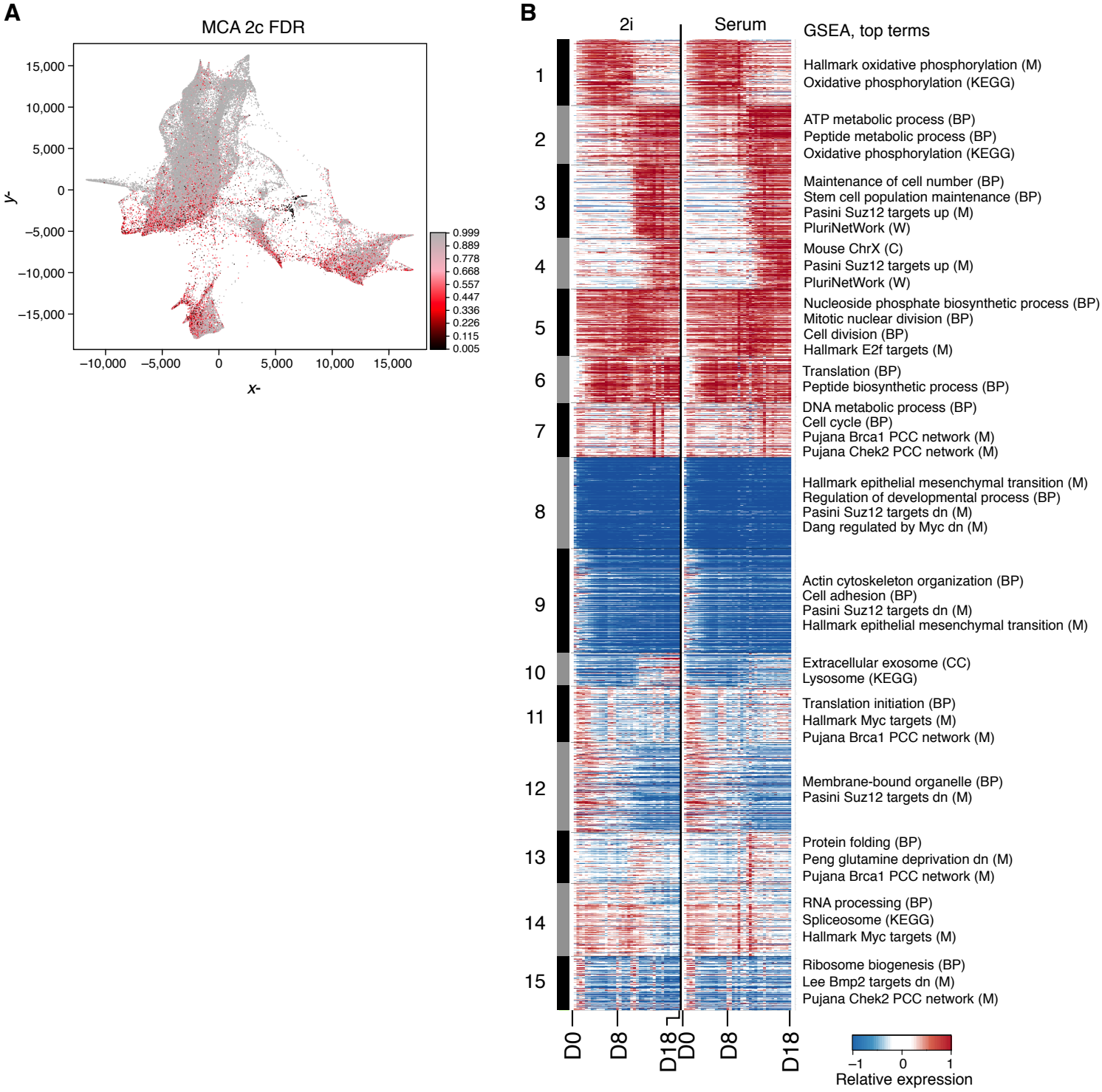
Figure S3


A
MCA 2c FDR
15,000
10,000
5,000
0
−5,000
−10,000
−15,000
y-
−10,000
−5,000
0
5,000
10,000
15,000
x-
0.999
0.889
0.778
0.668
0.557
0.447
0.336
0.226
0.115
0.005
B
2i
Serum
GSEA, top terms
1
Hallmark oxidative phosphorylation (M)
Oxidative phosphorylation (KEGG)
2
ATP metabolic process (BP)
Peptide metabolic process (BP)
Oxidative phosphorylation (KEGG)
3
Maintenance of cell number (BP)
Stem cell population maintenance (BP)
Pasini Suz12 targets up (M)
PluriNetWork (W)
4
Mouse ChrX (C)
Pasini Suz12 targets up (M)
PluriNetWork (W)
5
Nucleoside phosphate biosynthetic process (BP)
Mitotic nuclear division (BP)
Cell division (BP)
Hallmark E2f targets (M)
6
Translation (BP)
Peptide biosynthetic process (BP)
7
DNA metabolic process (BP)
Cell cycle (BP)
Pujana Brca1 PCC network (M)
Pujana Chek2 PCC network (M)
8
Hallmark epithelial mesenchymal transition (M)
Regulation of developmental process (BP)
Pasini Suz12 targets dn (M)
Dang regulated by Myc dn (M)
9
Actin cytoskeleton organization (BP)
Cell adhesion (BP)
Pasini Suz12 targets dn (M)
Hallmark epithelial mesenchymal transition (M)
10
Extracellular exosome (CC)
Lysosome (KEGG)
11
Translation initiation (BP)
Hallmark Myc targets (M)
Pujana Brca1 PCC network (M)
12
Membrane-bound organelle (BP)
Pasini Suz12 targets dn (M)
13
Protein folding (BP)
Peng glutamine deprivation dn (M)
Pujana Brca1 PCC network (M)
14
RNA processing (BP)
Spliceosome (KEGG)
Hallmark Myc targets (M)
15
Ribosome biogenesis (BP)
Lee Bmp2 targets dn (M)
Pujana Chek2 PCC network (M)
D0
D8
D18
D0
D8
D18
−1
0
1
Relative expression