

Fig. S4 Bisulfite sequencing for B19 fragment

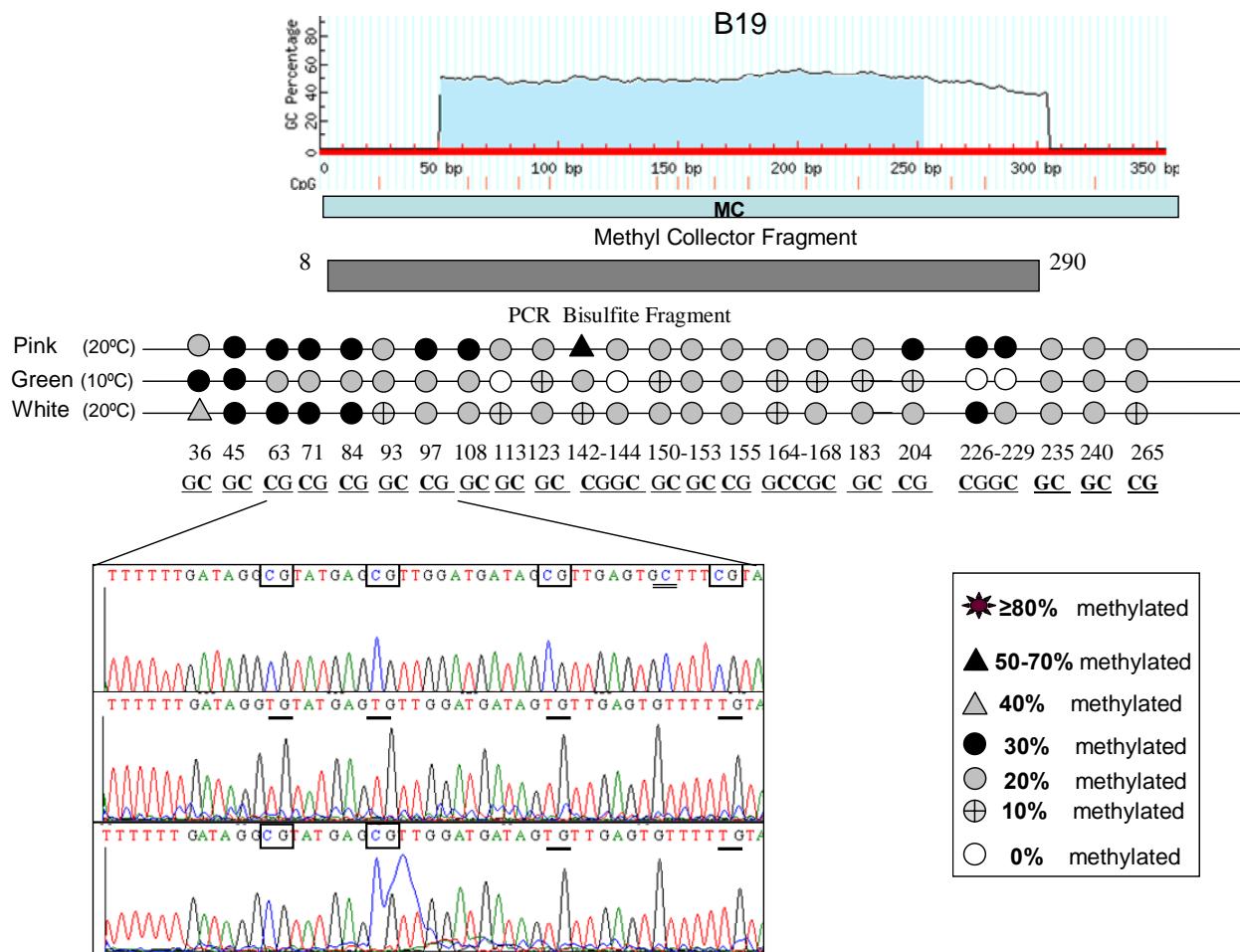


Figure S4. Bisulfite sequencing of the affinity precipitated B19 fragment

Prediction of the CpG Island and the design of primers for bisulfite sequencing were performed using 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.