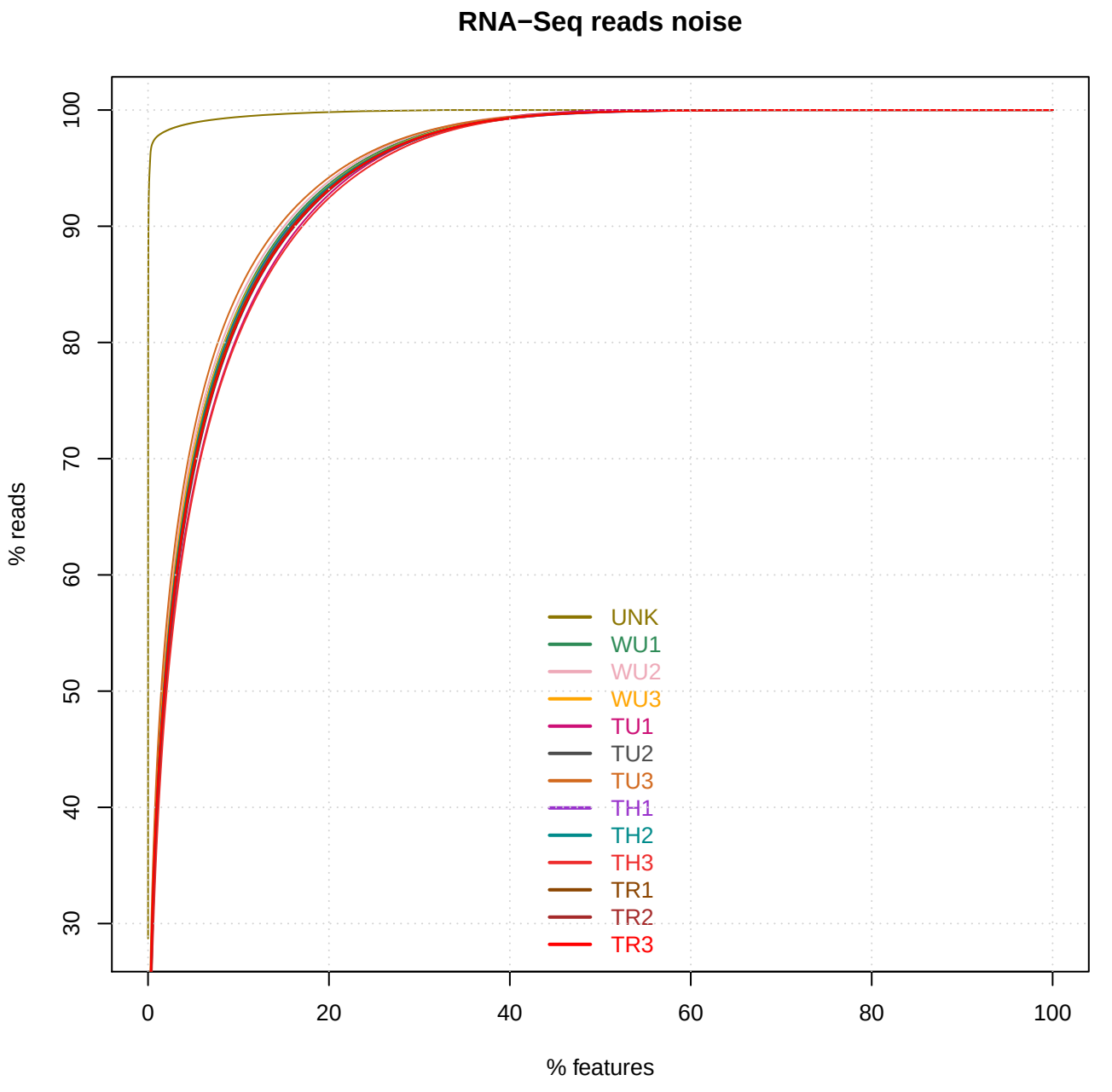
RNA−Seq reads noise
% reads
% features
100
90
80
70
60
50
40
30
0
20
40
60
80
100
UNK
WU1
WU2
WU3
TU1
TU2
TU3
TH1
TH2
TH3
TR1
TR2
TR3