

Supplemental Table 6. Multivariate linear models of fat mass (a), lean mass (b) and body mass index (c) in late infancy.

(a)						
	Total fat (g) at 12 months			Total fat (g) at 24 months		
	Beta	P	R ²	Beta	P	R ²
<i>NKX6.1</i> methylation	-0.456	0.002	21%	--	--	--
<i>GPR120</i> methylation	--	--	--	-0.693	0.002	33%
(b)						
	Lean mass (g) at 12 months			Lean mass (g) at 24 months		
	Beta	P	R ²	Beta	P	R ²
<i>SLC2A8</i> methylation	0.517	<0.0001	25%	--	--	--
<i>ONECUT1</i> methylation	--	--	--	0.537	0.001	29%
(c)						
	BMI Z-score Δ 0-24 months					
	Beta	P	R ²			
<i>SLC2A8</i> methylation	-0.661	<0.0001	42%			

(a) Non-predictive variables: *CHGA* methylation, *FABP5* methylation, *CTRP1* methylation, *CPT1A* methylation, *GAS6* methylation, *IGFBP4* methylation, *ONECUT1* methylation and *SLC2A8* methylation.

(b) Non-predictive variables: *NKX6.1* methylation, *GPR120* methylation, *CHGA* methylation, *FABP5* methylation, *CTRP1* methylation, *CPT1A* methylation, *GAS6* methylation and *IGFBP4* methylation.

(c) Non-predictive variables: *NKX6.1* methylation, *GPR120* methylation, *CHGA* methylation, *FABP5* methylation, *CTRP1* methylation, *CPT1A* methylation, *GAS6* methylation, *IGFBP4* methylation and *ONECUT1* methylation.