Figure S1 Differential methylation levels of HpyCH4IV sites and the corresponding MATscores. Differential methylation levels of HpyCH4IV sites for which the relative hypomethylation levels were analyzed using the D-REAM analysis. For each T-DMR, the corresponding MATscore is indicated along the horizontal axis. In each tissue comparison, the lowest MATscore of the HpyCH4IV sites, whose hypomethylation were confirmed using COBRA, was considered as a threshold value for detecting hypomethylated regions in the tissue comparison.