

Figure S6

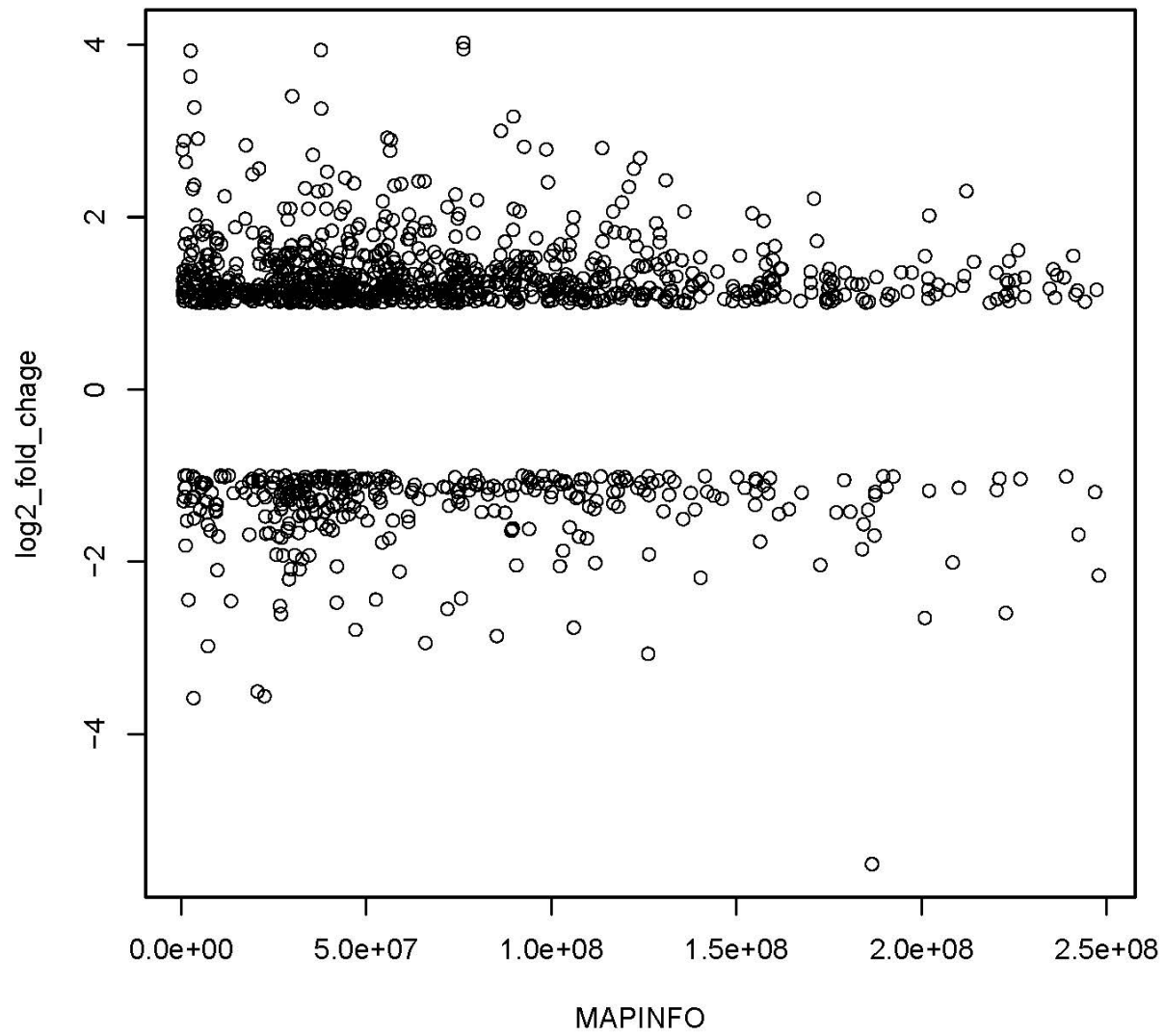


Figure S6: Differential methylated pattern analyses analyzed from the Infinium microarray analyses. Scatter plot of $\log_2(\text{fold_change})$ of the 1,202 sites illustrating differences in DNA methylation between control and PJ34-treated (PARylation inhibited) cells. A total of 1,202 CpG Sites with $\text{abs}(\log_2(\text{fold_change})) > 1$ were observed.