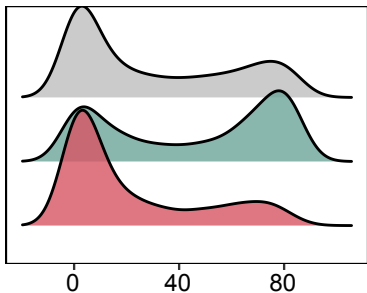


Broad conservation



Number of transcription factors

