Errata "Genome wide methylation analysis and obesity related traits" Emil Nilsson

Page List of papers	
	The status of paper IV is now "Submitted to SLEEP" Title of paper II is "Roux-en Y gastric bypass surgery induces genome-wide promoter-specific DNA methylation in whole blood of obese patients"
Abbreviations	BMIQ-Beta MIxture Quantile dilation
p. 11	Second sentence refers to obesity
p.12	Leptin levels decrease after sleep deprivation
p.12	hypoxia-inducible factor 3 alpha with lower case H
p.13	Heritability of circulating transcription factors should reference: Robert A S Ariëns, Marlies de Lange, Harold Snieder, May Boothby, Tim D Spector, Peter J Grant, Activation markers of coagulation and fibrinolysis in twins: heritability of the
p.14	prethrombotic state, <i>Lancet</i> 2002; 359: 667–71
p.19	First sentence refers to cytosine.
	The consort flowchart is found in paper II as figure 1. Weight was measured while subjects were wearing light clothes.
p.20	In paper I and II the ratio between the signal from the C and the sum of the C and T signals, was performed with the GenomeStudio 2009.2 (Illumina) software.
p.21	The experiment was carried out on a laptop on day 2 of the trial, both before and after the standardized breakfast.  In the third sentence, downstream is wrongfully capitalized.  In the middle of the page, "In addition an average", a comma is missing.
p.26	The supplementary table 1 is found in paper IV.  The word "maximum" should be omitted in the fourth sentence of the second section of the page.
p.27	First sentence: WNT related probes
p.29	Notch, bunched, and wnt signalling schould not be capitalized.
p.32	Second to last sentence: "Analysen visade att"