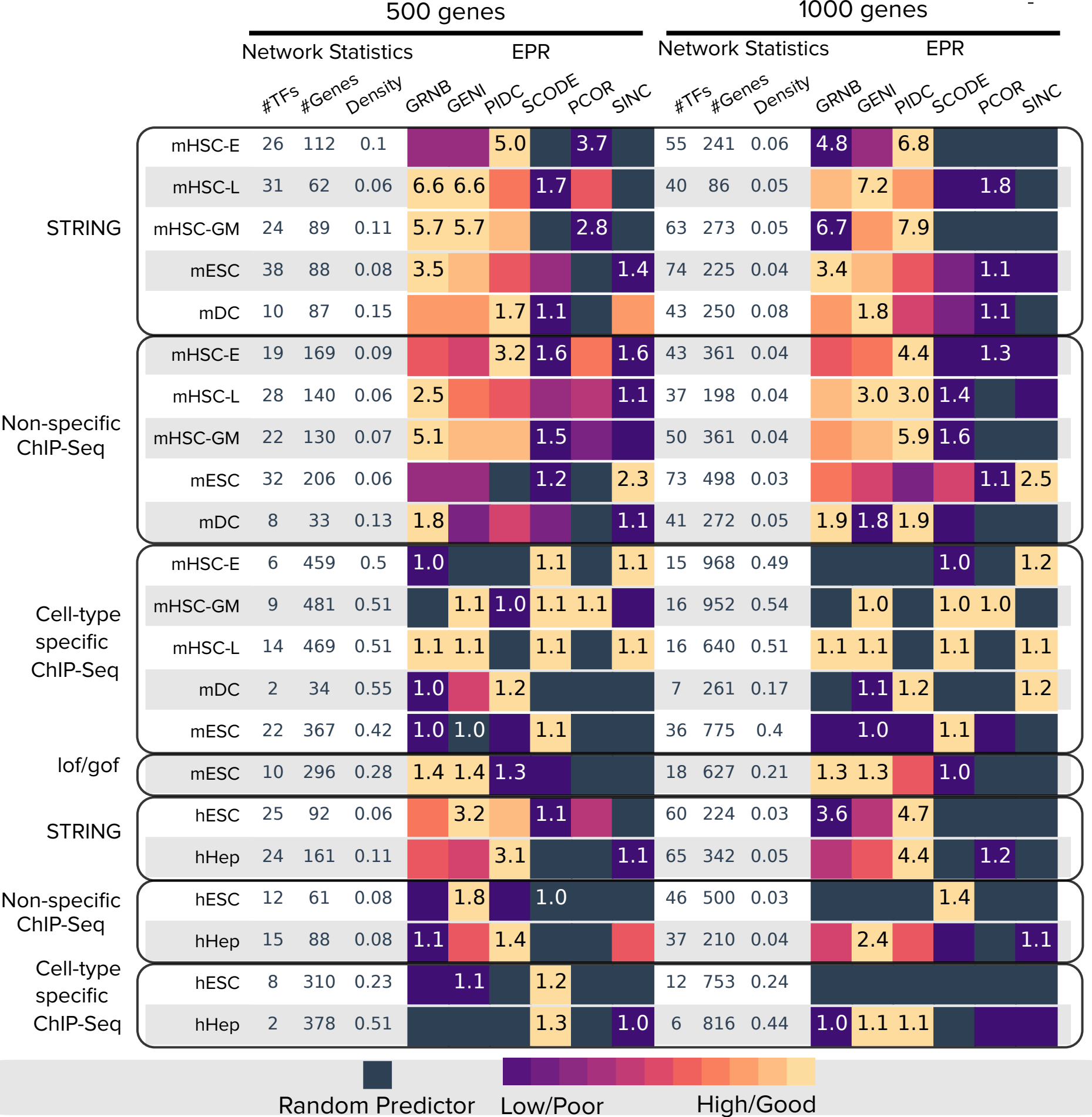
500 genes
1000 genes
Network Statistics
EPR
#TFs
#Genes
Density
GRNB
GENI
PIDC
SCODE
PCOR
SINC
STRING
mHSC-E 26 112 0.1 5.0 3.7 55 241 0.06 4.8 6.8
mHSC-L 31 62 0.06 6.6 6.6 1.7 40 86 0.05 7.2 1.8
mHSC-GM 24 89 0.11 5.7 5.7 2.8 63 273 0.05 6.7 7.9
mESC 38 88 0.08 3.5 1.4 74 225 0.04 3.4 1.1
mDC 10 87 0.15 1.7 1.1 43 250 0.08 1.8 1.1
Non-specific ChIP-Seq
mHSC-E 19 169 0.09 3.2 1.6 1.6 43 361 0.04 4.4 1.3
mHSC-L 28 140 0.06 2.5 1.1 37 198 0.04 3.0 3.0 1.4
mHSC-GM 22 130 0.07 5.1 1.5 50 361 0.04 5.9 1.6
mESC 32 206 0.06 1.2 2.3 73 498 0.03 1.1 2.5
mDC 8 33 0.13 1.8 1.1 41 272 0.05 1.9 1.8 1.9
Cell-type specific ChIP-Seq
mHSC-E 6 459 0.5 1.0 1.1 1.1 15 968 0.49 1.0 1.2
mHSC-GM 9 481 0.51 1.1 1.0 1.1 1.1 16 952 0.54 1.0 1.0 1.0
mHSC-L 14 469 0.51 1.1 1.1 1.1 1.1 16 640 0.51 1.1 1.1 1.1 1.1
mDC 2 34 0.55 1.0 1.2 7 261 0.17 1.1 1.2 1.2
mESC 22 367 0.42 1.0 1.0 1.1 36 775 0.4 1.0 1.1
lof/gof
mESC 10 296 0.28 1.4 1.4 1.3 18 627 0.21 1.3 1.3 1.0
STRING
hESC 25 92 0.06 3.2 1.1 60 224 0.03 3.6 4.7
hHep 24 161 0.11 3.1 1.1 65 342 0.05 4.4 1.2
Non-specific ChIP-Seq
hESC 12 61 0.08 1.8 1.0 46 500 0.03 1.4
hHep 15 88 0.08 1.1 1.4 37 210 0.04 2.4 1.1
Cell-type specific ChIP-Seq
hESC 8 310 0.23 1.1 1.2 12 753 0.24
hHep 2 378 0.51 1.3 1.0 6 816 0.44 1.0 1.1 1.1
Random Predictor
Low/Poor
High/Good