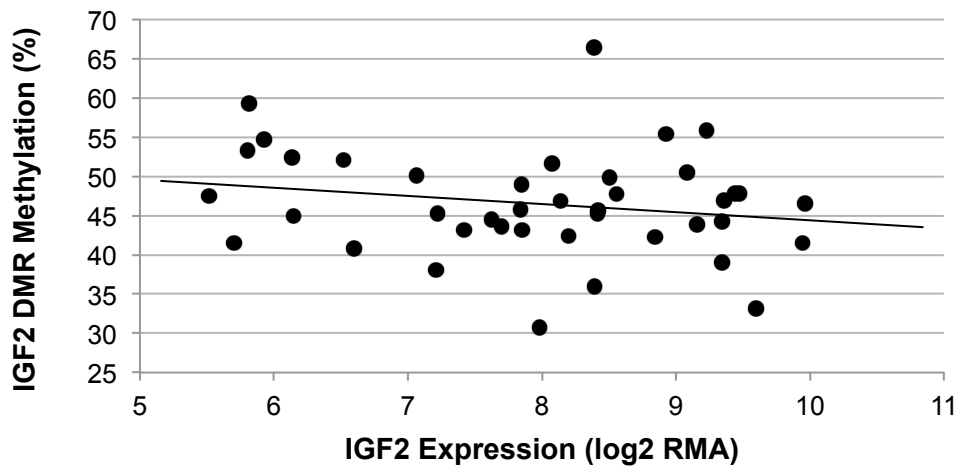


Supplementary Figure 1



Relationship between *IGF2* expression from Affymetrix U133 HTA microarray analysis, as \log_2 -transformed RMA normalized values, and *IGF2* DMR methylation measured by pyrosequencing ($\rho = -0.20$).