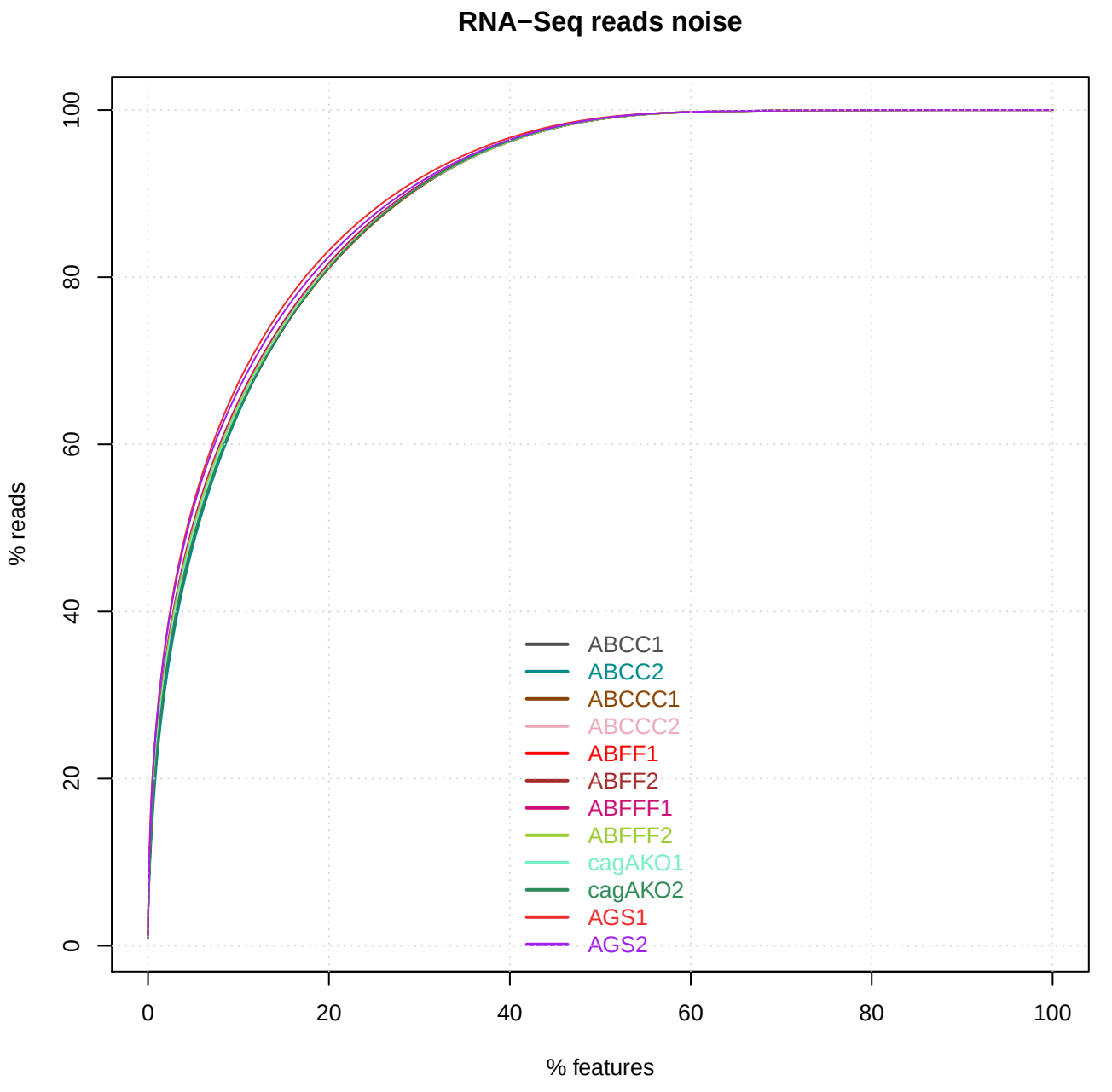
RNA−Seq reads noise
% reads
% features
0
20
40
60
80
100
ABCC1
ABCC2
ABCCC1
ABCCC2
ABFF1
ABFF2
ABFFF1
ABFFF2
cagAKO1
cagAKO2
AGS1
AGS2