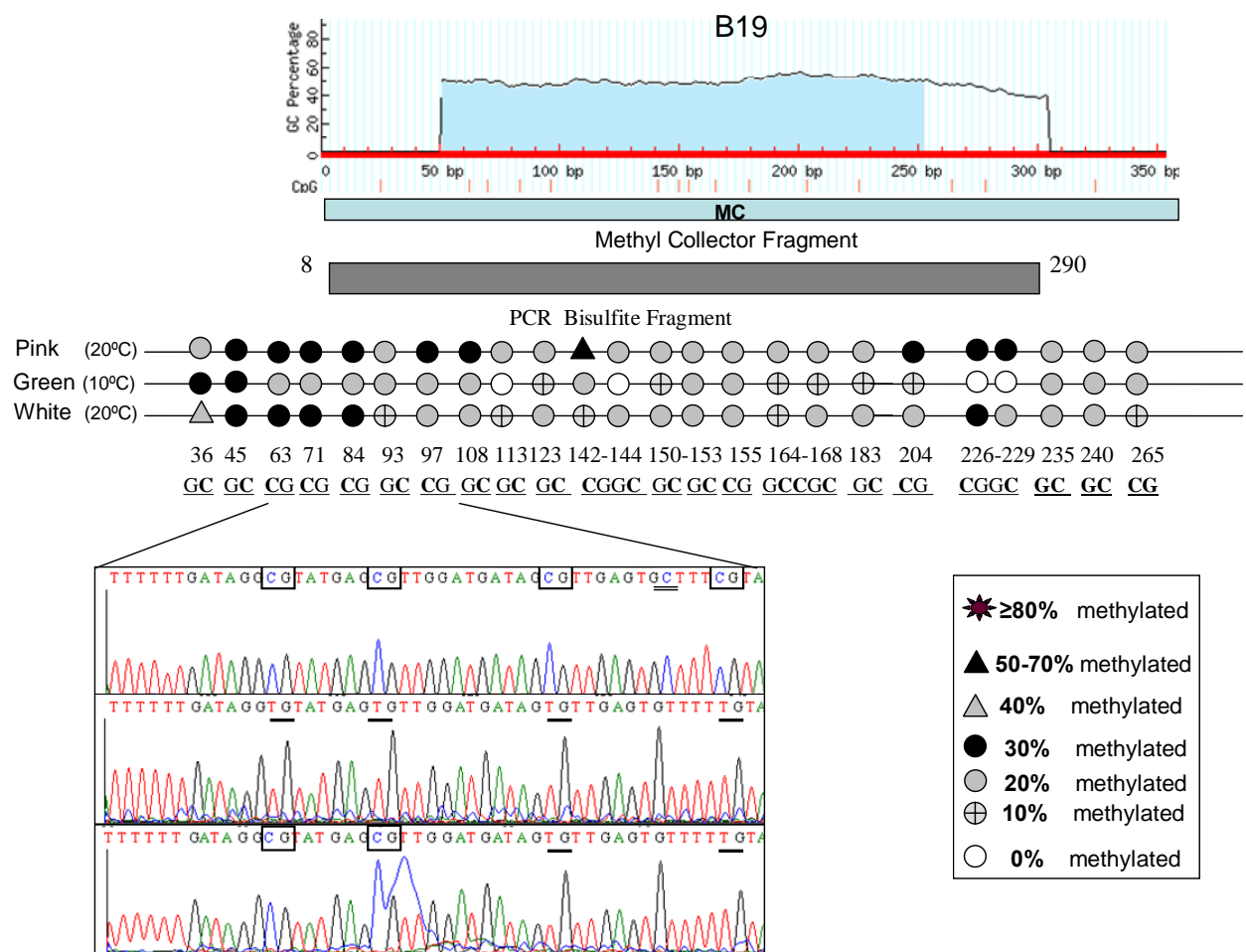


**Fig. S4 Bisulfite sequencing for B19 fragment**



**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.