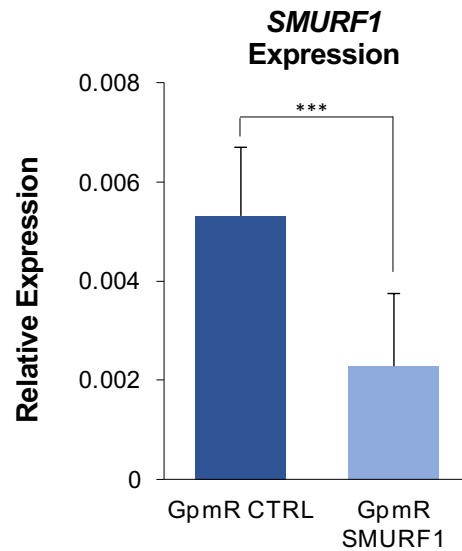
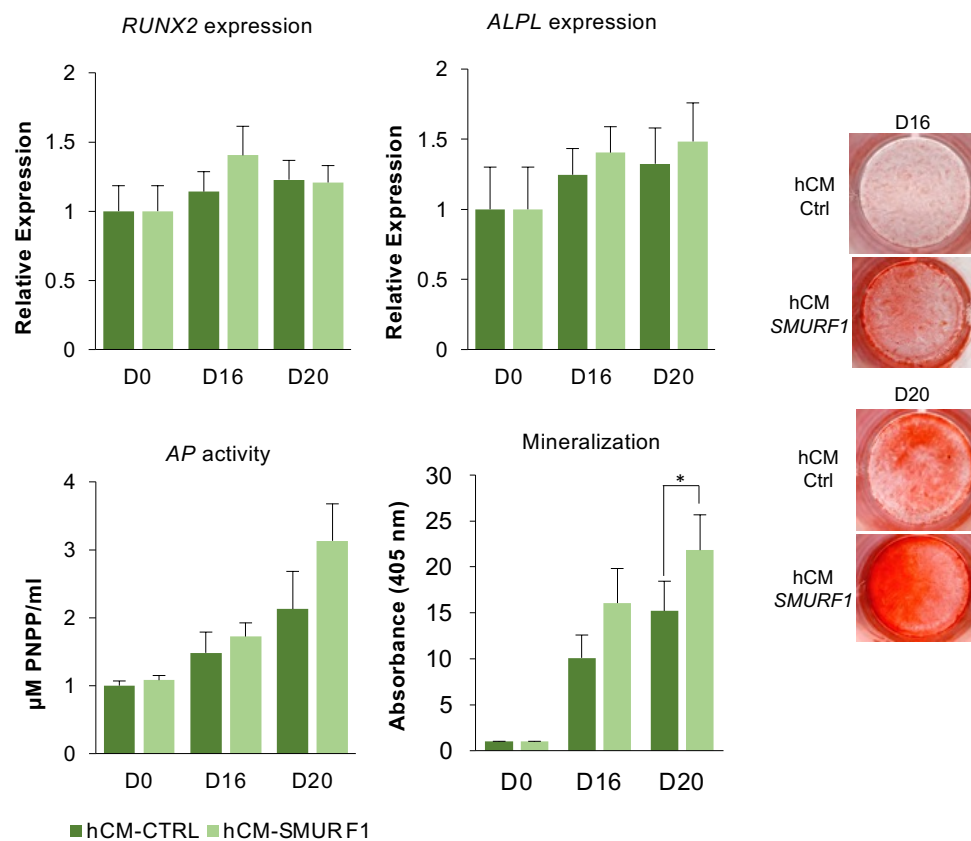
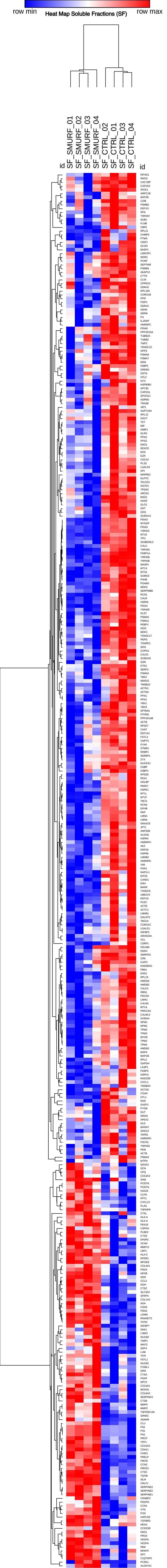


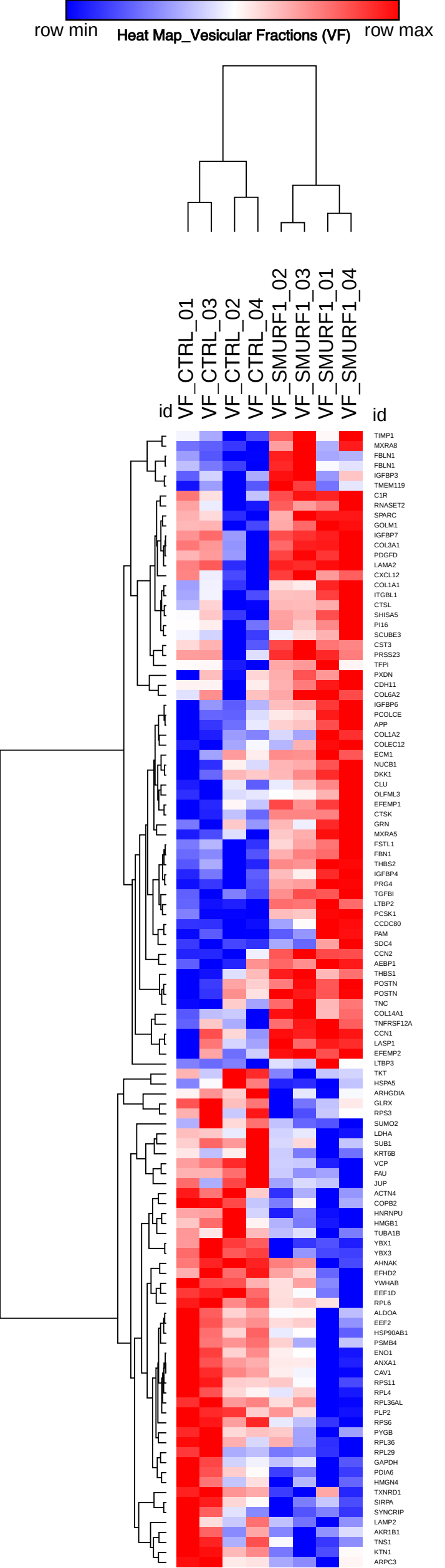


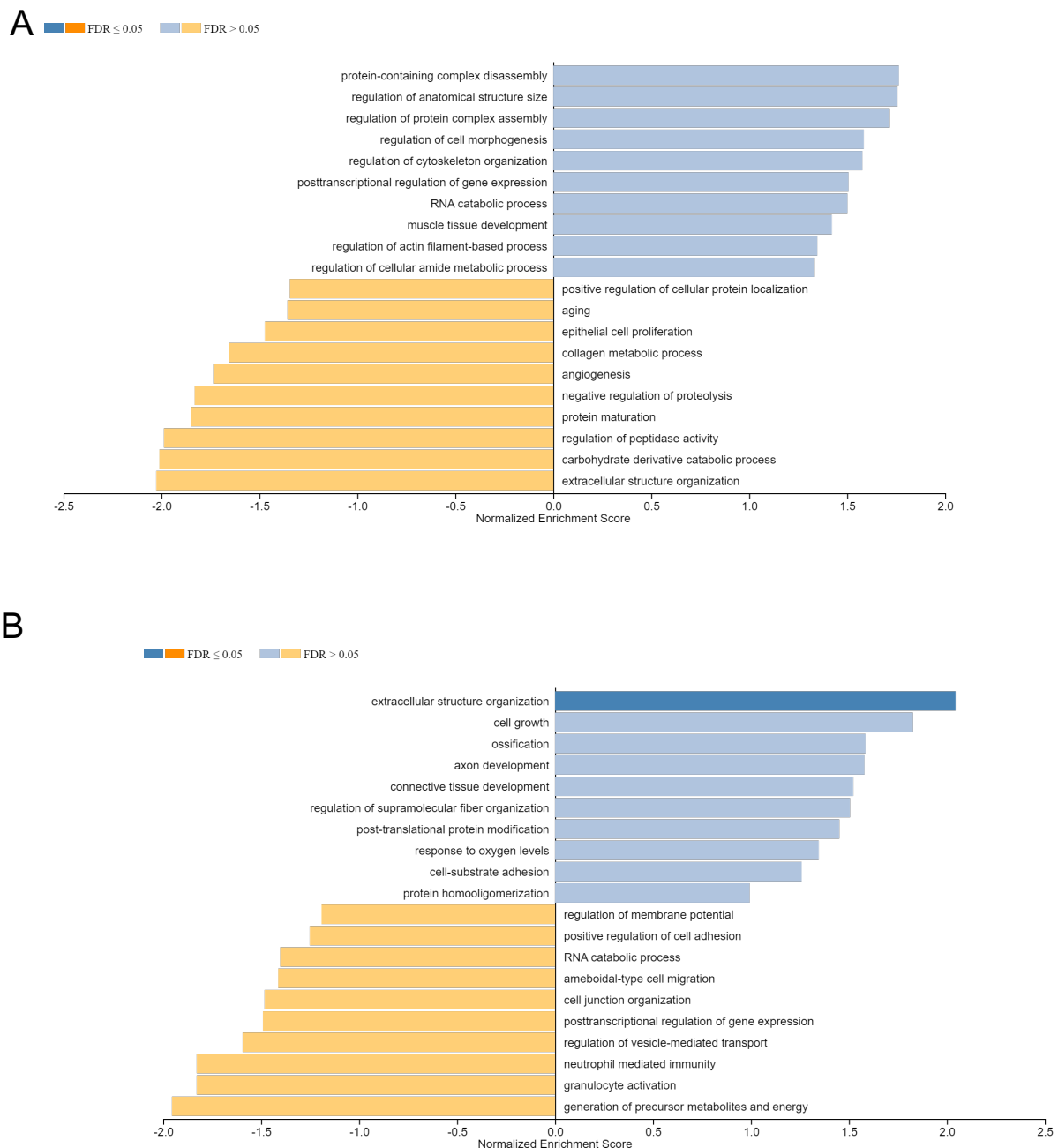
**Supplementary Figure 1. *SMURF1* silencing in ASC52telo human MSC cell line.** Graph shows *SMURF1* expression as seen by qPCR in cells transfected with 20nM of a control unspecific GapmeR or a GapmeR specifically designed for the silencing of *SMURF1* in human cells. Graph shows average results from three independent experiments. Error bars show Standard Error of the mean value. \*\*\* p-value <0.001. n=6.



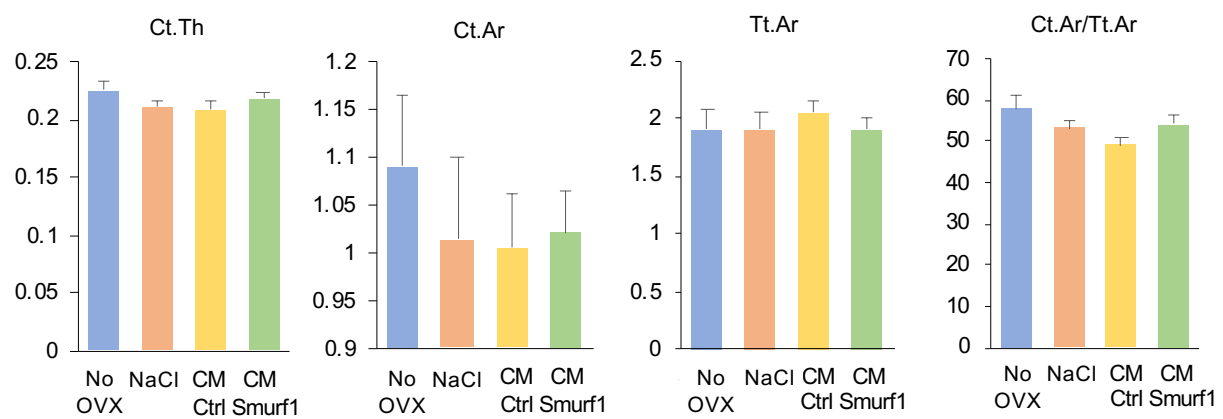
**Supplementary Figure 2.** ASC52telo osteogenic differentiation of 20 days following 48h-human Conditioned Media (hCM) preconditioning. No changes were observed in *RUNX2*. *ALPL* expresión and enzymatic activity show a non-significant increase in hCM-SMURF1. Statistically significant enhancemen in mineralization confirms osteogenic impact of hCM-SMURF1. Graph shows average results from four independent experiments . Error bars show Standard Error of the mean value. \* is p-value <0.05.







**Supplementary Figure 5.** Gene Ontology analysis of proteins significantly enriched in soluble and vesicular fractions of CM-SMURF1. A: Functions that can be associated with a regeneration processes like “regulation of cell morphogenesis” or “post-transcriptional regulation of gene expression” are identified in soluble fraction of CM-SMURF1 (blue bars). B: Functions suggesting a positive switch towards pro-osteogenic functions such as “extracellular matrix organization” or “ossification” are enriched in vesicular fraction of CM-SMURF1 (blue bars). Yellow bars highlights functions boosted in CM-Ctrl.



**Supplementary Figure 6.** Micro-CT cortical analysis reports no changes in any of the conditions under study. Cortical thickness (Ct.Th), Cortical Area (Ct.Ar), Total Area (Tt.Ar) and Cortical Area/Total Area (Ct.Ar/Tt.Ar). (n=5). Error bars show Standard Error of the mean value.