Description of Additional Supplementary Files

File Name: Supplementary Data 1
Description: Number of individuals in "old" and "young" cohorts

File Name: Supplementary Data 2
Description: Names of GTEx tissues with matching abbreviations used within figures

File Name: Supplementary Data 3
Description: Genes with significant age-related gene expression heteroscedasticity

File Name: Supplementary Data 4
Description: Genes with nonzero age-sex interaction term

File Name: Supplementary Data 5
Description: GO enrichment of $R^2$ of age across all tissues

File Name: Supplementary Data 6
Description: Tissues with matching COSMIC query and number of sequenced tumor samples