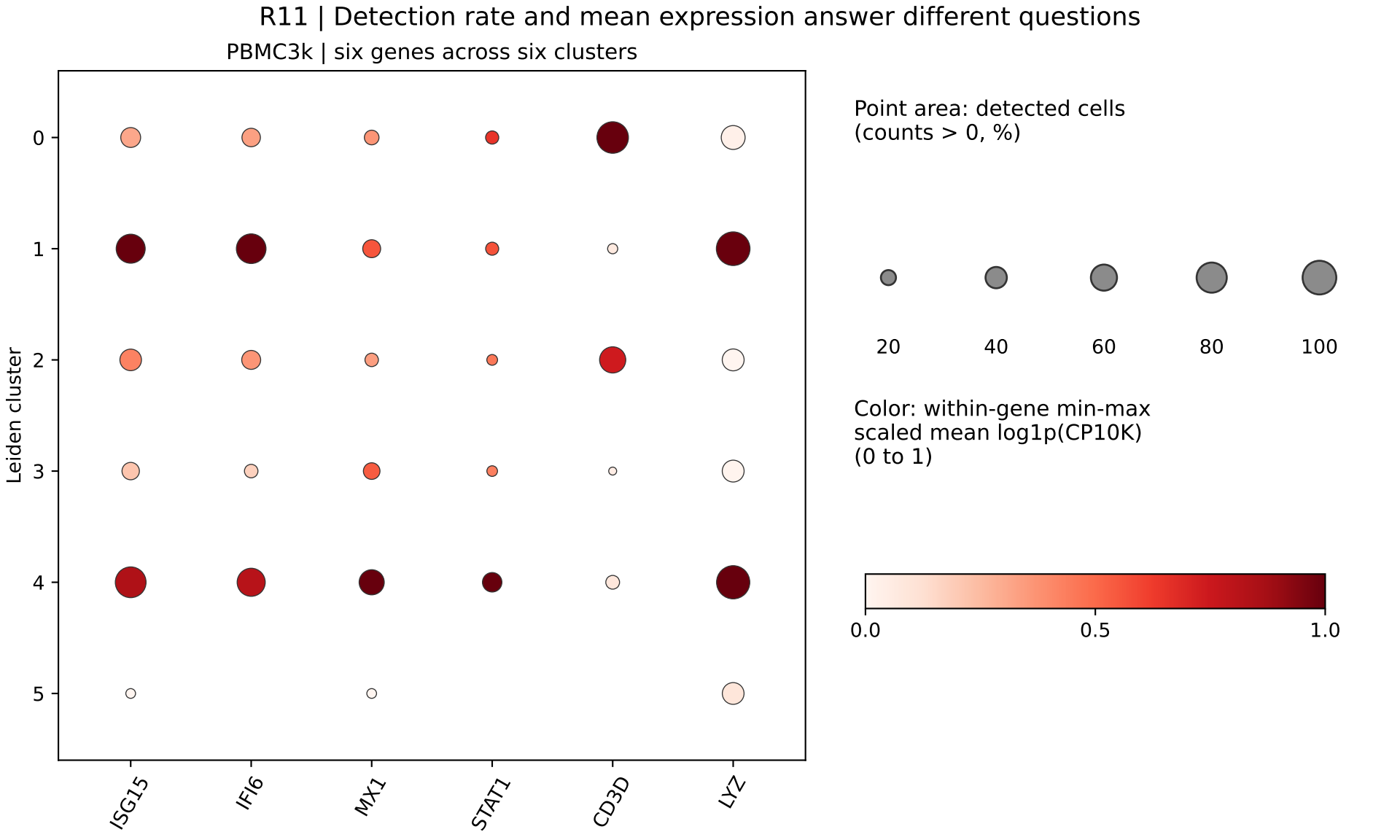
R11 | Detection rate and mean expression answer different questions
PBMC3k | six genes across six clusters
Leiden cluster
0
1
2
3
4
5
ISG15
IFI6
MX1
STAT1
CD3D
LYZ
Point area: detected cells
(counts > 0, %)
20
40
60
80
100
Color: within-gene min-max
scaled mean log1p(CP10K)
(0 to 1)
0.0
0.5
1.0