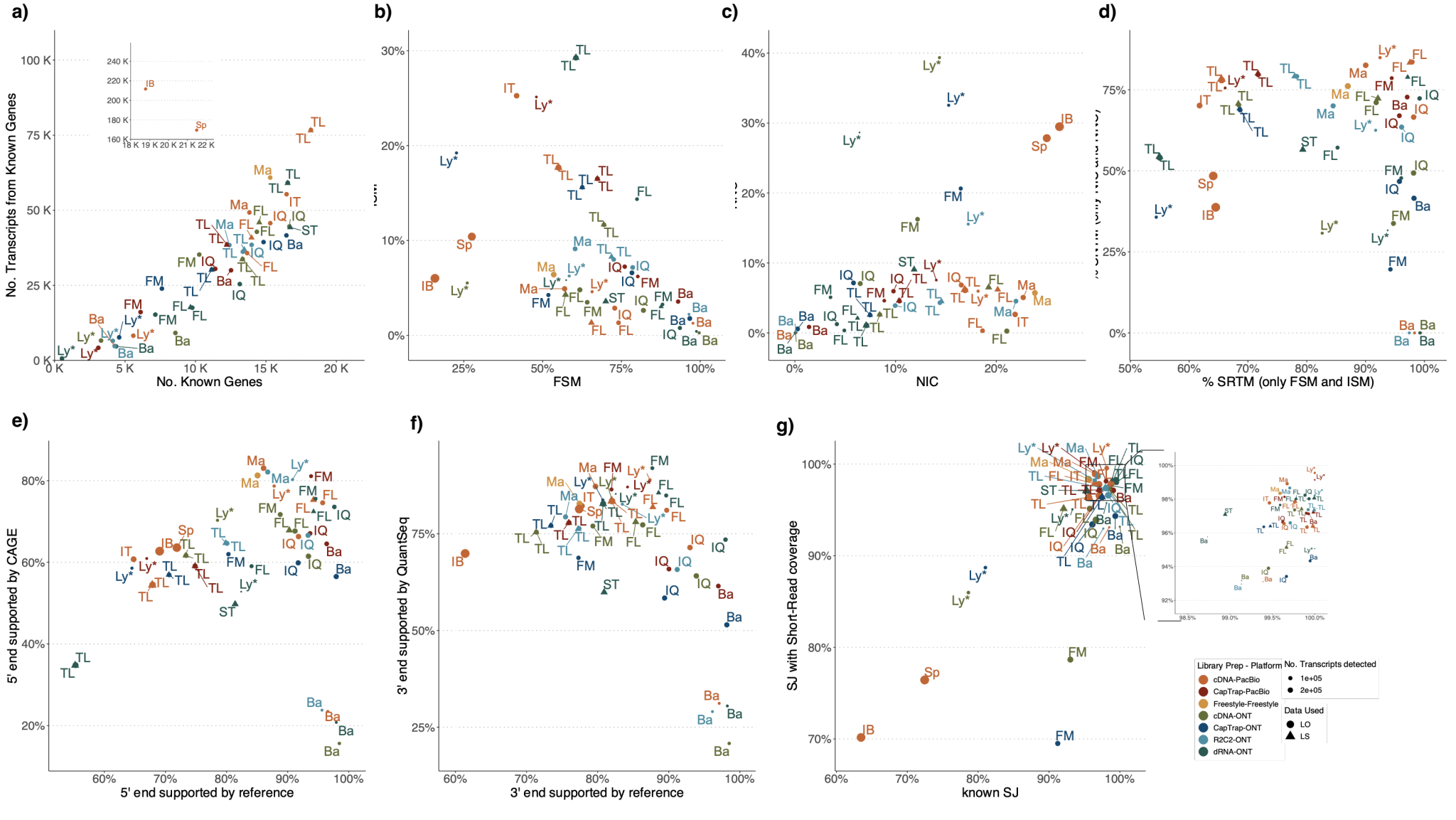
a)
No. Transcripts from Known Genes
No. Known Genes
b)
FSM
c)
NIC
d)
% SRTM (only FSM and ISM)
e)
5' end supported by CAGE
5' end supported by reference
f)
3' end supported by QuantSeq
3' end supported by reference
g)
SJ with Short-Read coverage
known SJ
Library Prep - Platform
cDNA-PacBio
CapTrap-PacBio
Freestyle-Freestyle
cDNA-ONT
CapTrap-ONT
R2C2-ONT
dRNA-ONT
No. Transcripts detected
1e+05
2e+05
Data Used
LO
LS