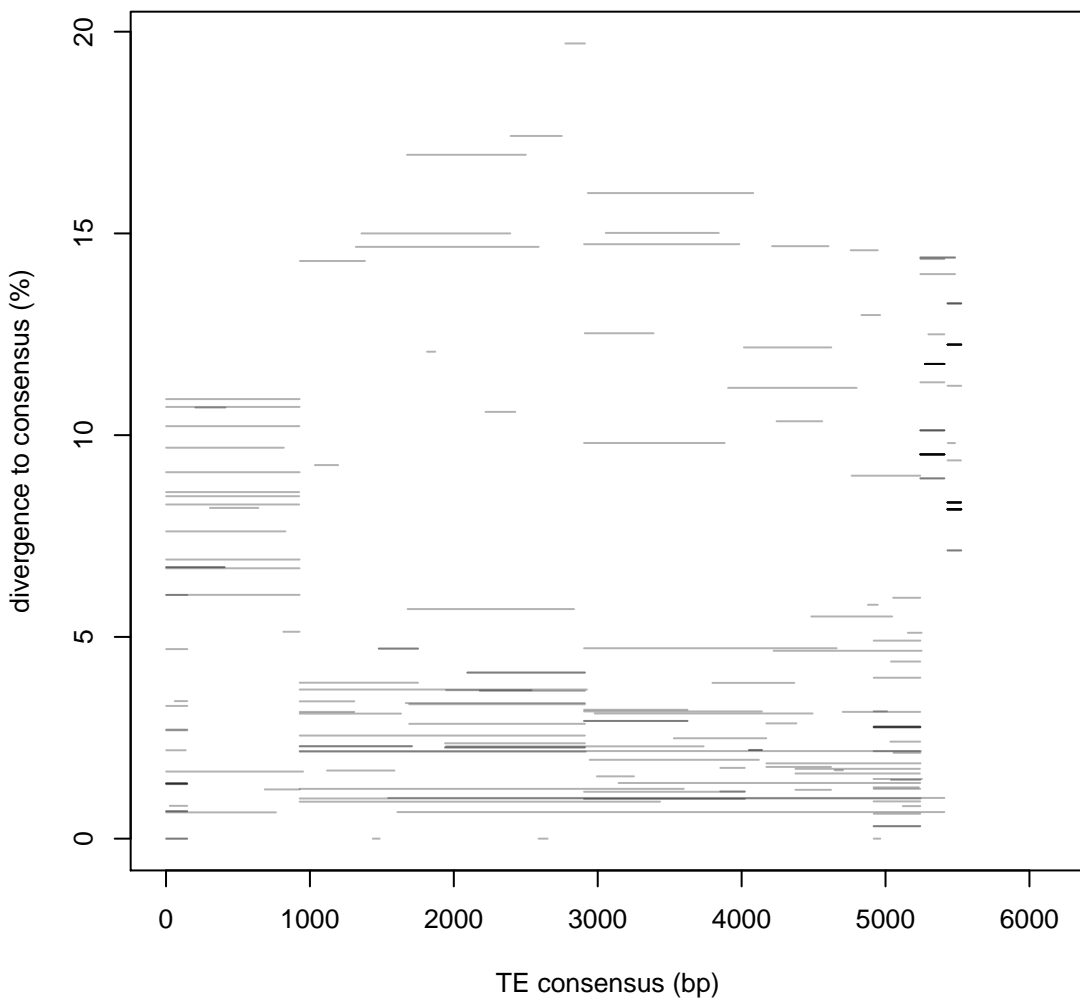
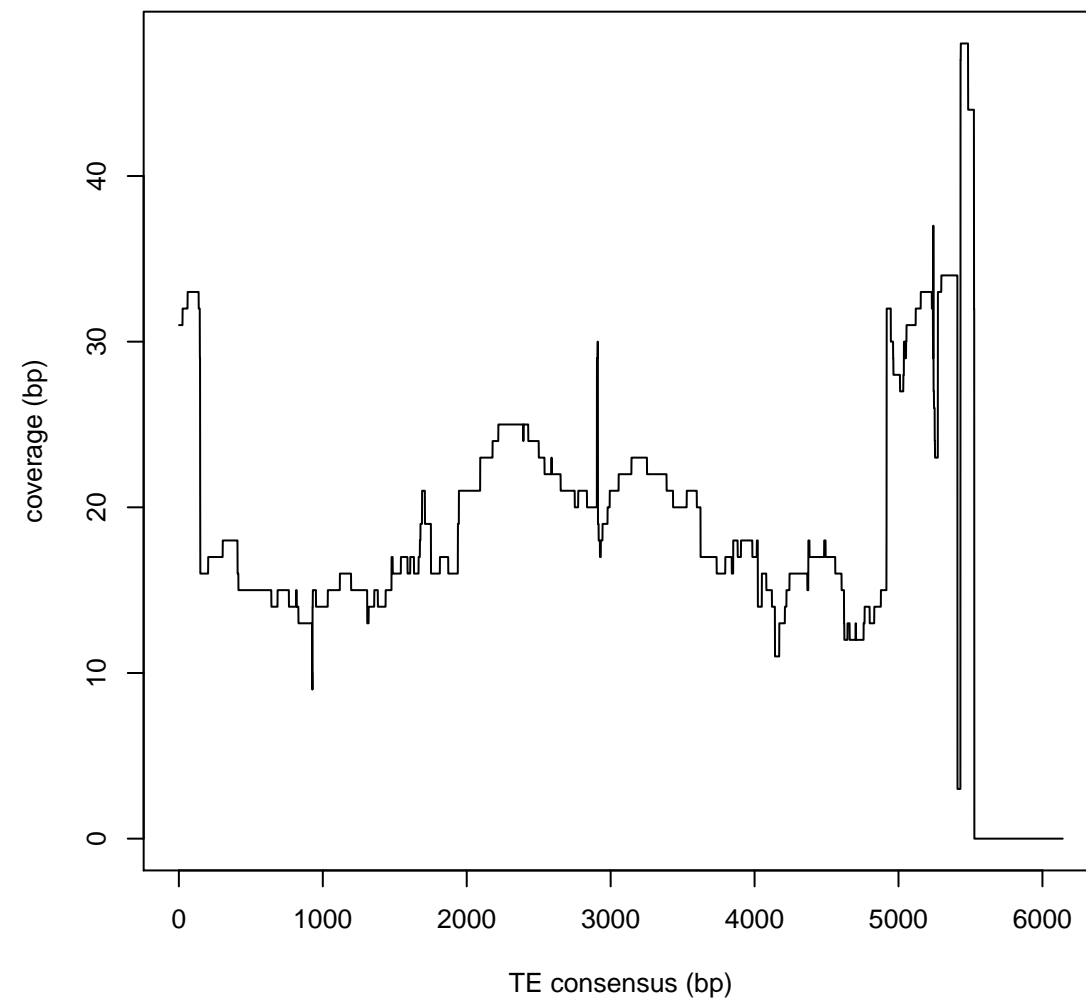


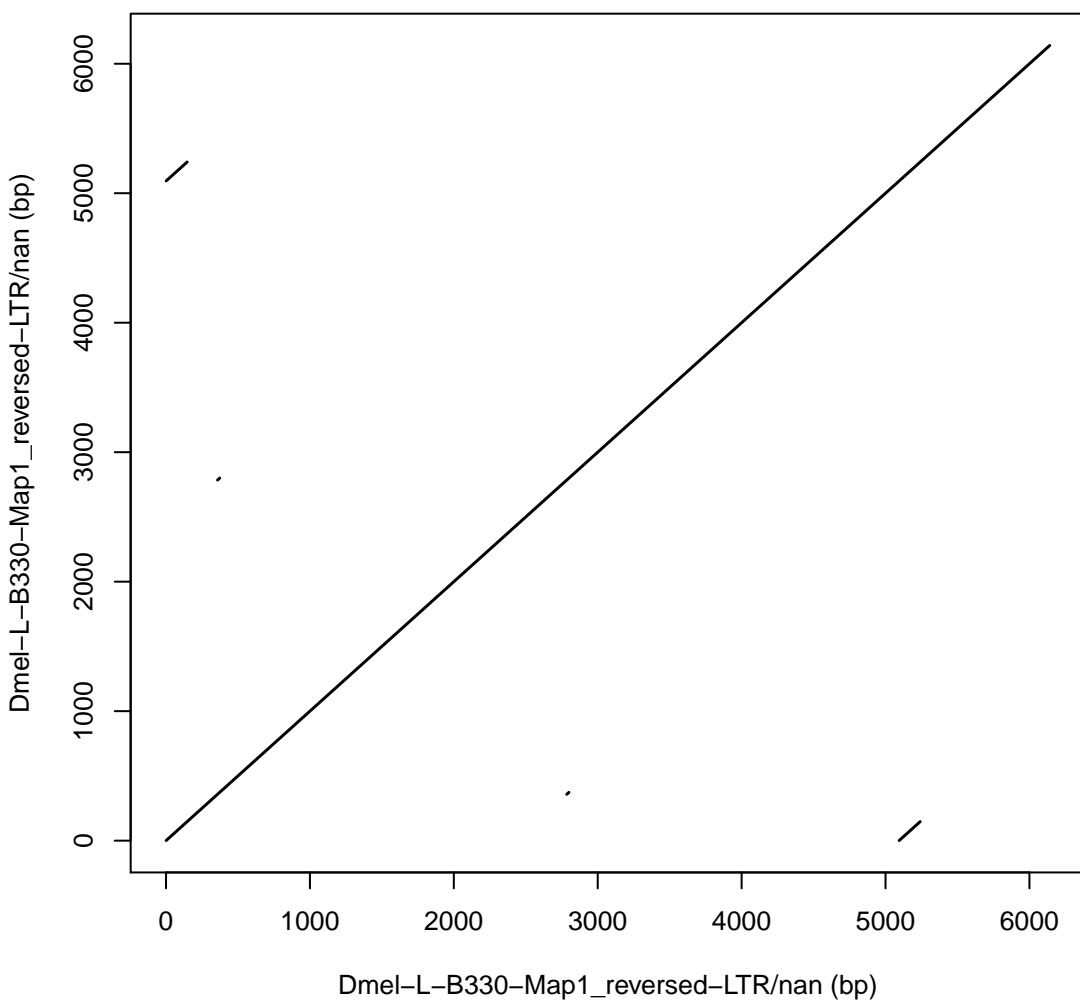
TE: Dmel-L-B330-Map1_reversed-LTR/nan
consensus size: 6141bp; fragments: 219; full length: 0 (>=5526.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

