**Figure legends**

**Supplementary Figure 1:** Methylation levels of *CDO1* (a), *SOX17* (b), and *TAC1* (c) measured in 11 pairs of tumours and adjacent normal tissues from NSCLC patients. Methylation levels are normalised to the reference gene *ACTB* and presented as square root Ct ratios. Outliers are indicated by bold circles located outside the whiskers of the boxplot. Statistical significance was computed using the nonparametric paired samples Wilcoxon test. \**p* < 0.10 (suggestive evidence), \*\**p* < 0.05 (moderate evidence).