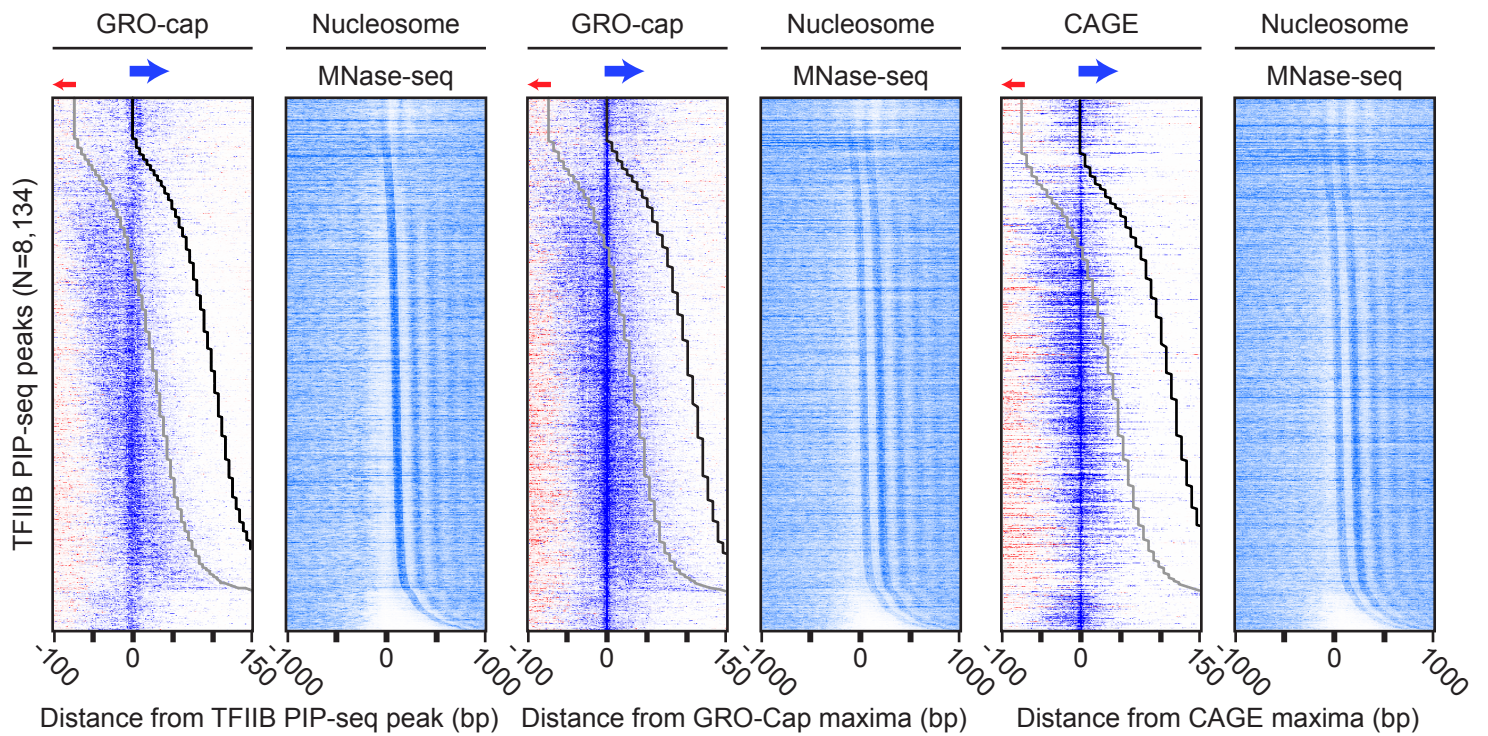


Supplemental Fig 7



Supplemental Figure 7. Dispersion validation. Heatmaps of GRO-cap and CAGE with three distinct reference points: TFIIB PIP-seq peak, GRO-Cap local maxima, and CAGE local maxima. All heatmaps were independently sorted with respect to closest downstream +1 nucleosome determined as previously described.