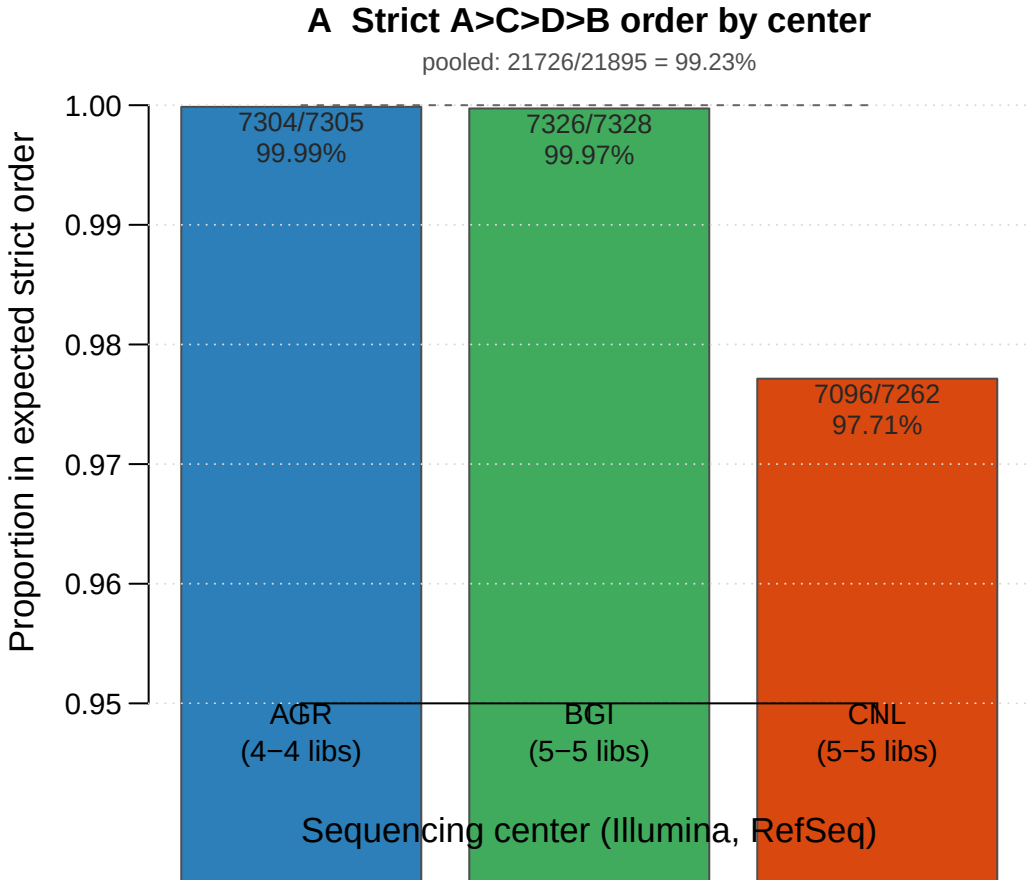
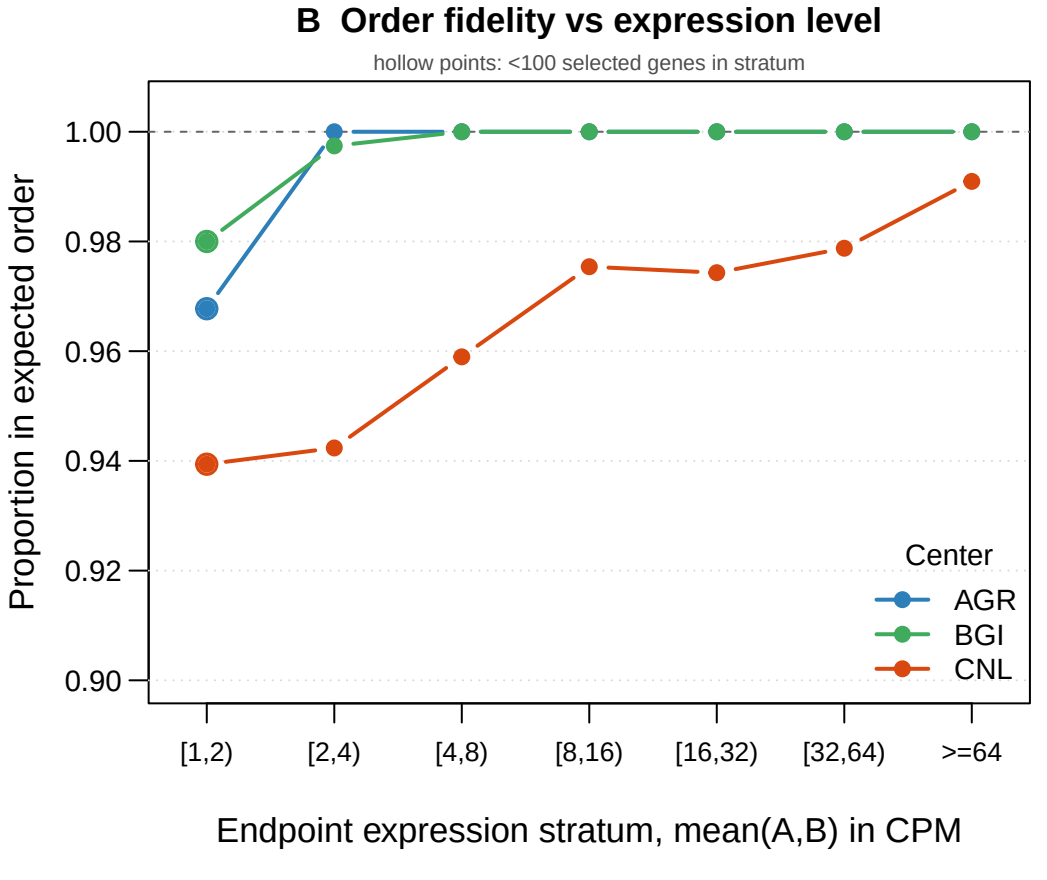

A Strict A>C>D>B order by center
pooled: 21726/21895 = 99.23%
Proportion in expected strict order
1.00
0.99
0.98
0.97
0.96
0.95
7304/7305
99.99%
7326/7328
99.97%
7096/7262
97.71%
AGR
(4−4 libs)
BGI
(5−5 libs)
CNL
(5−5 libs)
Sequencing center (Illumina, RefSeq)


B Order fidelity vs expression level
hollow points: <100 selected genes in stratum
Proportion in expected order
1.00
0.98
0.96
0.94
0.92
0.90
Center
AGR
BGI
CNL
[1,2)
[2,4)
[4,8)
[8,16)
[16,32)
[32,64)
>=64
Endpoint expression stratum, mean(A,B) in CPM