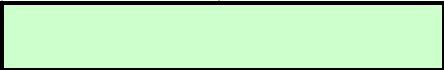

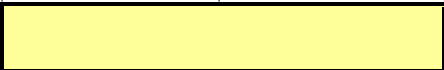




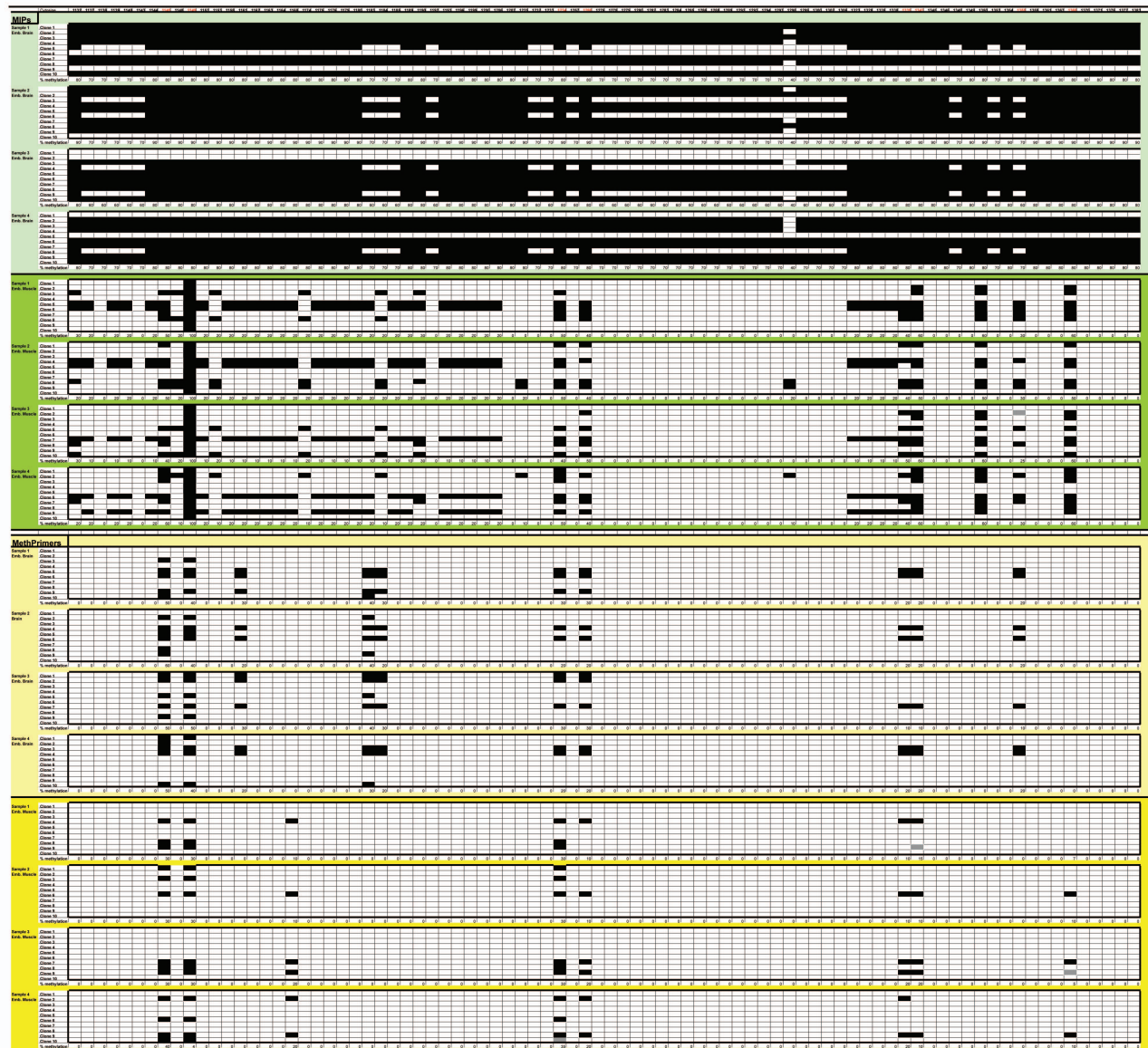


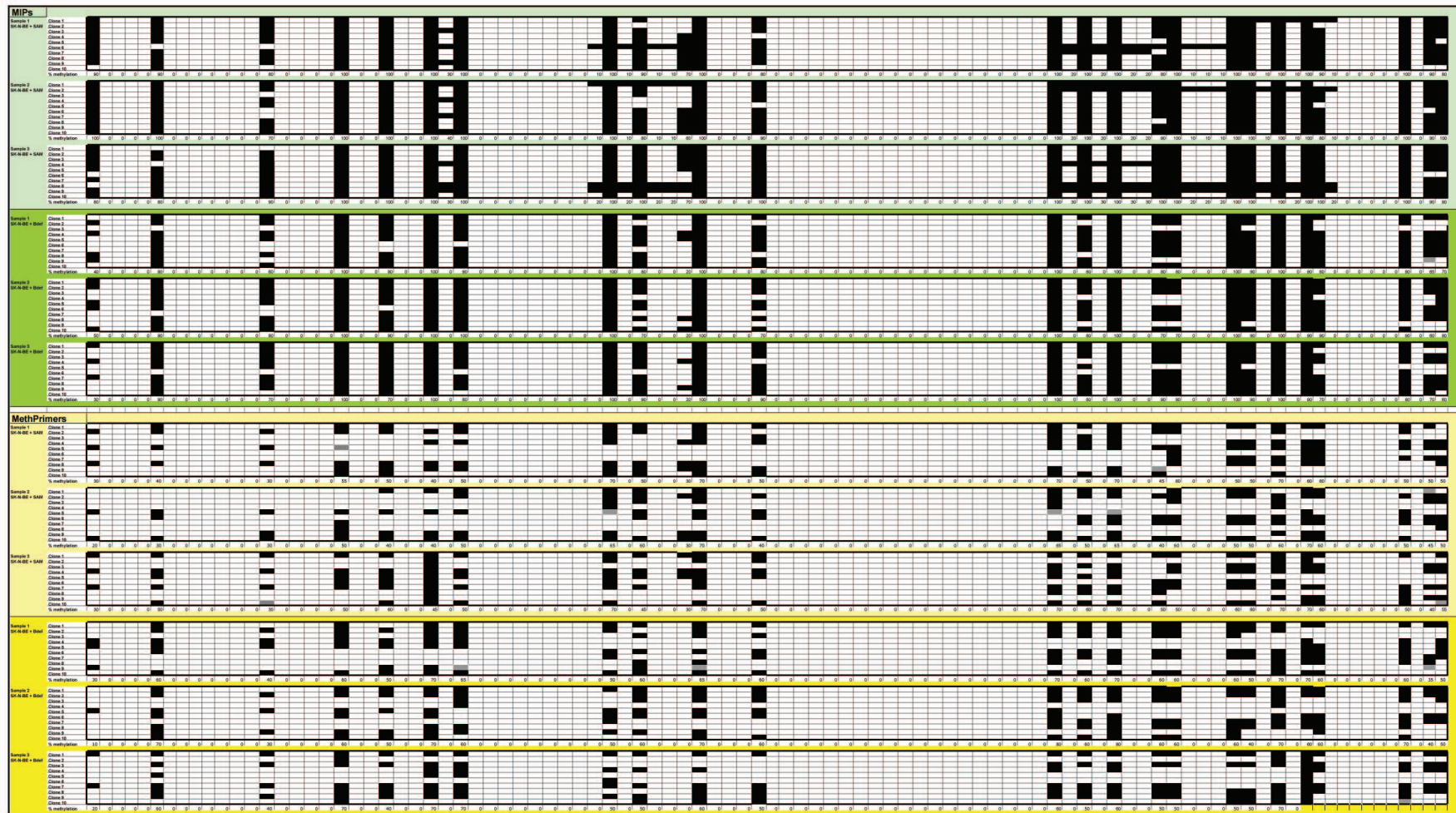
Supplementary Figure 5 graphic legend		
		Light green: High methylated conditions / analyzed by MIPs
		Dark green: Low methylated conditions / analyzed by MIPs
		Light yellow: High methylated conditions / analyzed by MethPrimers
		Dark yellow: Low methylated conditions / analyzed by MethPrimers
		Black box: methylated cytosine
		White box: unmethylated cytosine
		Grey box: undetermined
	<b>"Clone":</b>	Positive bacterial colonies sequenced per sample
	<b>"% methylation":</b>	% methylation for each cytosine in each sample
	<b>"Cytosine":</b>	position (bp) of cytosines in the promoter sequence (CpGs in red)



# Supplementary Figure 5b



# Supplementary Figure 5c



# Supplementary Figure 5d

