Supplementary Information

MicroRNA signature refine response prediction in CML

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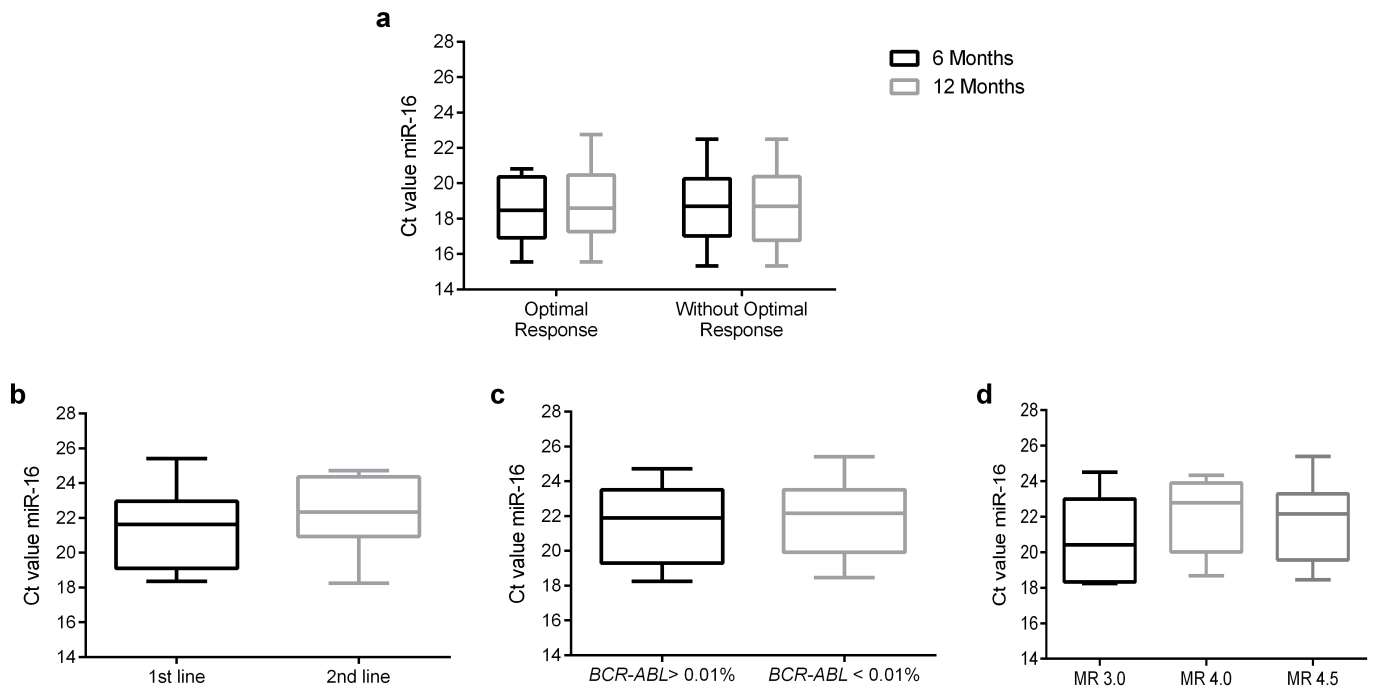
Supplementary Figures

Figure S1 – Ct variance of miR-16 in CML patients according the different comparative groups. The Ct value of miR-16 of CML patients at diagnosis according response levels (a). At follow-up measures, the Ct variance of this miR according to the number of TKI that patients were exposed (b), to *BCR-ABL1* values (c) and to molecular response rates (d). The results are represented as median with interquartile range. Any statistical difference was detected between groups.