



**Gene: SRY** ENSG00000279710

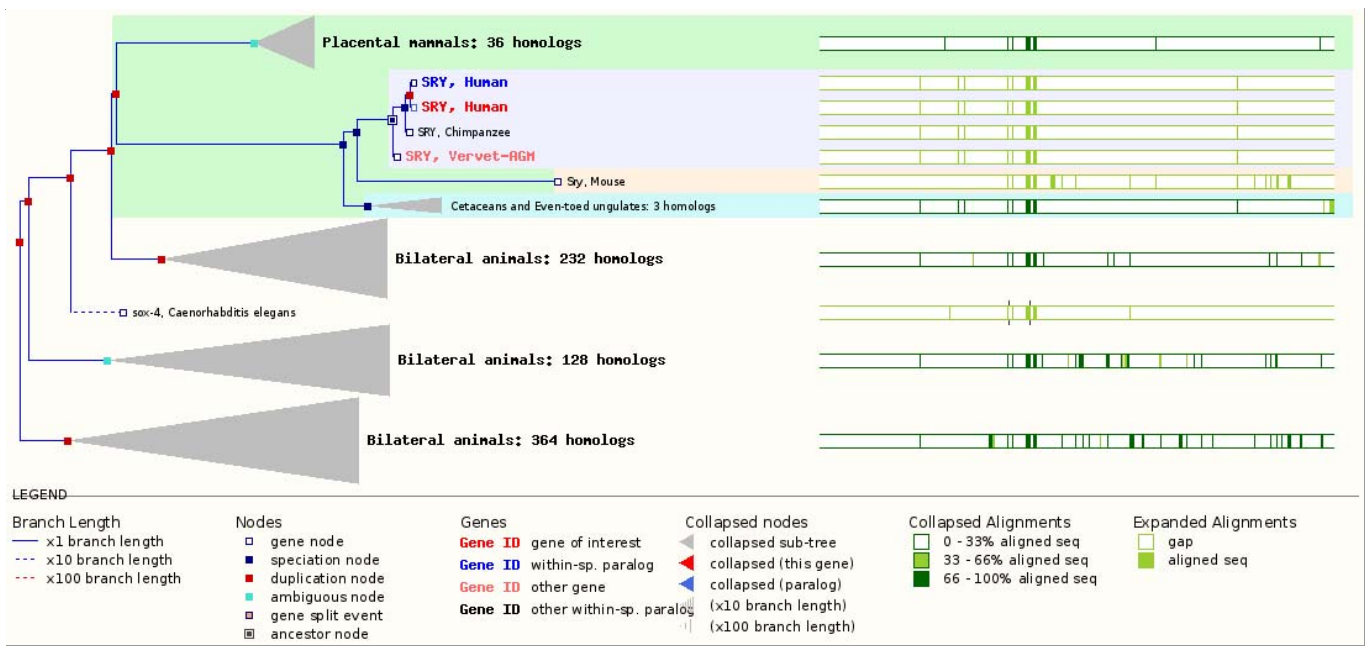
**Description** sex determining region Y [Source:HGNC Symbol;Acc:HGNC:11311]  
**Synonyms** TDF  
**Location** [Chromosome Y: 2,786,989-2,787,603](#) reverse strand.  
**INSDC coordinates** chromosome:GRCh38:CM000686.2:2786989:2787603:1  
**Transcripts** This gene has 1 transcript (splice variant) [Hide transcript table](#)

Name	Transcript ID	bp	Protein	Biotype	CCDS	Flags
SRY-201	<a href="#">ENST00000623823</a>	615	204 aa	Protein coding	<a href="#">CCDS14772</a>	GENCODE basic APPRIS PI

**Gene tree** ⓘ

GeneTree [ENSGT00760000118988](#)

**Number of genes** 769  
**Number of speciation nodes** 647  
**Number of duplication** 63  
**Number of ambiguous** 58  
**Number of gene split events** 0



- View options:**
- [View current genes only](#)
  - [View paralogs of current genes](#)
  - [View all duplication nodes](#)
  - [View fully expanded tree](#)
  - [Switch off highlighting](#)

Use the 'configure page' link in the left panel to set the default. Further options are available from menus on individual tree nodes.