



Figure 1–figure supplement 4. Abf1 and Reb1 motifs based on CUT&RUN and ORGANIC ChIP-seq are similar.

The MEME motif-finding program was applied to (A-B) 1''-32'' pooled CUT&RUN ≤ 120 bp data and (C) 600 mM Abf1 and (D) 80 mM Reb1 ORGANIC data, and log-odds sequence logos are shown. Note the close correspondence between motifs determined using CUT&RUN and ORGANIC. (E) Percentage of peak calls with motifs. For each ≤ 120 bp dataset, peaks were called using thresholds set to recover similar numbers of peaks (stringent ~ 650 and relaxed ~ 1100).