

Figure S5

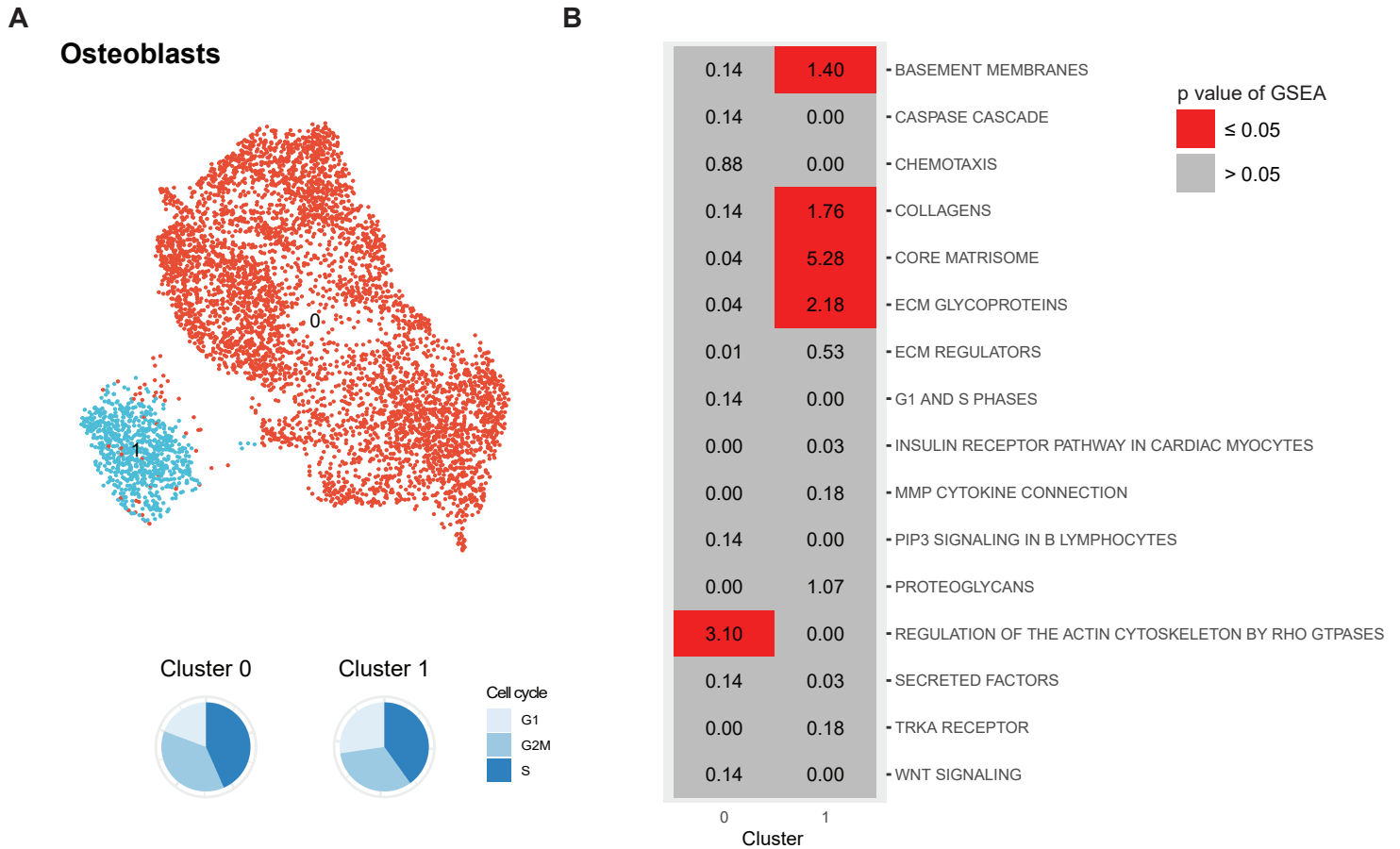


Figure S5. Osteoblast cells demonstrate phenotypic heterogeneity. A) UMAP analysis of normal human osteoblast cells in cell culture. B) Pathway enrichment analysis against msigdb C2 canonical pathways on genes differentially upregulated in each cluster relative remaining cells in the dataset.