Pregnant mother recruited (Central Taiwan; n=430)

117 excluded (non-compliance)

Urine sample collected (third trimester; n=313)

17 excluded (non-compliance)

Live birth (n=313)

Loss to follow up (n=5)
From 9 twins one was chosen randomly and other not included (n=9)

Cord blood samples collected (newborns; n=299)

Cord blood DNA-methylation (n=64 available; 485,577 CpGs)

Final DNA methylation data (n=64; 385,183 CpGs)

Measuring DNA methylation (Illumina 450K)

Figure S1. Subject recruitment and preprocessing of DNA methylation data in Taiwanese birth cohort