Figure S1. Chemical cleavage mapping for *cen1* and *cen3*. See legend to Figure 1A.

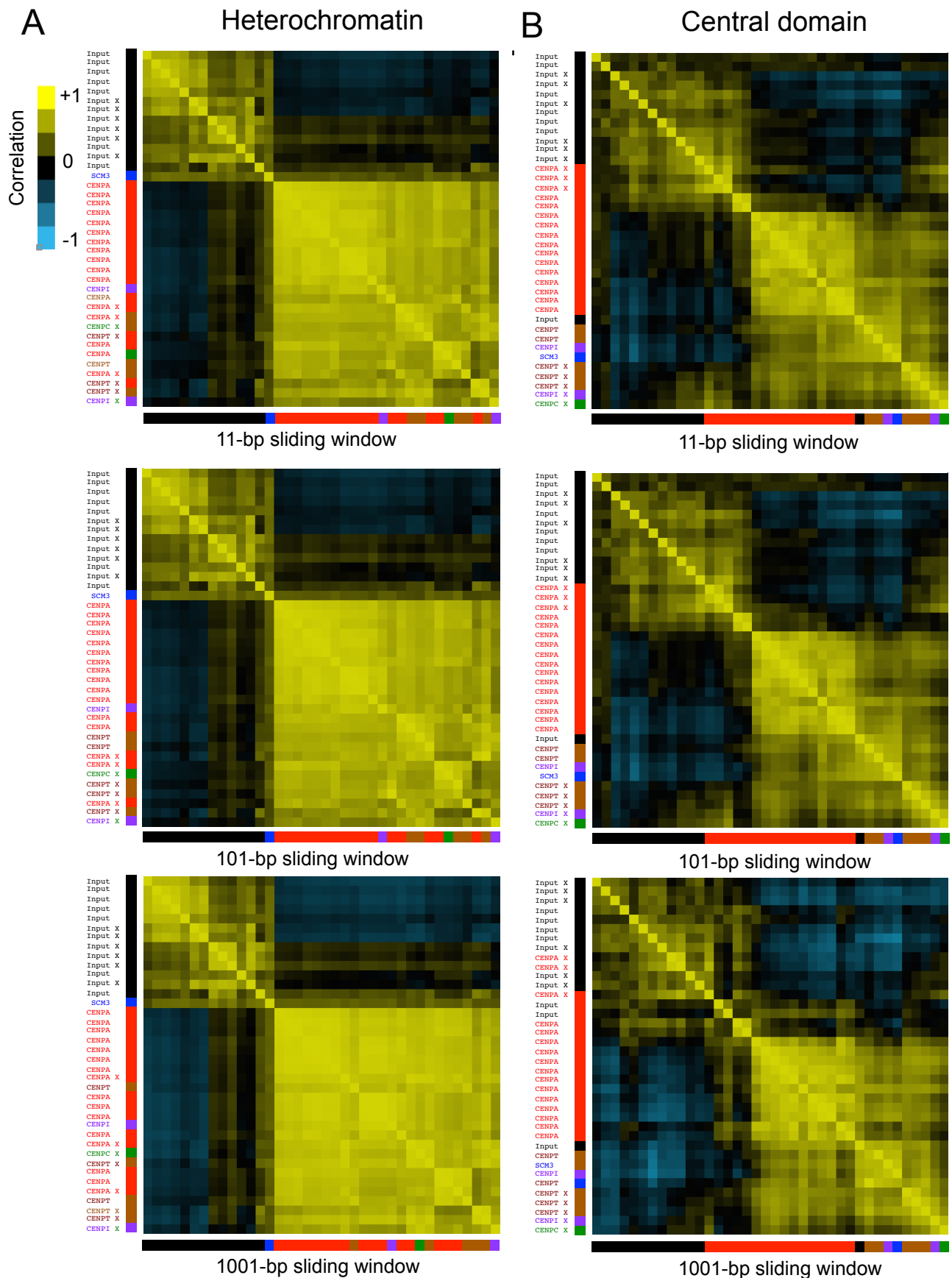J. Thakur, P. B. Talbert, and S. Henikoff

Figure S2. Correlation matrix heat maps with smoothing. See the legend to Figure 5. Triangular smoothing was applied using sliding windows as indicated.
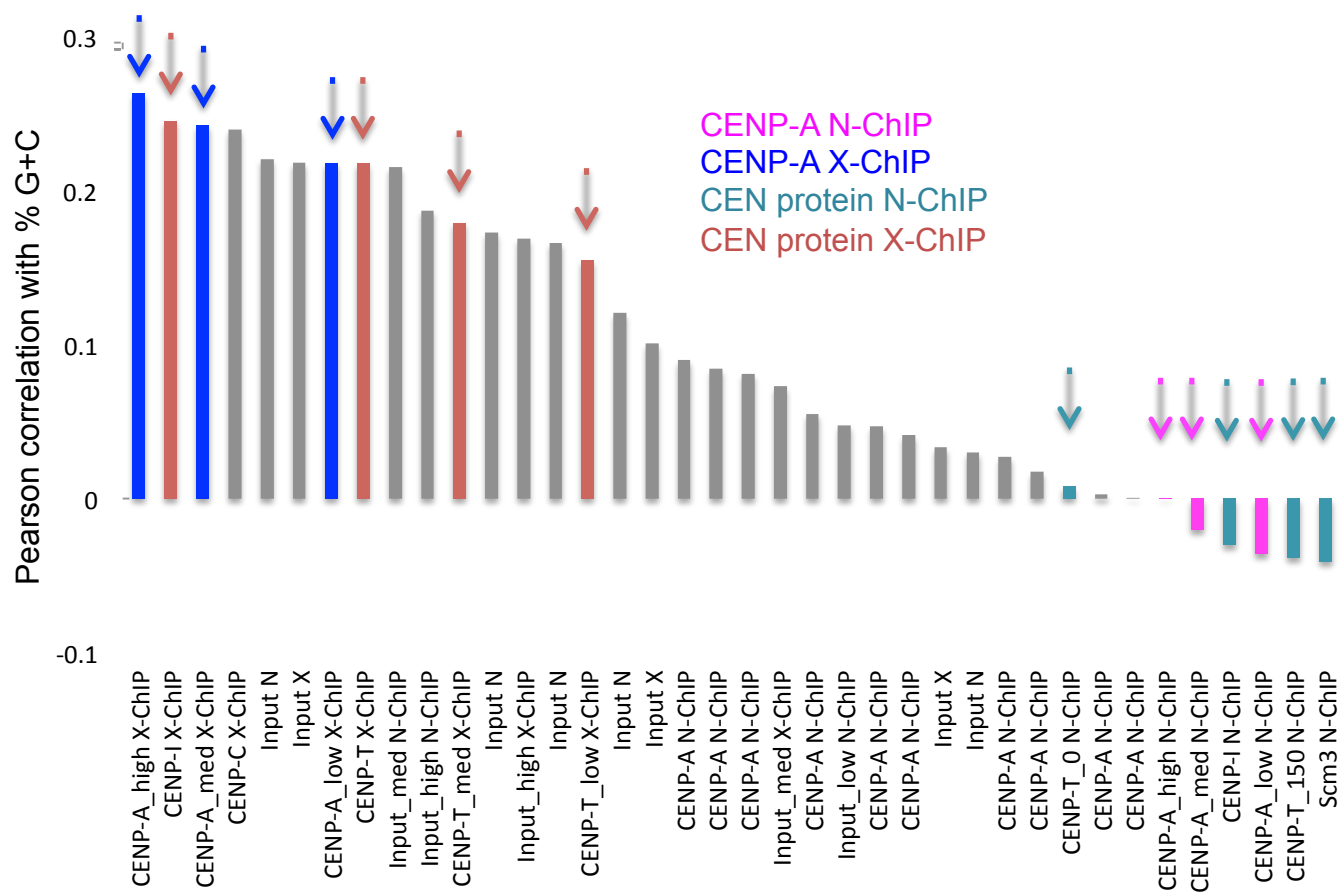
J. Thakur, P. B. Talbert, and S. Henikoff

Figure S3. Correlation of GC content with signal in 38 ChIP datasets. Arrows correspond to datasets displayed in Figure 4.

J. Thakur, P. B. Talbert, and S. Henikoff

Table S1 *S pombe* **strains used in the study.**

Strain	Genotype	Source
FY20724	*h minus cnp1-FLAG::kanR*	NBRP, Japan PMID: 21445296
PE199	*h minus leu1 ade6 cnp20-FLAG::kanR*	Takeshi Sakuno PMID: 19758558
FY8393	*h minus leu1 mis6-HA[LEU2]*	NBRP, Japan PMID: 9230309
FY21240	*h minus scm3-FLAG[hph]*	NBRP, Japan PMID: 19217403
FY20571	*h minus lys::cnp1-HA Lys+*	NBRP, Japan PMID: 10864871
FY8426	*h minus leu1 ura4 mis6-GFP[LEU2]*	NBRP, Japan PMID: 9230309
hy067	*h minus leu1-32 ura4DS/E ade6-210 cnt1::ura4+ pREP41-myc-cnp3*	Songtao Jia PMID: 19910462
H4S47C	*h minus his3-D1 leu1-32 ura4-D18 ade6-M210, hhf1S47C, hhf2S47C*	Ji-Ping Wang PMID: 24277842

J. Thakur, P. B. Talbert, and S. Henikoff