



Figure 4-Supplemental 1. Heatmap of DNA methylation of MYH1, PCDHB16, PCDHB15 and BCL2L10 in the first set of 10 metastatic melanoma patient sample and 20 primary patient melanoma.

The median DNA methylation of the analysed CpG for each gene is indicated as percentage and overall survival in months. Primary samples are divided by a small black line indicating the cut-off at one year between the short and long overall survival (SS and LS).