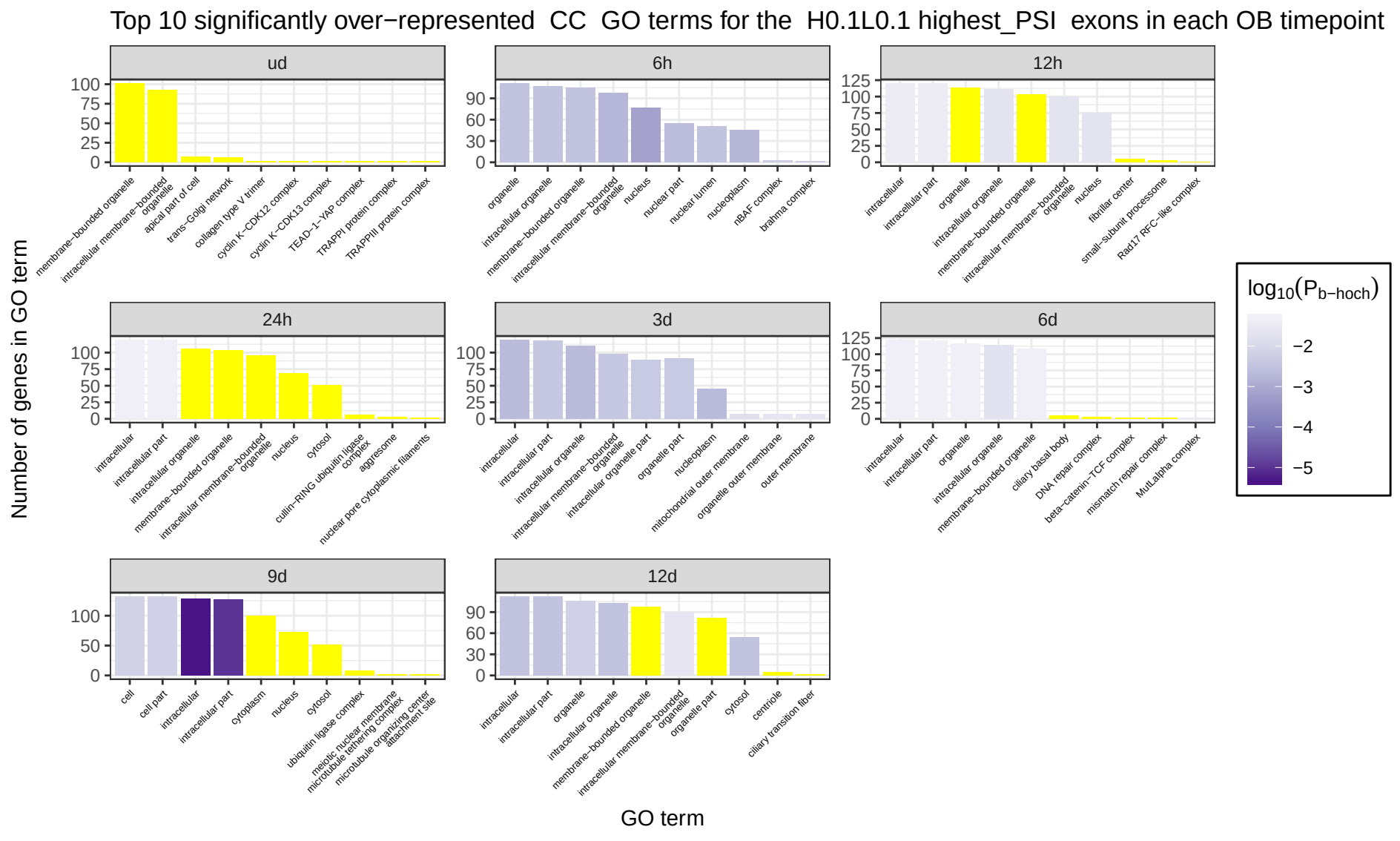
Top 10 significantly over−represented CC GO terms for the H0.1L0.1 highest_PSI exons in each OB timepoint
ud
membrane−bounded organelle
intracellular membrane−bounded organelle
apical part of cell
trans−Golgi network
collagen type V trimer
cyclin K−CDK12 complex
cyclin K−CDK13 complex
TEAD−1−YAP complex
TRAPPI protein complex
TRAPPII protein complex
6h
organelle
intracellular organelle
membrane−bounded organelle
intracellular membrane−bounded organelle
nucleus
nuclear part
nuclear lumen
nucleoplasm
nBAF complex
brahma complex
12h
intracellular
intracellular part
organelle
intracellular organelle
membrane−bounded organelle
intracellular membrane−bounded organelle
nucleus
fibrillar center
small−subunit processome
Rad17 RFC−like complex
24h
intracellular
intracellular part
intracellular organelle
membrane−bounded organelle
intracellular membrane−bounded organelle
nucleus
cytosol
cullin−RING ubiquitin ligase complex
aggresome
nuclear pore cytoplasmic filaments
3d
intracellular
intracellular part
intracellular organelle
intracellular membrane−bounded organelle
intracellular organelle part
organelle part
nucleoplasm
mitochondrial outer membrane
organelle outer membrane
outer membrane
6d
intracellular
intracellular part
organelle
intracellular organelle
membrane−bounded organelle
ciliary basal body
DNA repair complex
beta−catenin−TCF complex
mismatch repair complex
MutLalpha complex
9d
cell
cell part
intracellular
intracellular part
cytoplasm
nucleus
cytosol
ubiquitin ligase complex
meiotic nuclear membrane microtubule tethering complex
microtubule organizing center attachment site
12d
intracellular
intracellular part
organelle
intracellular organelle
membrane−bounded organelle
intracellular membrane−bounded organelle
organelle part
cytosol
centriole
ciliary transition fiber
Number of genes in GO term
GO term
log10(Pb−hoch)
−2
−3
−4
−5