**Table 2.** Regression models linking number of pregnancies to telomere length (models 1-4) and DNAmAge (models 5-8).

|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
|  | **Telomere Length~** | | | | **DNAmAge~** | | | |
|  | (1) | (2)† | (3)† | (4)† | (5) | (6)† | (7)† | (8)† |
| **Chronological Age (yrs.)** | -0.047\*\* | -0.029+ | -0.028+ | -0.029+ | 0.485 | 0.667 | 0.656 | 0.645 |
| **No. Pregnancies** | -0.014\* | -0.013\* | -0.014\* | -0.016\* | 0.363\* | 0.326\* | 0.459\*\* | 0.510\*\* |
| **SES** |  | -0.006 | -0.006 | -0.004 |  | -0.180 | -0.214+ | -0.291+ |
| **Currently Pregnant** |  |  | 0.011 | 0.011 |  |  | -1.472\*\* | -1.460\*\* |
| **No. Pregnancies x SES** |  |  |  | -0.004 |  |  |  | 0.106 |
| **Intercept** | 1.826\*\* | 1.337\*\* | 1.332\*\* | 1.343\*\* | 14.818 | 10.319 | 10.611 | 10.850 |
| **Observations** | 821 | 821 | 821 | 821 | 397 | 397 | 397 | 397 |
| **Adjusted R2** | 0.015 | 0.063 | 0.062 | 0.062 | 0.011 | 0.041 | 0.067 | 0.067 |

†Includes controls for principal components of genetic variation to control for population structure, and average urbanicity (complete results shown in Table S2). +p<0.1;\*p<0.05;\*\*p<0.01;\*\*\*p<0.001