

Supplementary figure set (exclusion peak correction)

Exclusion peak correction for GIM screens. For each microarray hybridization file, the horizontal axis represents the relative genomic position of the affected gene, while the vertical axis represents, in \log_2 transformed values, the measured growth defect. The title indicates the identifier of the screen replicate. For example « YDR283C GCN2 DEL 2 » corresponds to the screen done with the deletion (« DEL ») of the GCN2 gene, systematic name « YDR283C », for replicate number 2.

In red, original values, in black, values after correction using an estimation of recombination frequencies.

















































































