

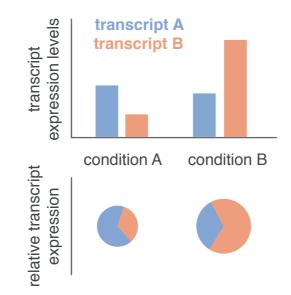
self-contained html (+ txt, json)

OUTPUT

high resolution plots

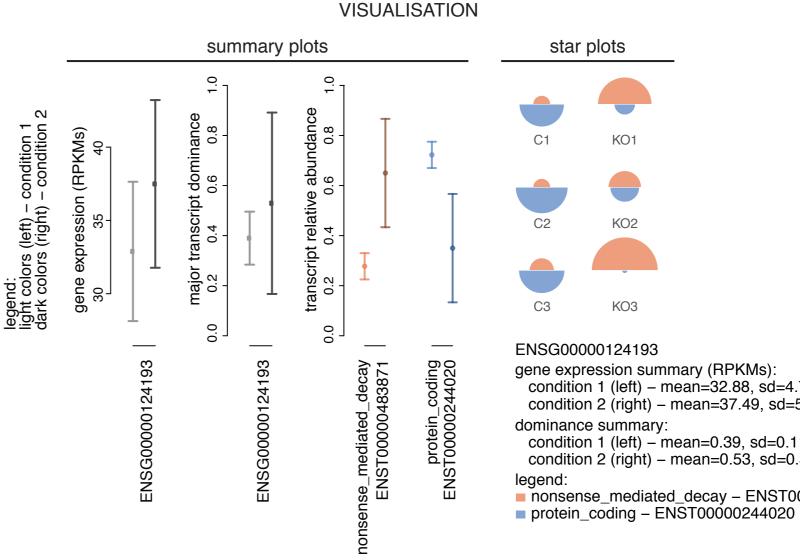
SwitchSeq





ANNOTATION

- i. transcript biotype [Ensembl]
- ii. principal transcripts [APPRIS]
- iii. transcript expression breadth
- iv. % overlap in the coding sequence [EMBOSS needle] link to protein structure [UniPDB]



tviz

