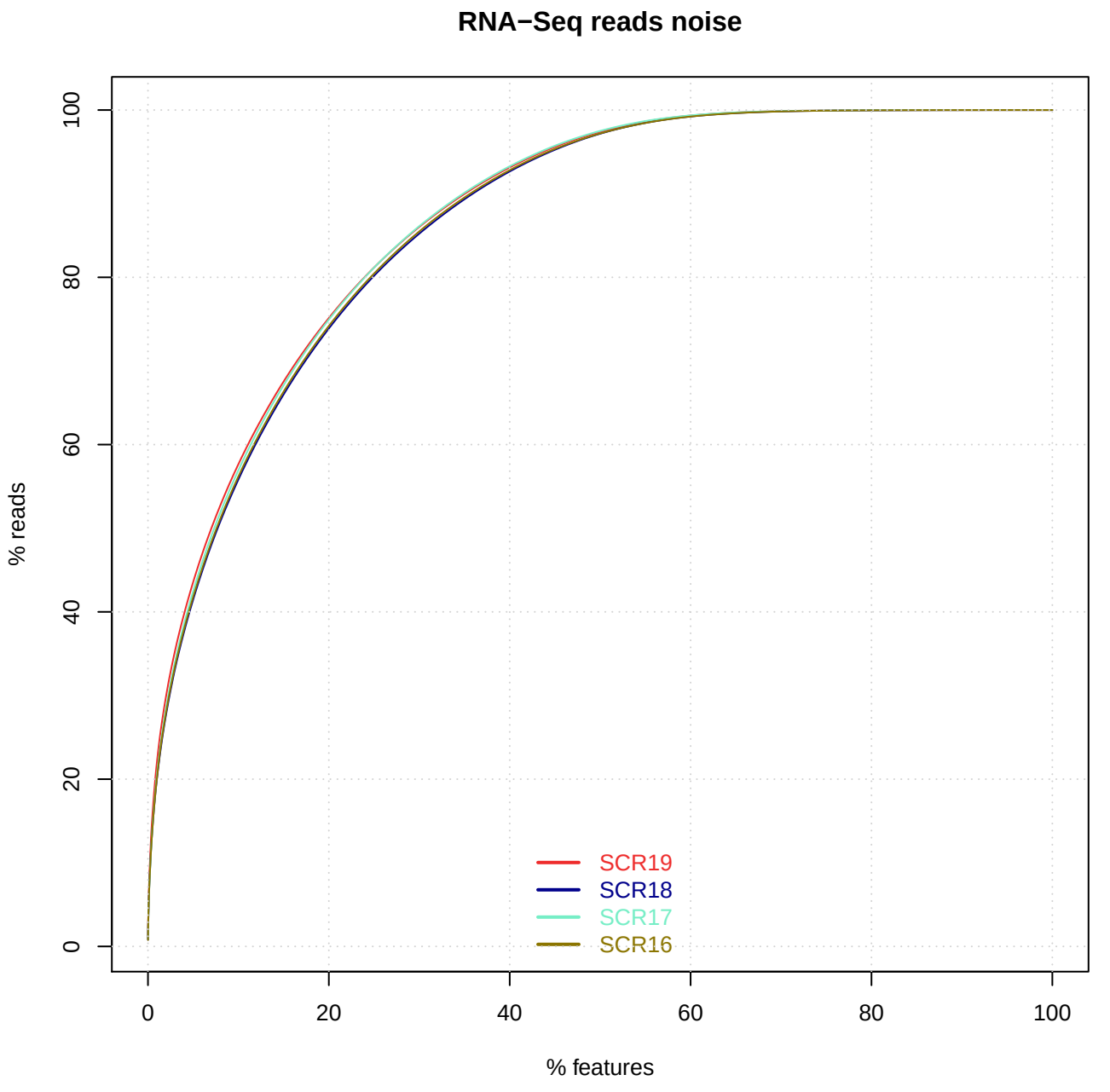
RNA−Seq reads noise
% reads
% features
0
20
40
60
80
100
SCR19
SCR18
SCR17
SCR16