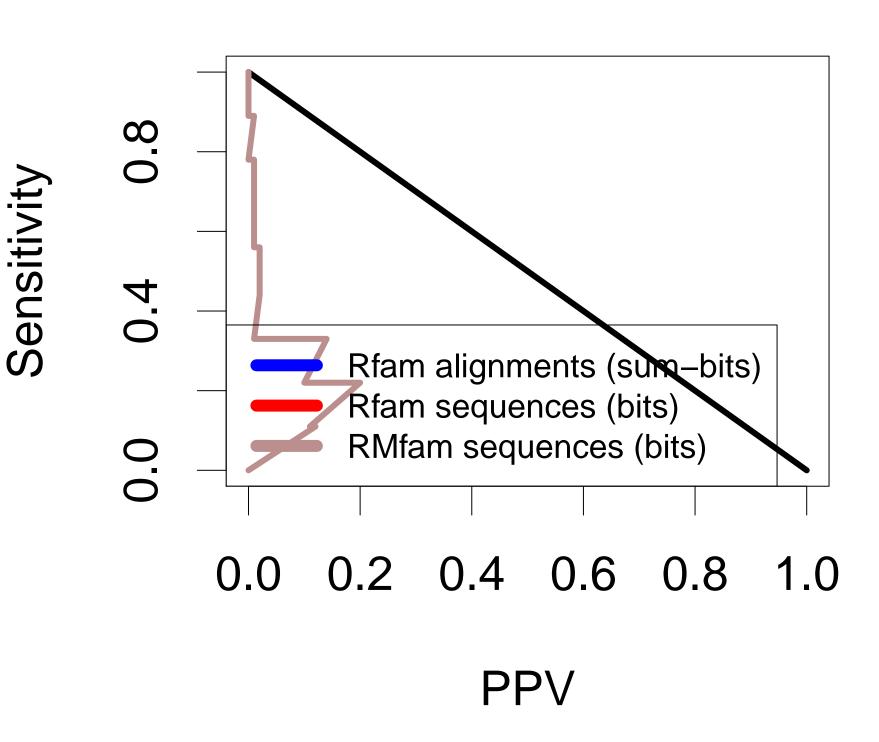
ROC curve: HuR_binding

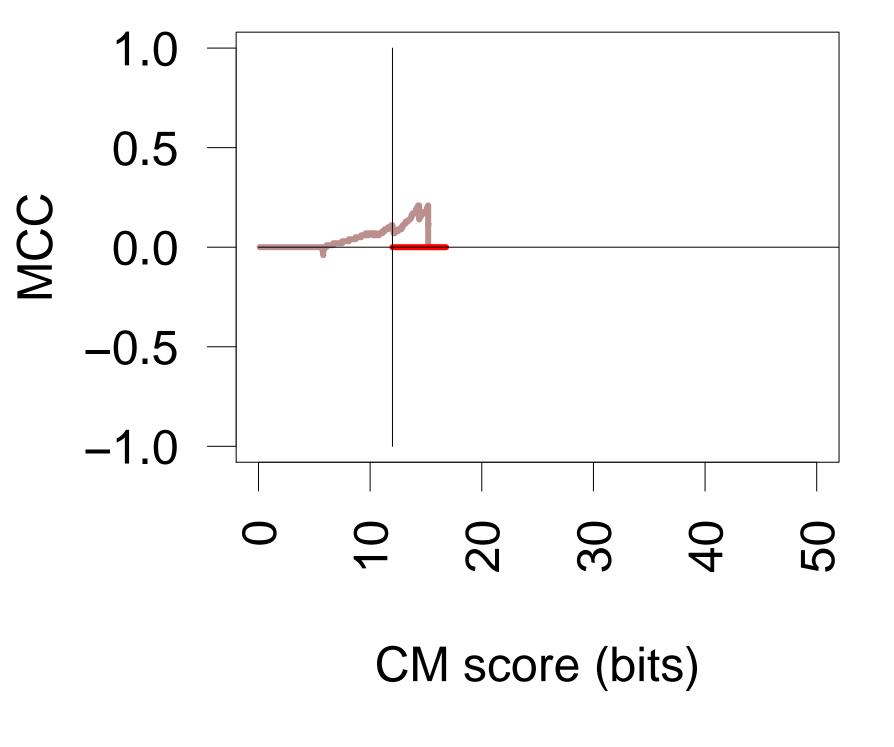
Rfam alignments (sum-bits) Rfam sequences (bits) RMfam sequences (bits) 0.0 0.2 0.4 0.6 0.8 1.0 Specificity

Sensitivity

ROC-like curve: HuR_binding



Score vs MCC: HuR_binding



Performance: HuR_binding

