

Fig. S3 Bisulfite sequencing of the M10 fragment

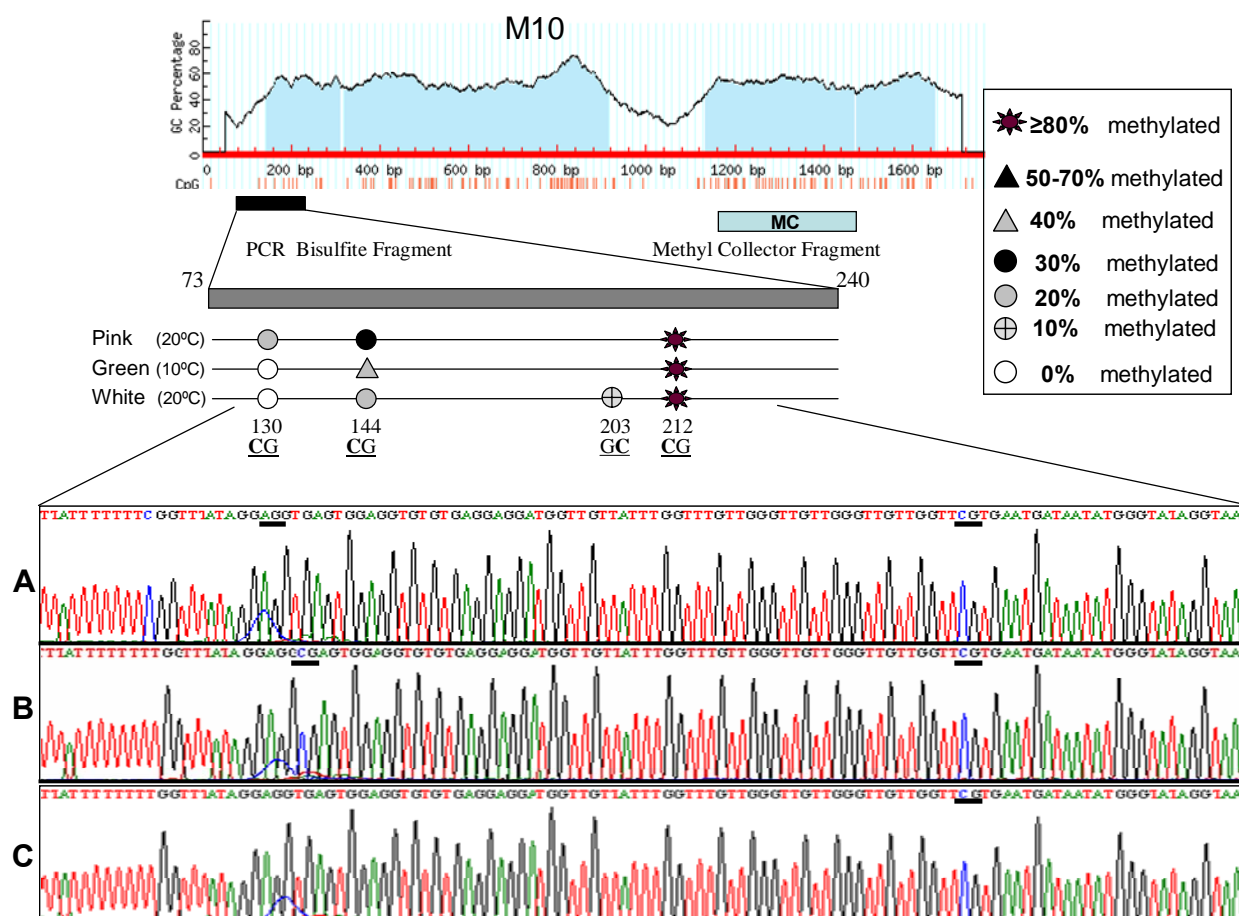


Figure S3 Bisulfite sequencing of the extended region of the affinity precipitated M10 fragment gi|171789667: 1315-1713 Contig56007. Prediction of the CpG Island and primers designed for the bisulfite sequencing was performed by the MethPrimer program <http://www.urogene.org/methprimer/index1.html>. Quantification of the methylation percentage was calculated by direct bisulfite sequencing of 10 independent clones.