Sup. Fig.3: DNA methylation at three CpG sites in the MGC33302 gene (also known as MFSD8/CLN7) was correlated with paternal stress during preschool in girls only.

Three CpG sites in the first intron of the MGC33302 (alias MFSD8, encoding a putative lysosomal transporter) were negatively associated in girls with paternal stress at preschool. Specifically, these included the HCS site cg03482087 (*rho* = - .61, -5.4% DNA methylation change, slope = -.08 (95% CI = -.11 - -.04), FDR = 0%), and the 2 MCS sites cg03883519 (*rho* = - .41, -6.4% DNA methylation change, slope = -.09 (95% CI = -.18 - 0), FDR = 7.4%) and cg15557833 (*rho* = - .34, -5.8% DNA methylation change, slope = -.08 (95% CI = -.15 - -.02), FDR = 13.7%). No statistically significant association was found in the full group or boys-only analysis