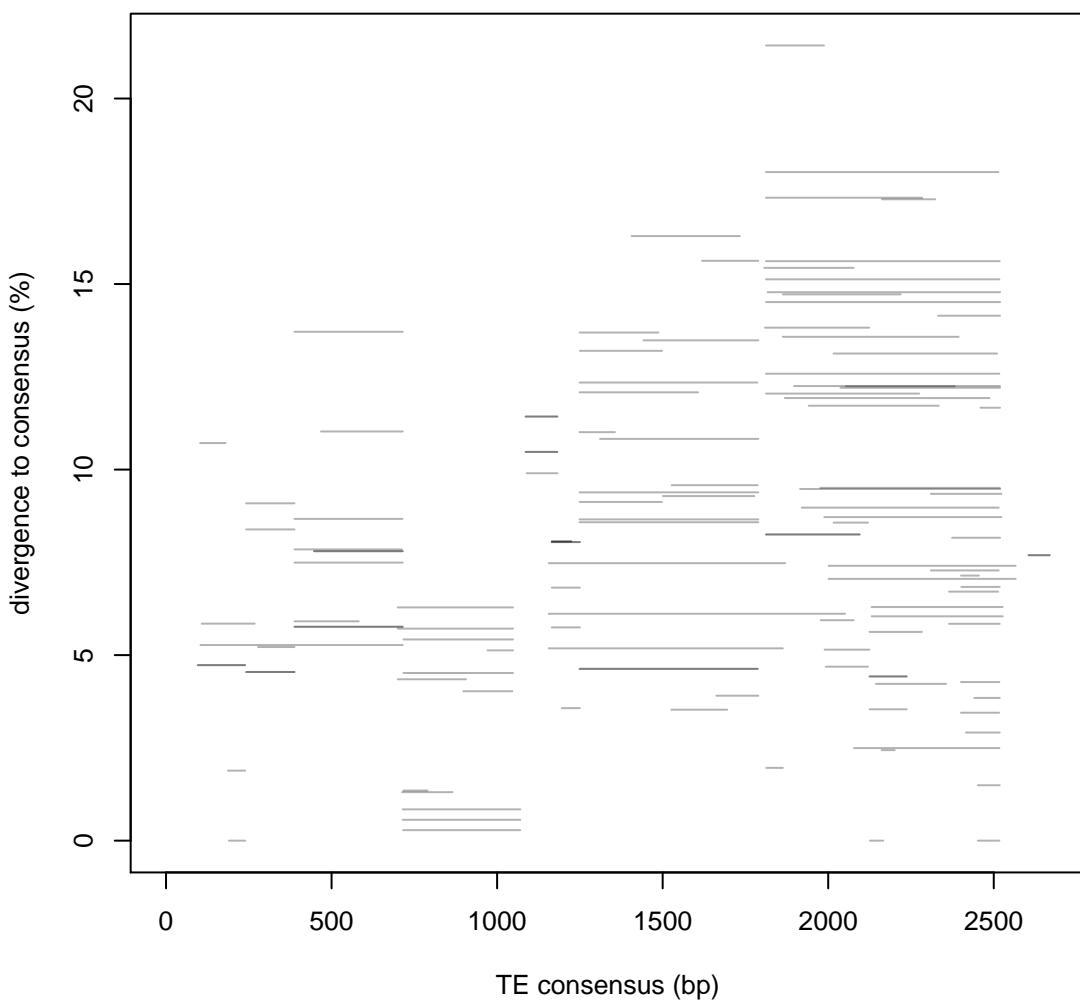
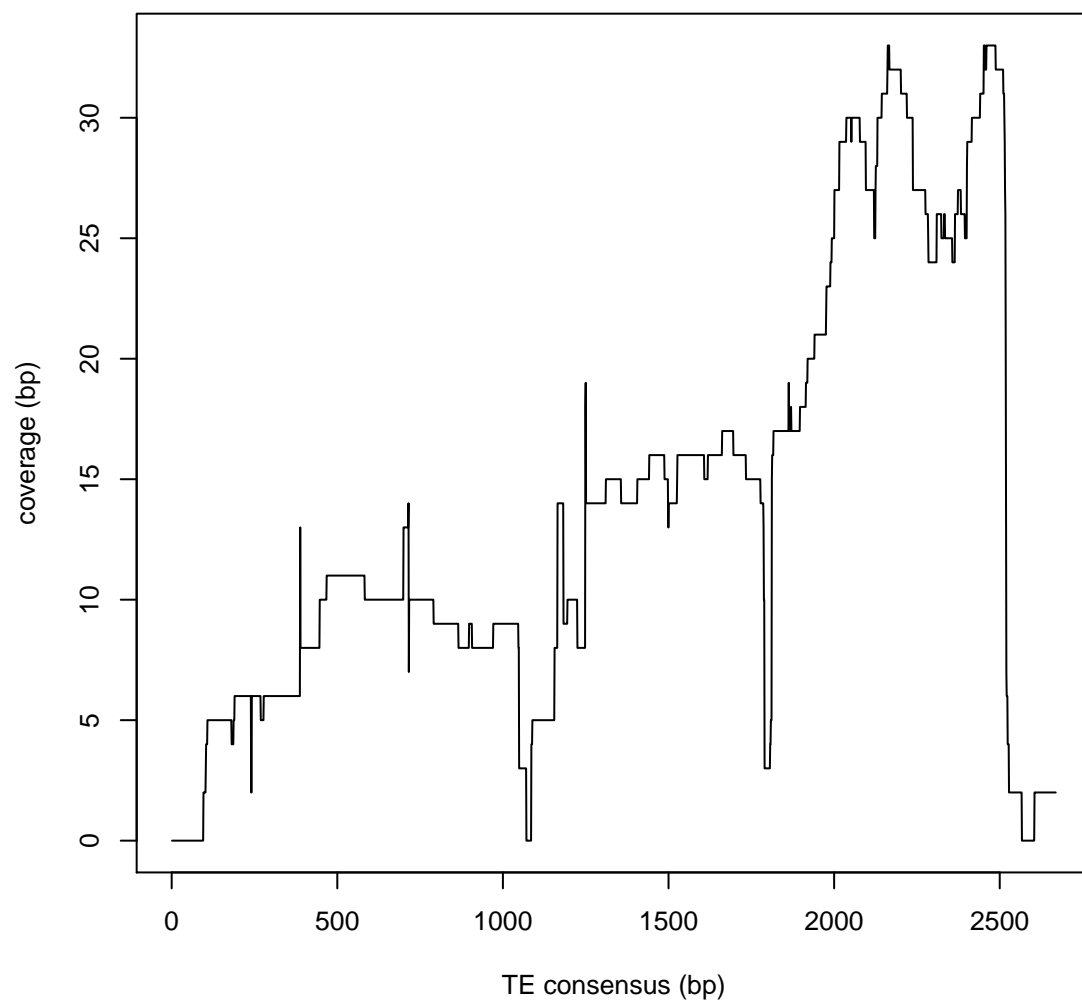


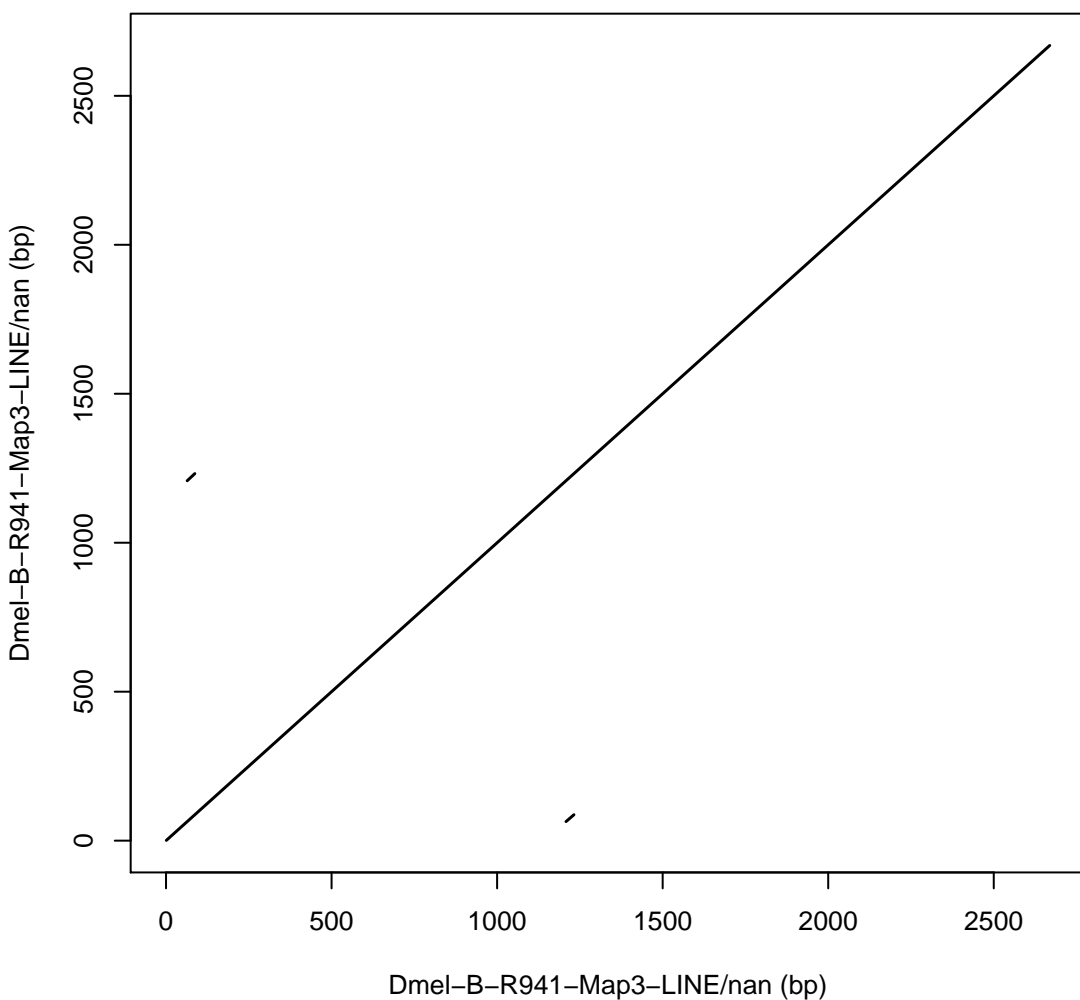
TE: Dmel-B-R941-Map3-LINE/nan
consensus size: 2669bp; fragments: 128; full length: 0 (>=2402.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

