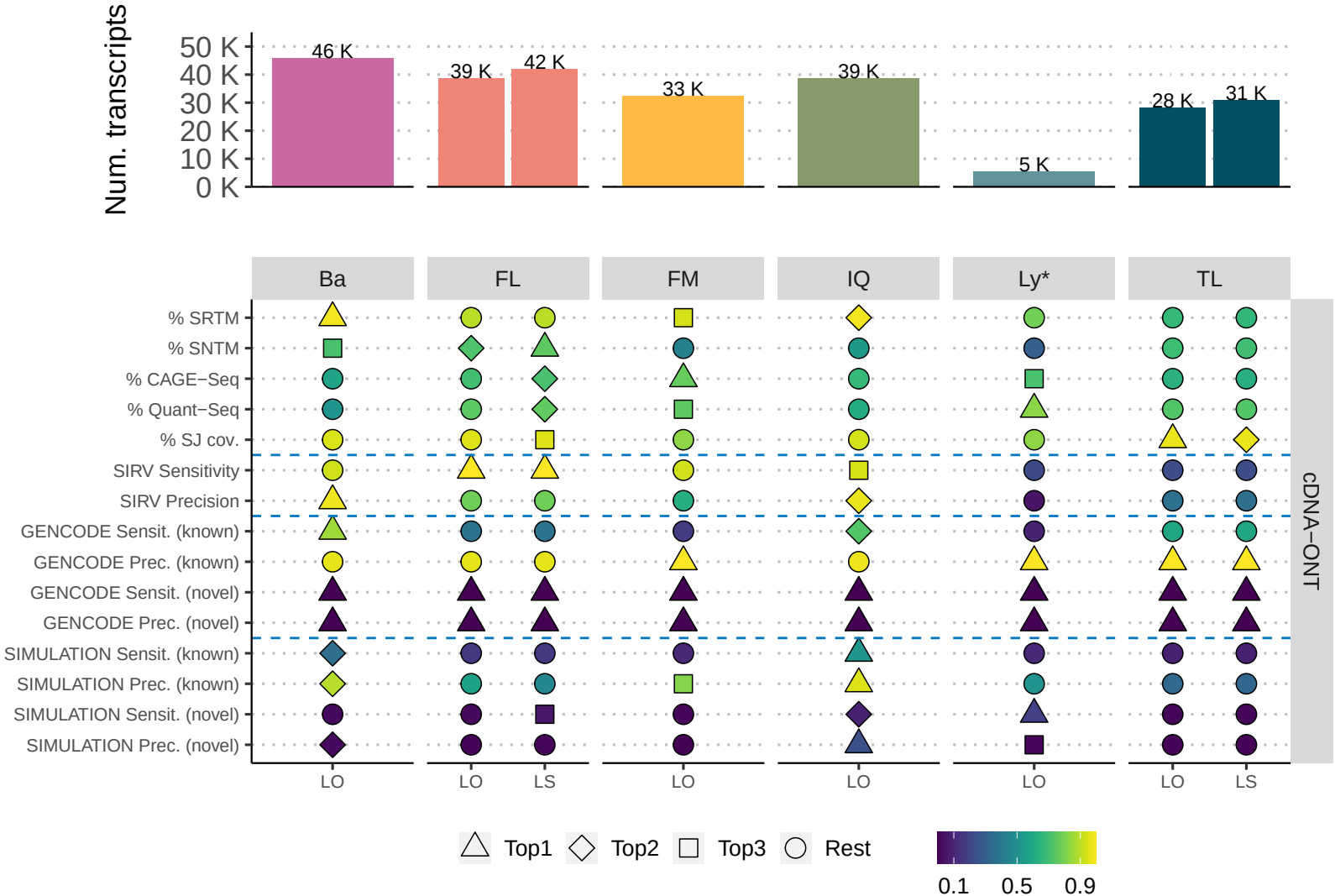
Num. transcripts
50 K
40 K
30 K
20 K
10 K
0 K
46 K
39 K
42 K
33 K
39 K
5 K
28 K
31 K
Ba
FL
FM
IQ
Ly*
TL
% SRTM
% SNTM
% CAGE−Seq
% Quant−Seq
% SJ cov.
SIRV Sensitivity
SIRV Precision
GENCODE Sensit. (known)
GENCODE Prec. (known)
GENCODE Sensit. (novel)
GENCODE Prec. (novel)
SIMULATION Sensit. (known)
SIMULATION Prec. (known)
SIMULATION Sensit. (novel)
SIMULATION Prec. (novel)
cDNA−ONT
LO
LO
LS
LO
LO
LO
LO
LS
Top1
Top2
Top3
Rest
0.1
0.5
0.9