Figure S6 – Nucleosome positions (left panel) and Abf1 chromatin occupancy (right panel) in relation to TSS's is related to the variation in the width of NFR. Nucleosome data and Abf1 ChIP-seq data from Figure 6A are plotted at GCBS-associated TSS's as a heatmap. The intensity is a measure of occupancy in reads indicated to the right of each heatmap. Plots were generated using SeqPlots and transferred into the R-statistical environment. Here the nucleosome traces were sorted by NFR-size from widest (top) to narrowest (bottom).