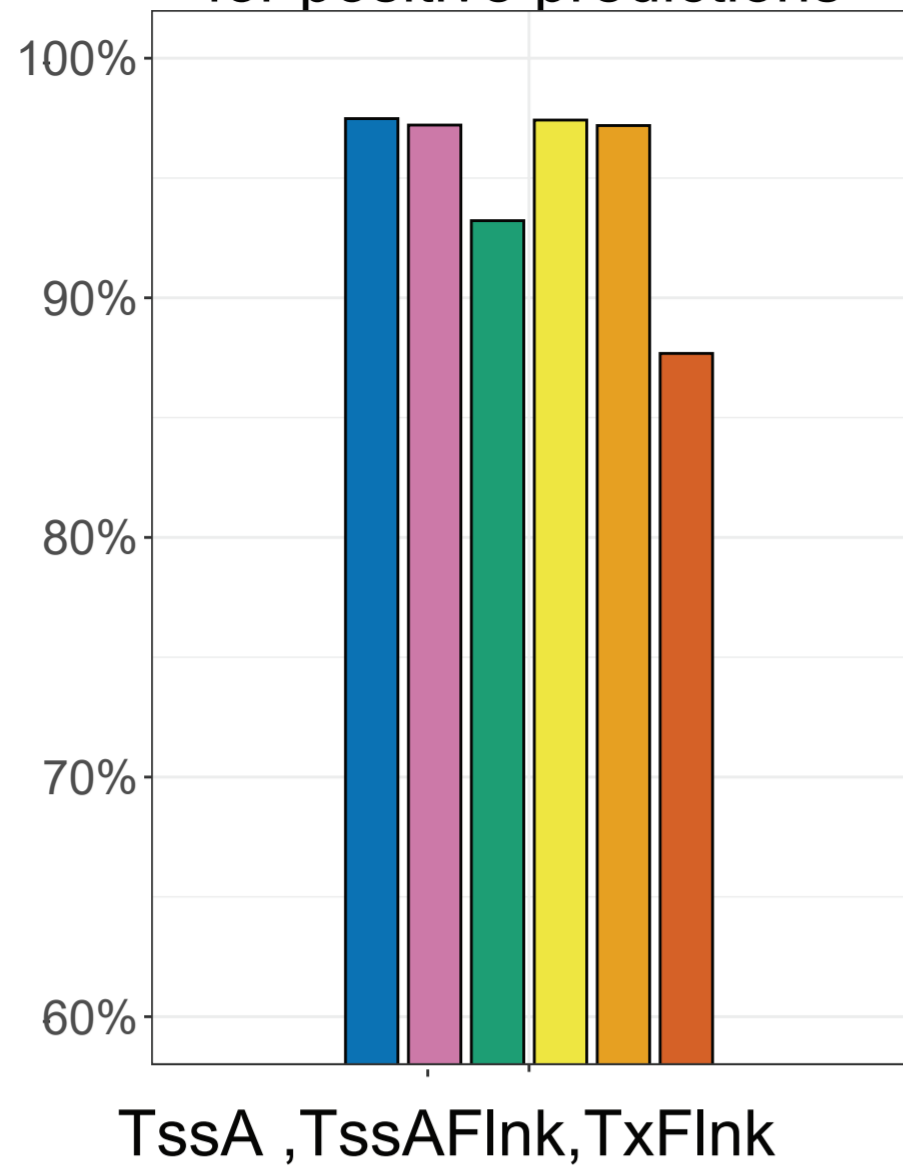
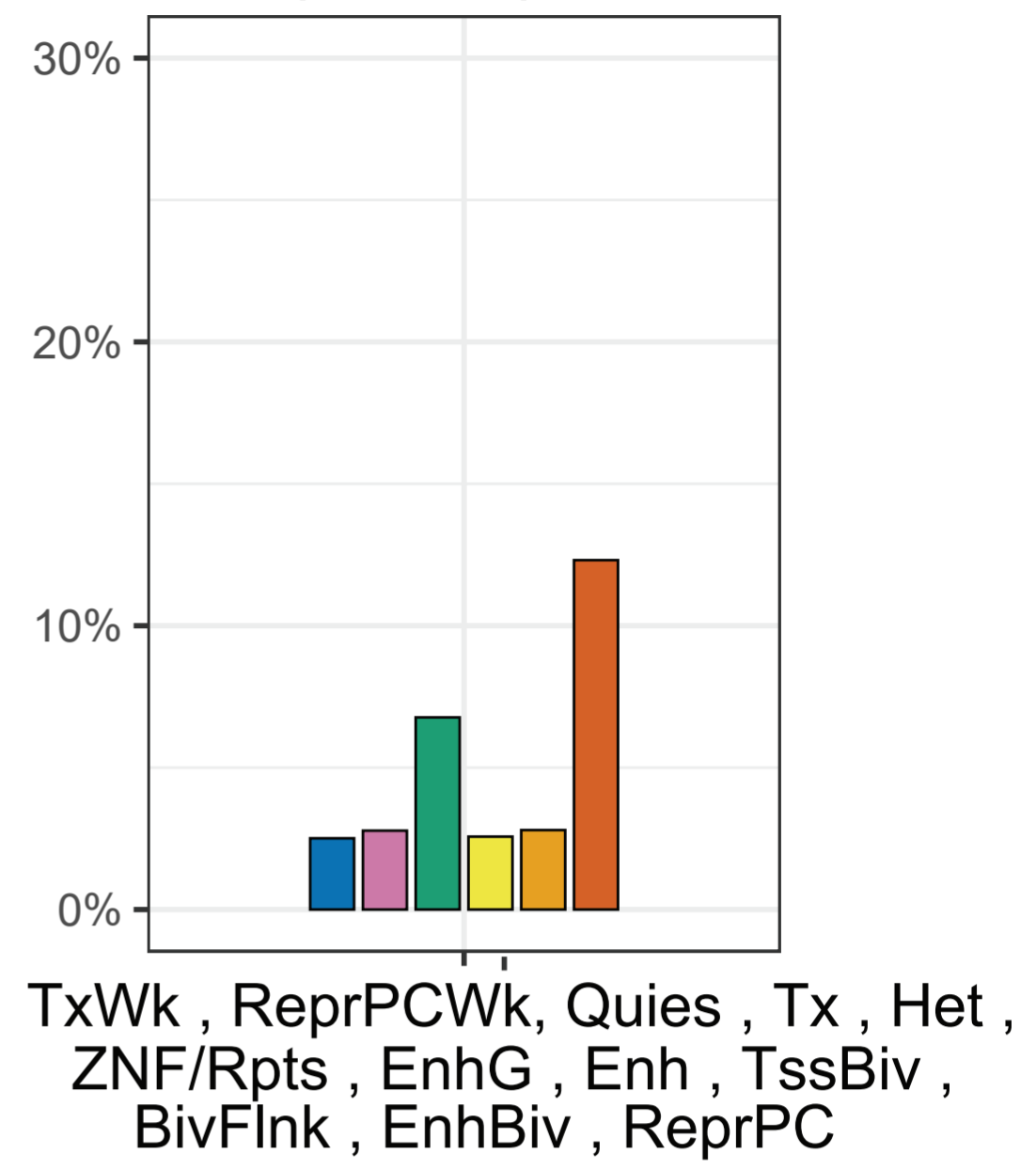


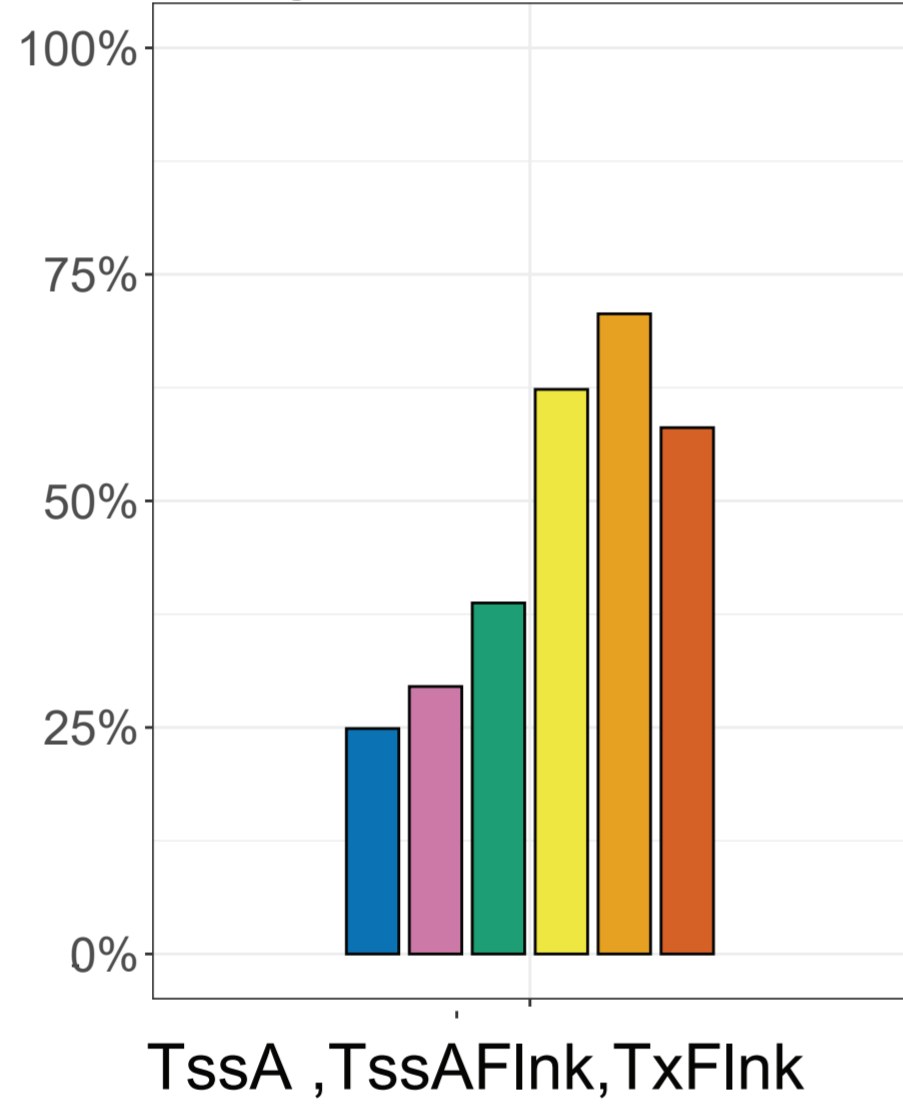
A Active Transcription ChromHMM Chromatin States for positive predictions



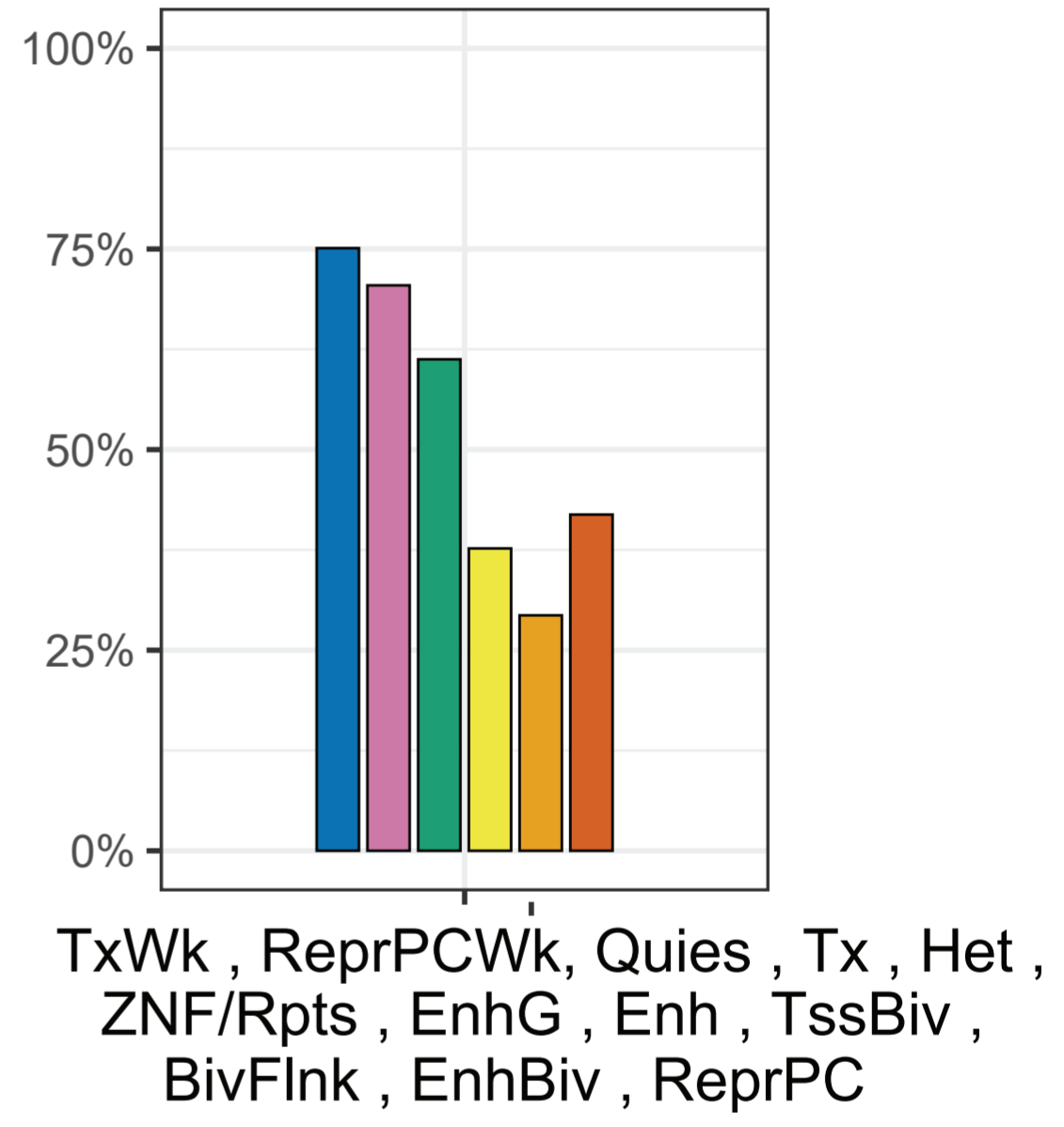
Weak Transcription/Repressed ChromHMM Chromatin States for positive predictions



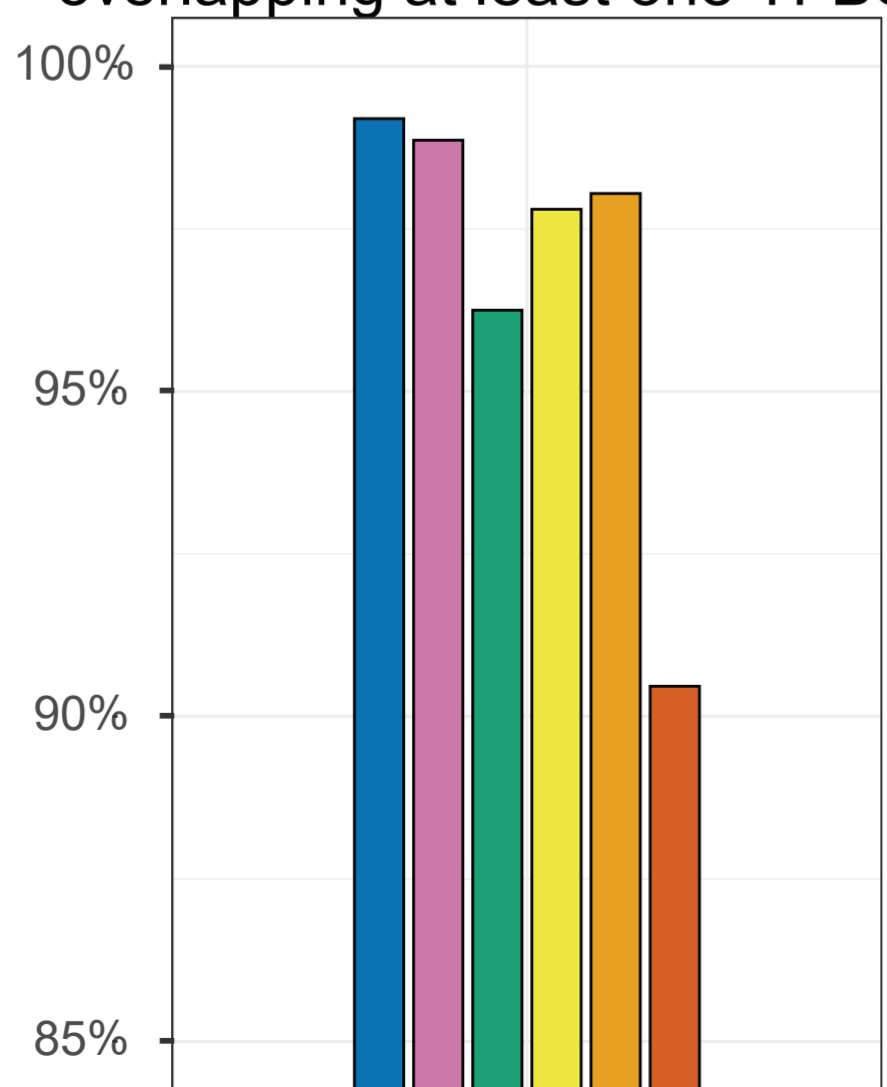
B Active Transcription ChromHMM Chromatin States for negative predictions



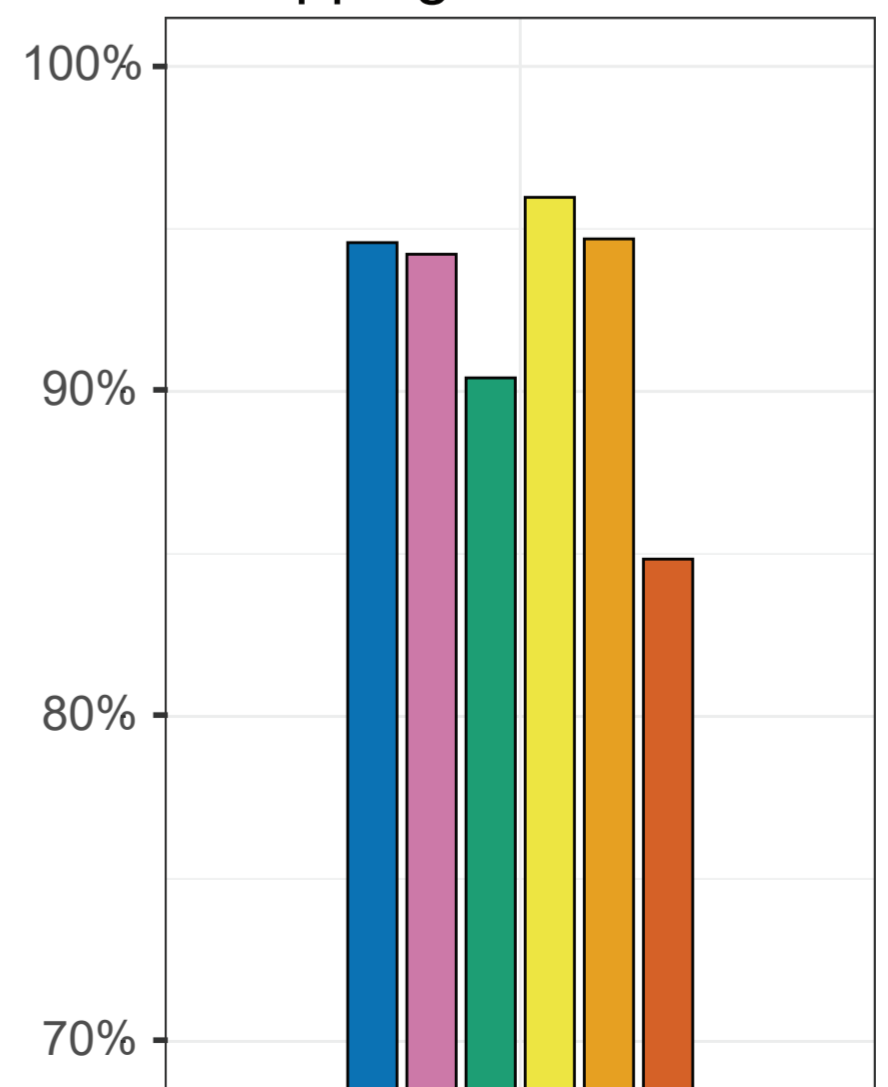
Weak Transcription/Repressed ChromHMM Chromatin States for negative predictions



C Percentage of each algorithm's predictions overlapping at least one TFBS



D Percentage of each algorithm's predictions overlapping H3K4me3 Peaks



■ DeepTSS ■ TOMETOOLS ■ RECLU
■ ADAPT-CAGE ■ PARACLUS ■ CAGER