Supplementary Table S1. Studies investigating the association of BMI and Horvath's DNA methylation aging markers.

| Study | Population | Sample Size | Aging Biomarker | Tissue | Type of Estimate | Effect Size | Test Statistics |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $\begin{gathered} \text { Simpkin AJ, } \\ 2017 \end{gathered}$ | British women | 790 | $\Delta$ Age | buccal | $\beta$ coefficient | 0.085 years per $\mathrm{kg} / \mathrm{m} 2$ | $\begin{gathered} \text { 95\% CI: }(0.014, \\ 0.156) \end{gathered}$ |
| $\begin{gathered} \text { Simpkin AJ, } \\ 2017 \end{gathered}$ | British women | 152 | $\Delta$ Age | blood | $\beta$ coefficient | 0.044 years per $\mathrm{kg} / \mathrm{m} 2$ | $\begin{gathered} 95 \% \text { CI: }(-0.065, \\ 0.15) \end{gathered}$ |
| $\begin{gathered} \text { Horvath S, } \\ 2014 \end{gathered}$ | German | 141 | Chronological age-adjusted DNAm Age | liver | $\beta$ coefficient | 0.168 year per $\mathrm{kg} / \mathrm{m} 2$ | $\mathrm{SE}=0.046$ |
| Quach A, 2017 | African American postmenopausal women | 1058 | DNAm AA (EEAA) | blood | $\beta$ coefficient | $\begin{gathered} 0.11 \text { years per } \\ \mathrm{kg} / \mathrm{m} 2 \end{gathered}$ | $\mathrm{p}=0.003$ |
| Quach A, 2017 | African American postmenopausal women | 1058 | DNAm AA (IEAA) | blood | $\beta$ coefficient | $\begin{gathered} 0.04 \text { years per } \\ \mathrm{kg} / \mathrm{m} 2 \end{gathered}$ | $\mathrm{p}=0.17$ |
| $\begin{gathered} \text { Dugue PA, } \\ 2018 \end{gathered}$ | Overweight Australian adults vs. Lean adults | 1384 | DNAm AA | blood | $\beta$ coefficient | 0.40 years | $\mathrm{SE}=0.31$ |
| $\begin{gathered} \text { Dugue PA, } \\ 2018 \end{gathered}$ | Obese I Australian adults vs. Lean adults | 451 | DNAm AA | blood | $\beta$ coefficient | 0.15 years | $\mathrm{SE}=0.42$ |
| $\begin{gathered} \text { Dugue PA, } \\ 2018 \end{gathered}$ | Obese II Australian adults vs. Lean adults | 105 | DNAm AA | blood | $\beta$ coefficient | 2.38 years | $\mathrm{SE}=0.73$ |
| $\begin{gathered} \text { Nevalainen T, } \\ 2017 \end{gathered}$ | young adults | 183 | DNAm Age | blood | correlation coefficient | 0.11 | $\mathrm{p}=0.138$ |
| $\begin{aligned} & \text { Nevalainen T, } \\ & 2017 \end{aligned}$ | middle-aged adults | 183 | DNAm Age | blood | correlation coefficient | 0.218 | $\mathrm{p}=0.0001$ |
| $\begin{gathered} \text { Nevalainen T, } \\ 2017 \\ \hline \end{gathered}$ | nonagenarian individuals | 119 | DNAm Age | blood | correlation coefficient | -0.115 | $\mathrm{p}=0.211$ |

Abbreviations: CI, confidence interval; BMI, body mass index; $\triangle$ Age, the discrepancy between DNA methylation age and chronological age; DNAm AA, DNA methylation age acceleration, the residual resulting from regression DNA methylation age on chronological age in a linear model; $\beta$, regression coefficient from investigating the relationship between BMI and Horvath's DNA methylation aging markers; r, Spearman's rank correlation coefficient from investigating the relationship between BMI and Horvath's DNA methylation aging markers.

