TE: Dmel−B−G903−Map4_reversed−LINE/nan
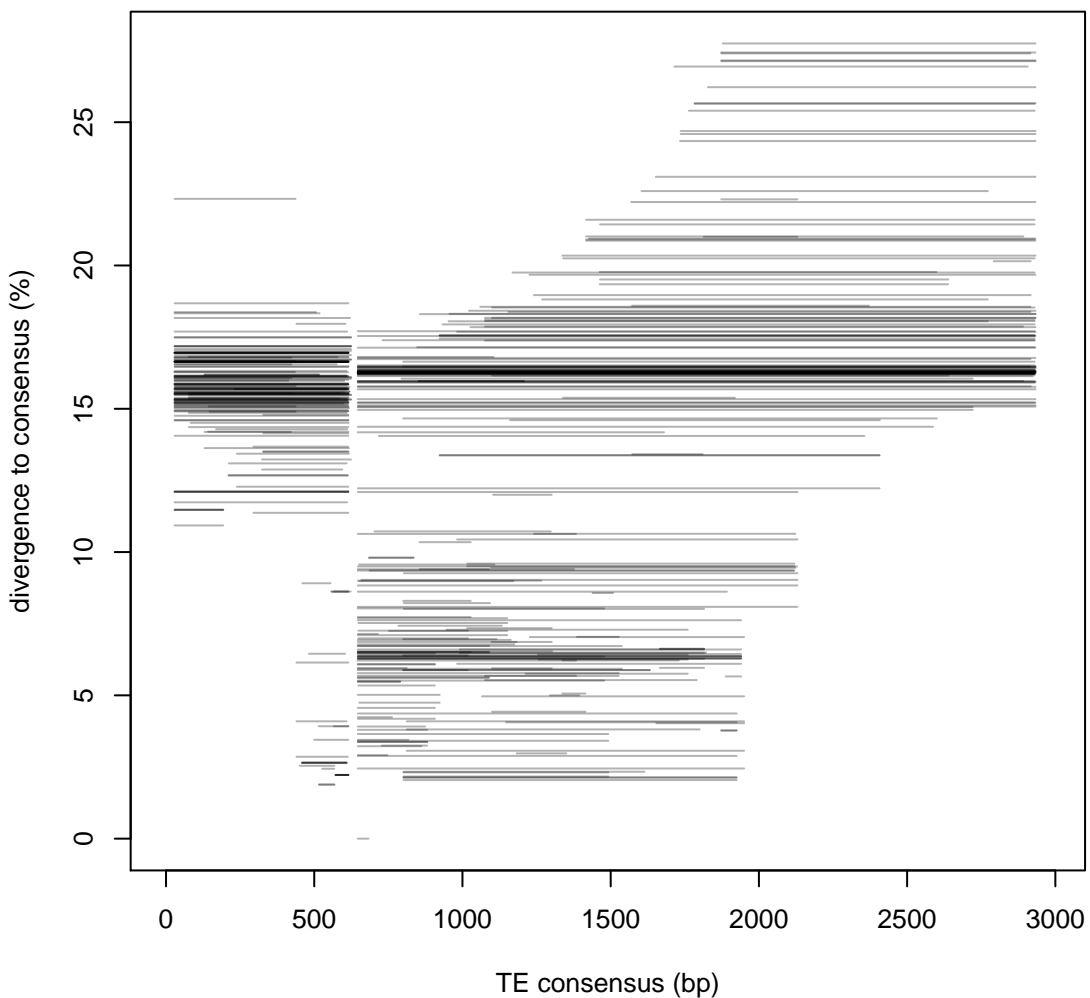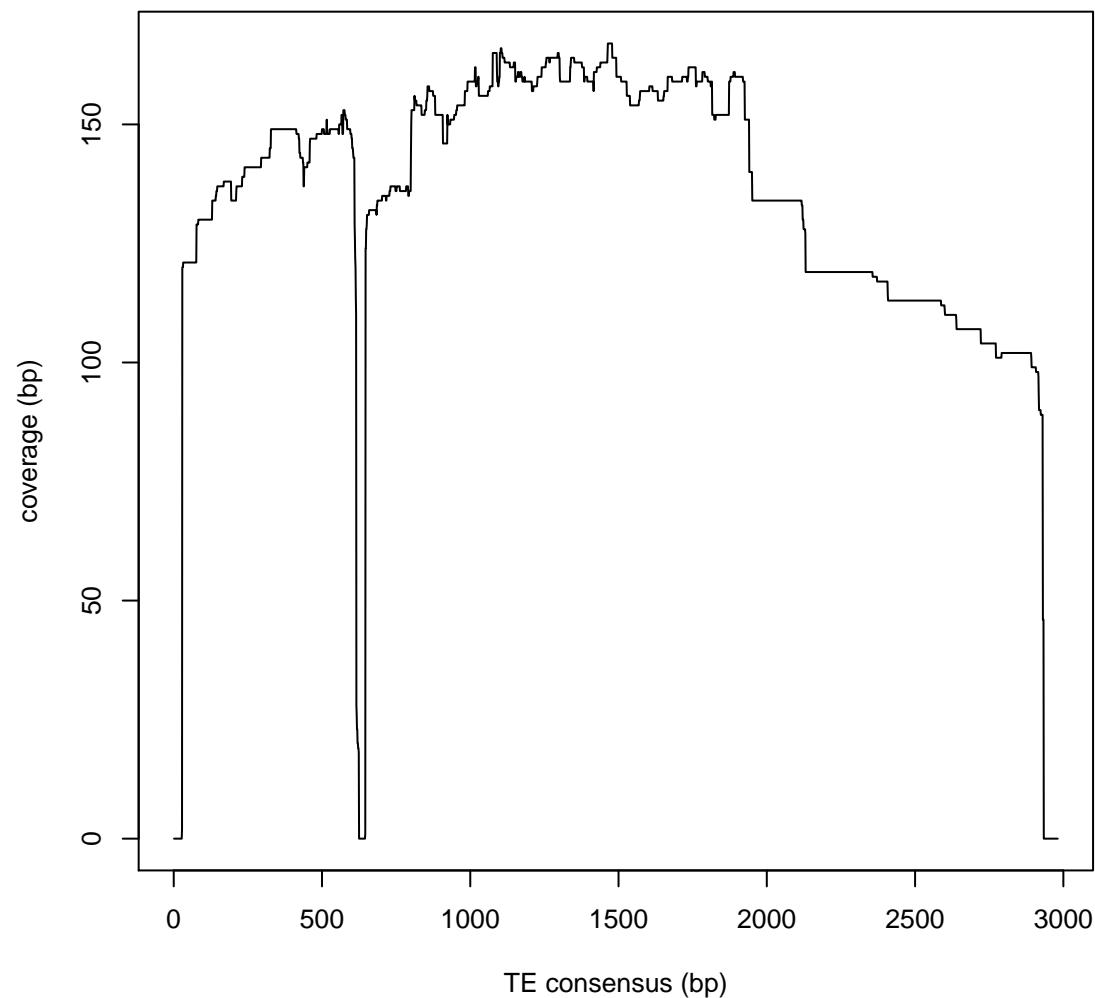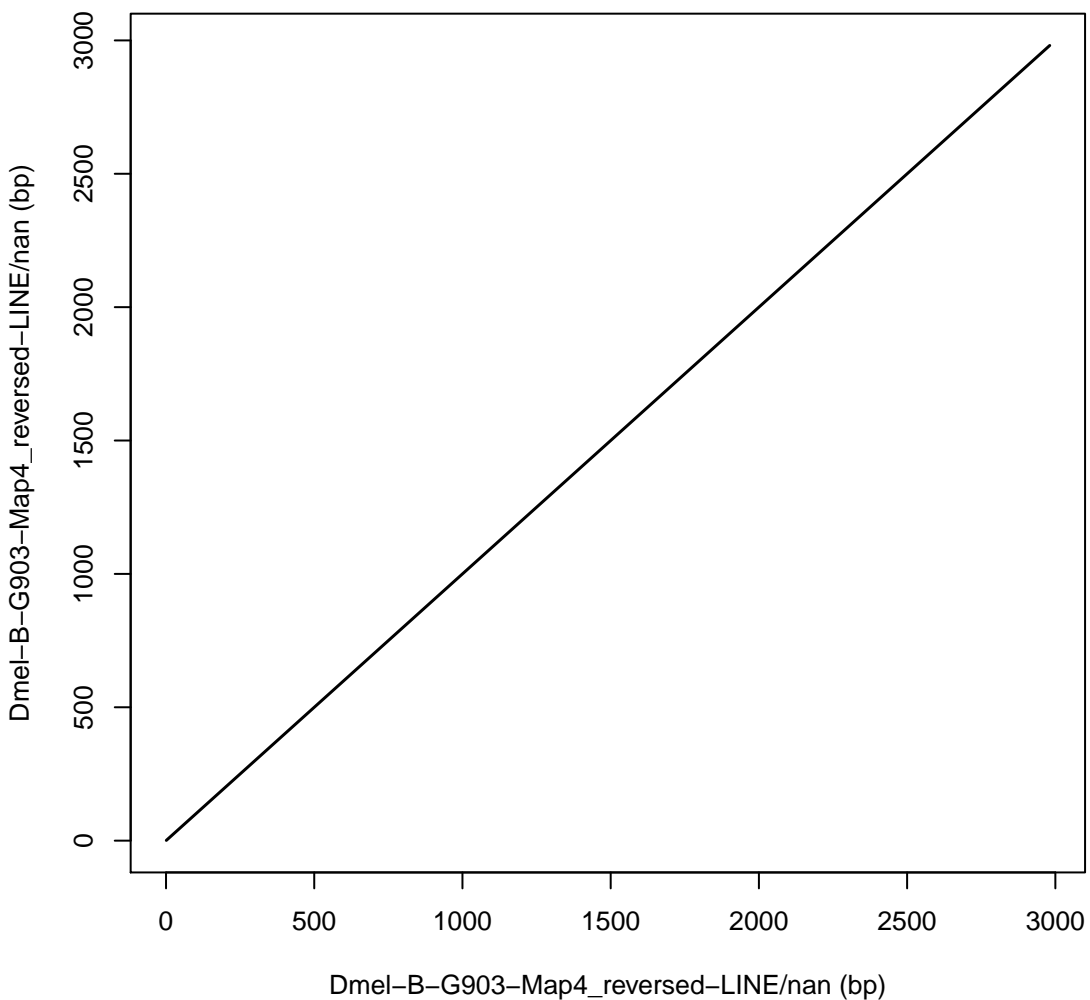### consensus size: 2981bp; fragments: 441; full length: 0 (>=2682.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected