

Table S5. Thresholds applied in EpiClass analysis of WGBS data from plasma

Sample cohort	Marker	Methylation density ^a
Lung cancer ^a	GALR1	35%
	TLX1	10%
	ZNF154	10%
Hepatocellular carcinoma ^b	GALR1	15%
	TLX1	0%
	ZNF154	35%

^a The methylation density threshold was determined based on a +/- 600bp interval on each side of the CpG of interest

^b The methylation density threshold was determined based on a +/- 500bp interval on each side of the CpG of interest