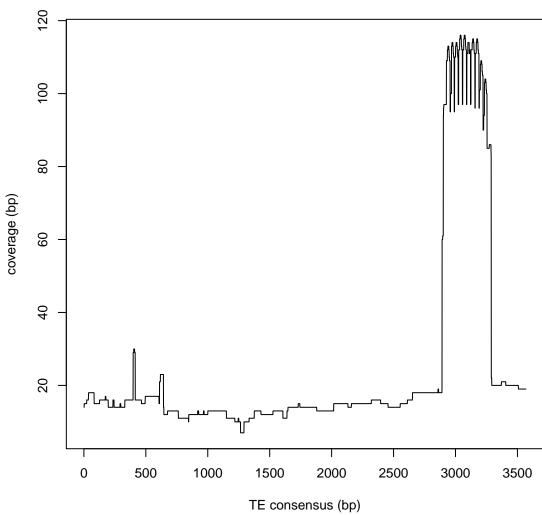
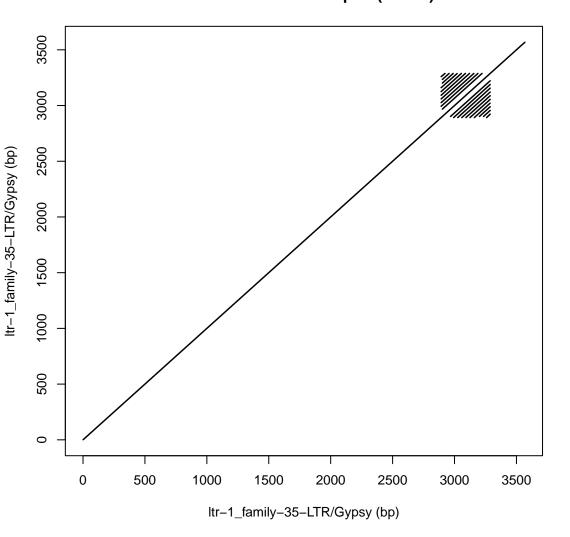
TE: ltr-1_family-35-LTR/Gypsy consensus size: 3568bp; fragments: 332; full length: 1 (>=3211.2bp) ∞ divergence to consensus (%) coverage (bp) α TE consensus (bp)

TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

