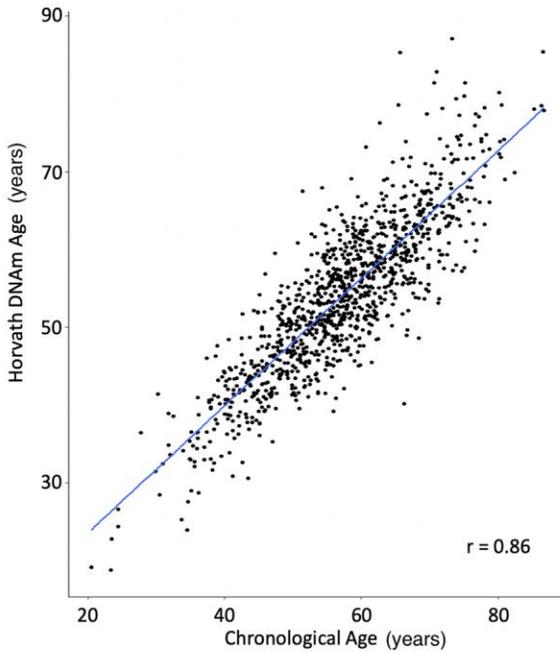
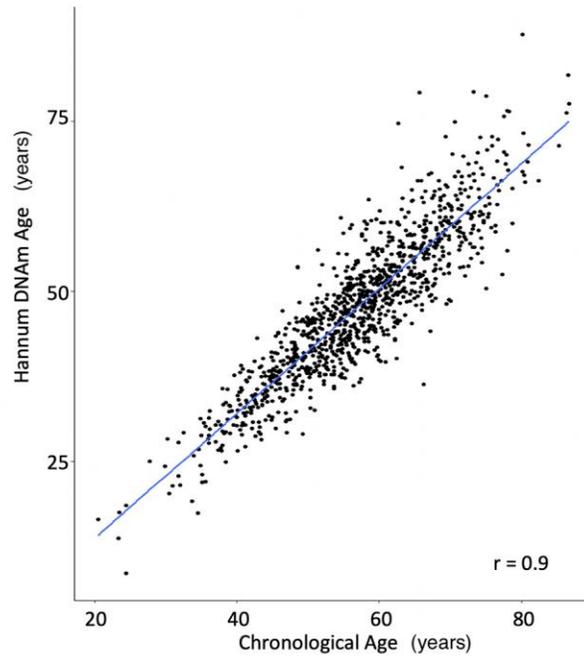


SUPPLEMENTARY FIGURES

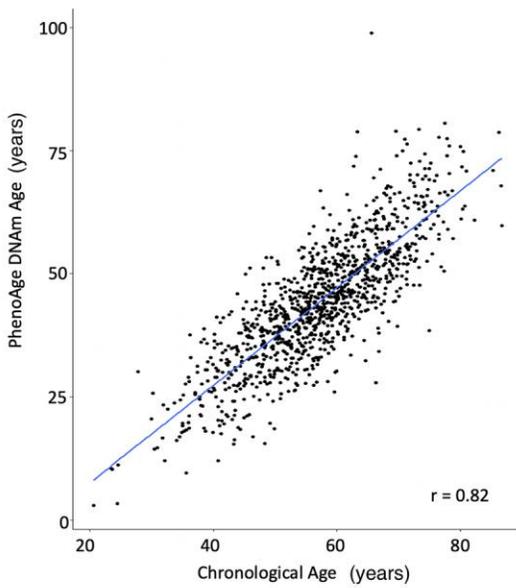
**A** HorvathAge



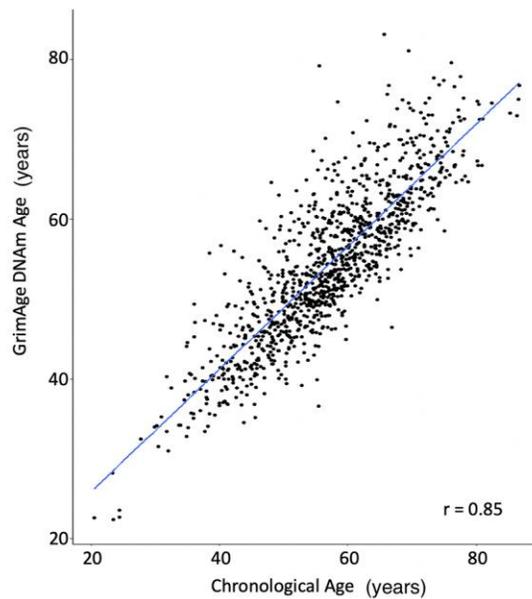
**B** HannumAge



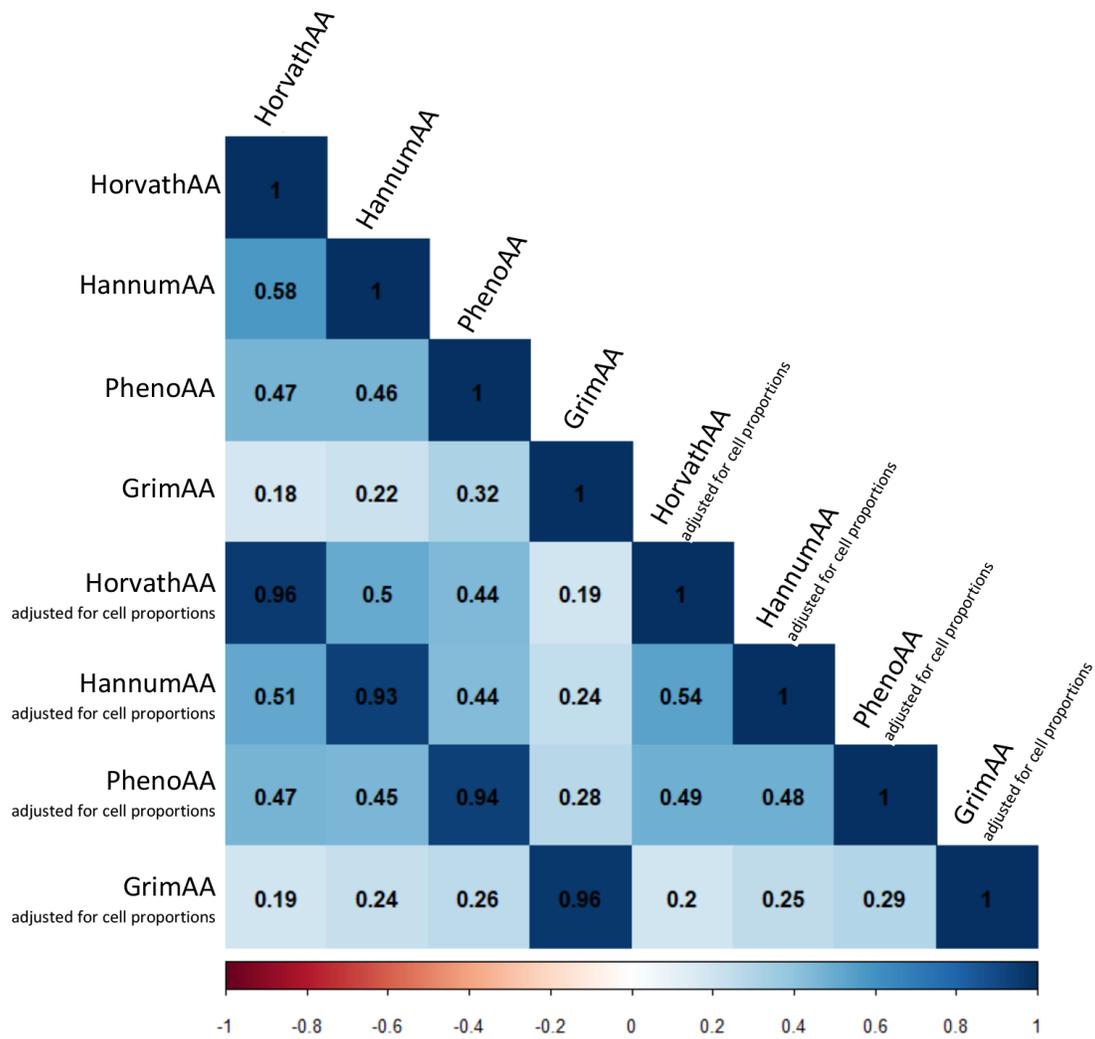
**C** PhenoAge



**D** GrimAge



**Supplementary Figure 1. Correlation between chronological age and DNAm age estimated by four epigenetic clocks.** Scatterplots with Pearson correlation coefficients between chronological age and DNA methylation age estimated by the HorvathAge (A), HannumAge (B), PhenoAge (C), and GrimAge (D) clocks. DNAm: DNA methylation; r: Pearson correlation coefficient.



Supplementary Figure 2. Pearson correlation coefficients for DNA methylation age acceleration metrics estimated by four epigenetic clocks.