



Supplemental Figure 9. Expression of cell aging genes. Heatmap based on microarray results showing the median-centered, log₂ expression values of genes associated with the GO term “cell aging” in H1 ESCs, untargeted OI iPSCs (iPSC-OI12-FVc3), four gene-targeted iPSC lines, three OI MSC lines, and five gene-targeted iMSC lines.