

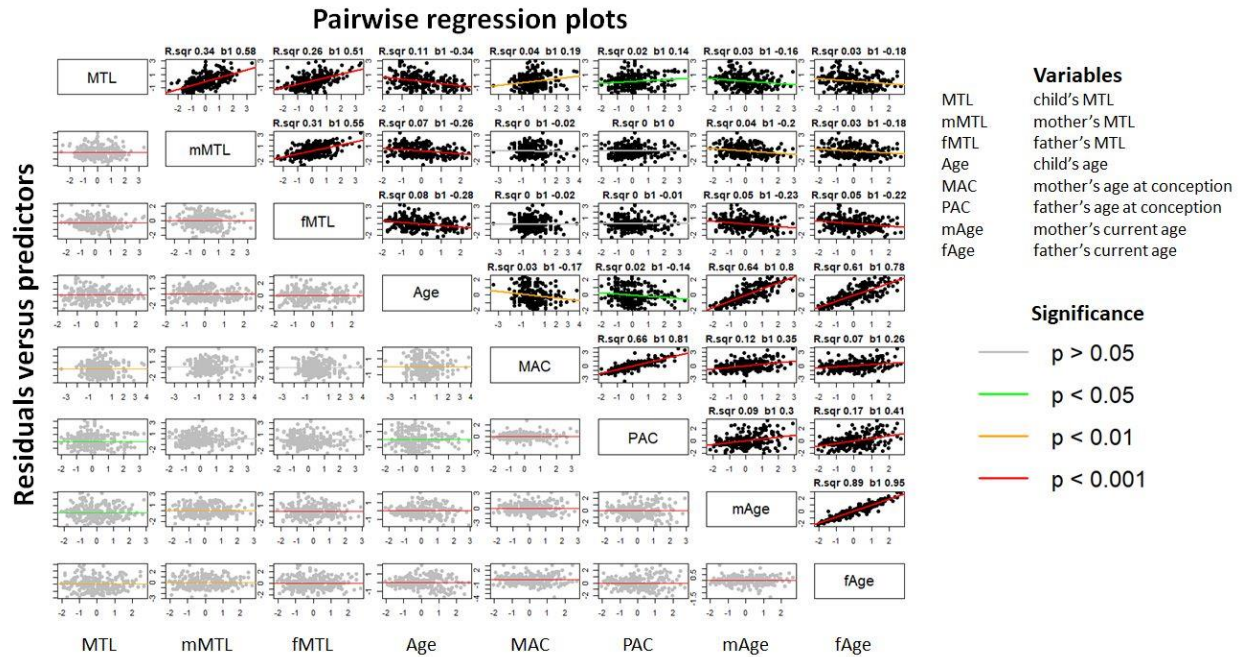
## Supplementary document 1

### **WGS-based telomere length analysis in Dutch family trios implicates stronger maternal inheritance and a role for RRM1 gene**

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**Figure S1. Pairwise correlations between offspring mean telomere length (MTL) and meta-variables.**

The right triangle shows correlation plots with fitted correlation lines and linear regression statistics (R squared and correlation coefficient). The left triangle shows residuals of each correlation with respect to the predictive variable. We see that mother's and father's ages, as well as PAC and MAC show strong correlation, which should be considered in multiple linear regression analysis (see the main text for details).