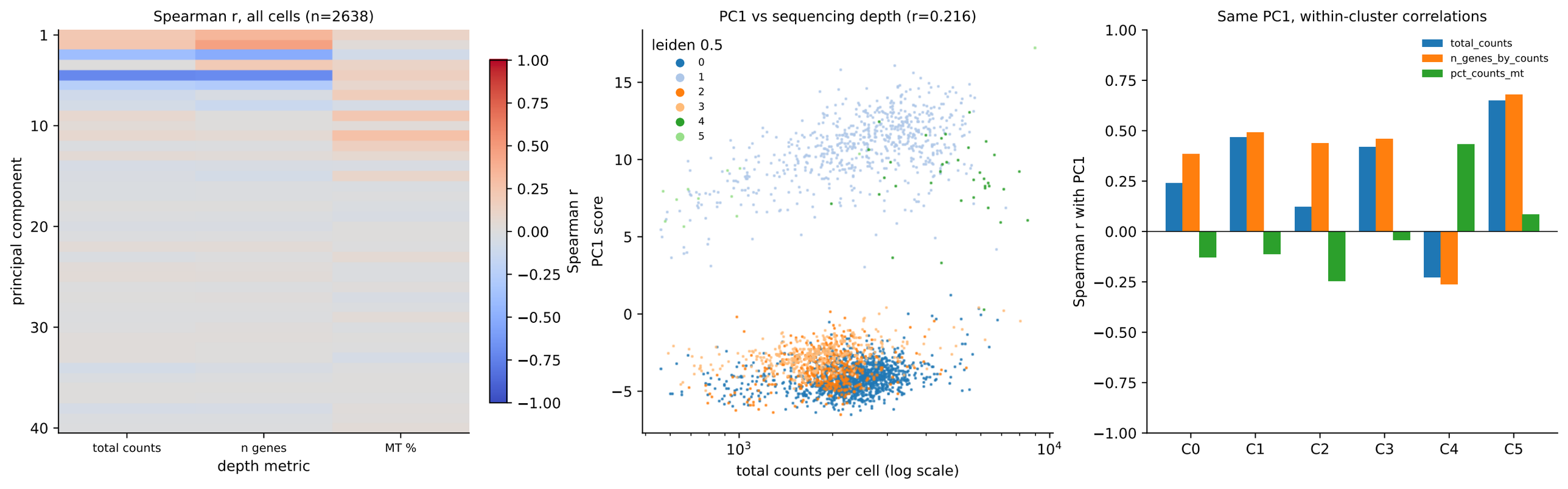

Spearman r, all cells (n=2638)
principal component
depth metric
total counts
n genes
MT %
Spearman r
PC1 vs sequencing depth (r=0.216)
PC1 score
leiden 0.5
total counts per cell (log scale)
Same PC1, within-cluster correlations
Spearman r with PC1
total_counts
n_genes_by_counts
pct_counts_mt
C0
C1
C2
C3
C4
C5