Supplementary Figure 1:

Legend: No correlation is seen between beta values from the Illumina Infinium array and % methylation from pyrosequencing (Pearson’s P=0.6, coefficient=0.1, N=12).

Supplementary Figure 2:

Legend: Box plots showing no significant change in RGN methylation levels in the BE-BED-EAC sequence. N(BE) = 13, N(BED) = 6, N(EAC) = 23. (BE=Barrett’s esophagus, BED=Barrett’s esophagus with dysplasia, EAC=esophageal adenocarcinoma).