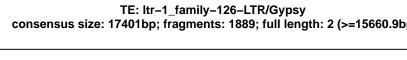
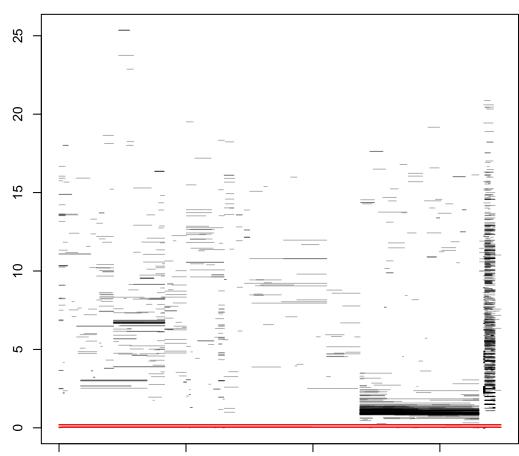
consensus size: 17401bp; fragments: 1889; full length: 2 (>=15660.9bp)

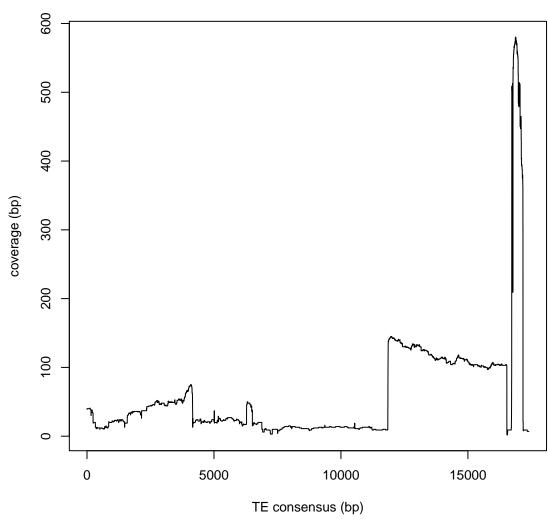




divergence to consensus (%)

0

## TE consensus genomic coverage



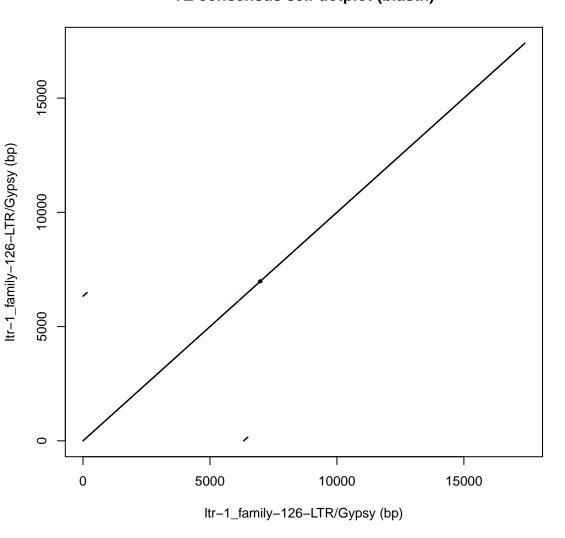
## TE consensus self dotplot (blastn)

TE consensus (bp)

10000

15000

5000



## TE consensus structure and protein hits

