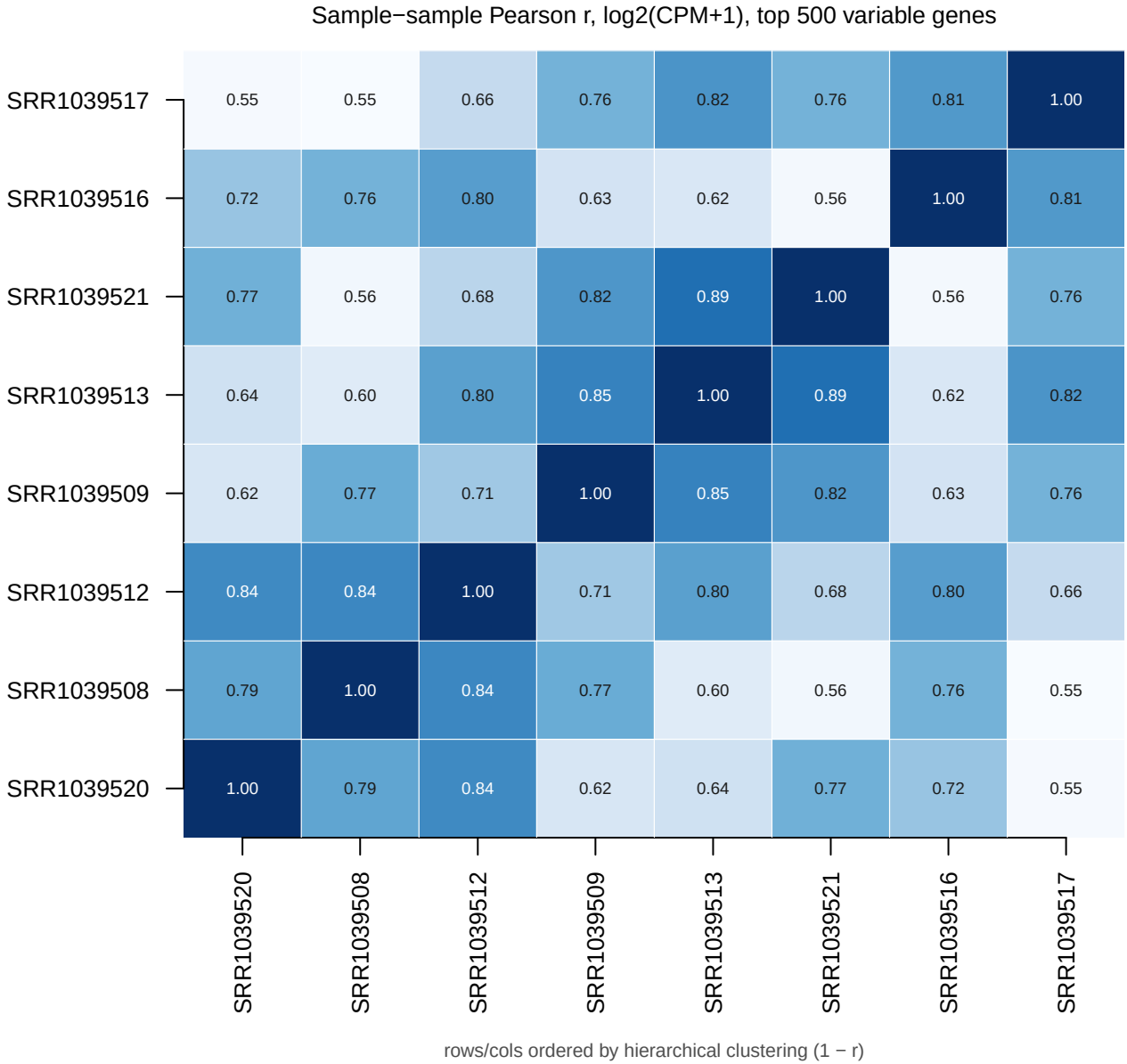
Sample−sample Pearson r, log2(CPM+1), top 500 variable genes
SRR1039517
SRR1039516
SRR1039521
SRR1039513
SRR1039509
SRR1039512
SRR1039508
SRR1039520
0.55 0.55 0.66 0.76 0.82 0.76 0.81 1.00
0.72 0.76 0.80 0.63 0.62 0.56 1.00 0.81
0.77 0.56 0.68 0.82 0.89 1.00 0.56 0.76
0.64 0.60 0.80 0.85 1.00 0.89 0.62 0.82
0.62 0.77 0.71 1.00 0.85 0.82 0.63 0.76
0.84 0.84 1.00 0.71 0.80 0.68 0.80 0.66
0.79 1.00 0.84 0.77 0.60 0.56 0.76 0.55
1.00 0.79 0.84 0.62 0.64 0.77 0.72 0.55
SRR1039520
SRR1039508
SRR1039512
SRR1039509
SRR1039513
SRR1039521
SRR1039516
SRR1039517
rows/cols ordered by hierarchical clustering (1 − r)