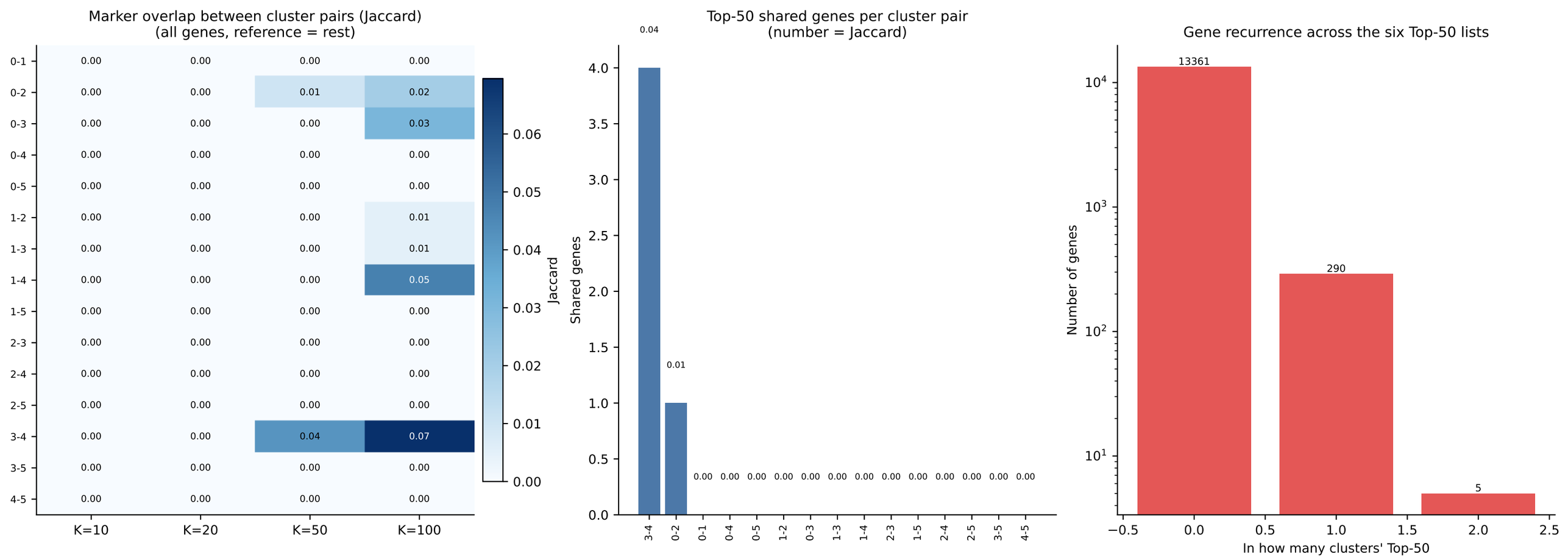
Marker overlap between cluster pairs (Jaccard)
(all genes, reference = rest)
0-1 0.00 0.00 0.00 0.00
0-2 0.00 0.00 0.01 0.02
0-3 0.00 0.00 0.00 0.03
0-4 0.00 0.00 0.00 0.00
0-5 0.00 0.00 0.00 0.00
1-2 0.00 0.00 0.00 0.01
1-3 0.00 0.00 0.00 0.01
1-4 0.00 0.00 0.00 0.05
1-5 0.00 0.00 0.00 0.00
2-3 0.00 0.00 0.00 0.00
2-4 0.00 0.00 0.00 0.00
2-5 0.00 0.00 0.00 0.00
3-4 0.00 0.00 0.04 0.07
3-5 0.00 0.00 0.00 0.00
4-5 0.00 0.00 0.00 0.00
K=10
K=20
K=50
K=100
Jaccard
0.00
0.01
0.02
0.03
0.04
0.05
0.06
Top-50 shared genes per cluster pair
(number = Jaccard)
Shared genes
0.0
0.5
1.0
1.5
2.0
2.5
3.0
3.5
4.0
0.04
0.01
0.00
3-4
0-2
0-1
0-4
0-5
1-2
0-3
1-3
1-4
2-3
1-5
2-4
2-5
3-5
4-5
Gene recurrence across the six Top-50 lists
Number of genes
10^1
10^2
10^3
10^4
13361
290
5
−0.5
0.0
0.5
1.0
1.5
2.0
2.5
In how many clusters' Top-50