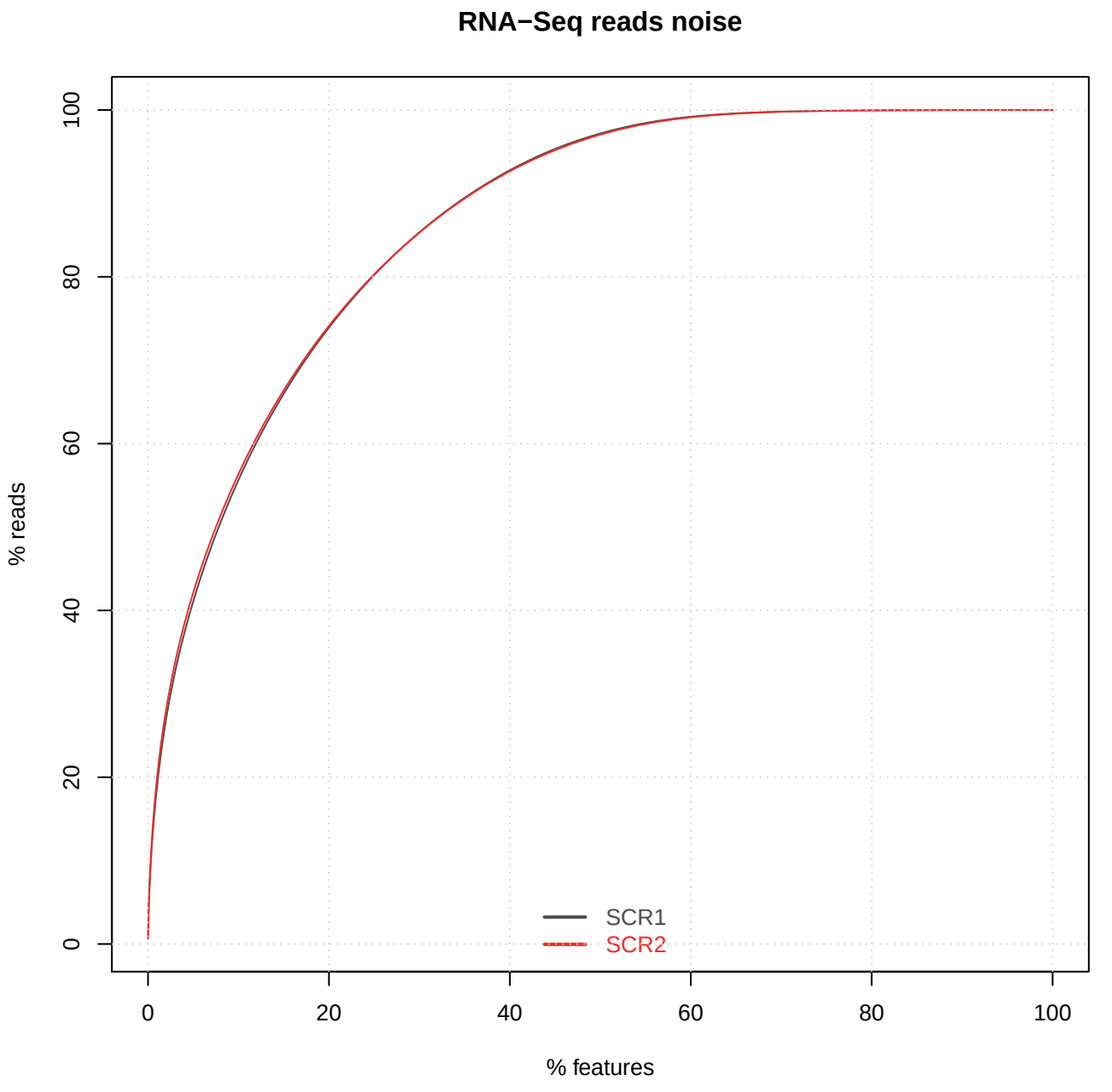
RNA−Seq reads noise
% reads
% features
0
20
40
60
80
100
SCR1
SCR2