



Supplementary Figure 4 - UpSet plot (Lex et al., 2014) representing the number of intersections of differentially expressed genes (DEGs) amongst different datasets (Smith et al., 2012; CP_hamstrings, Smith et al., 2009; CP_flexors_wrist, Samani et al., 2022; miR-486_KO, and current study; C2C12 myoblast treated with miR-486 mimic). Horizontal bars represent the total number of DEG within the datasets. The vertical bars represent the number of DEG in the highlighted comparisons (filled black circles along the x axis). A line linking multiple filled circles represents a column in which those DEGs are shared among more than one group comparison.