

Table S5

MA	coverage > 3				posteriorMax >=0.99			
	AIIC	CG	CHG	CHH	AIIC	CG	CHG	CHH
MA1_1	13417233	2344610	2513665	8558958	23428047	3030753	3797292	16600002
MA1_3	25009841	3450911	3937253	20731677	29141823	3753716	4183152	21204955
MA3	25679815	3086050	3542387	19051378	24357974	3655356	4131622	16570996

Previous methods
MethylStar

Table S5: Pre-processing of WGBS data using MethylStar increases the number of high-confident cytosines that can be used for epimutation analysis compared with previous pre-processing approaches.