

Supplemental Information

SAF-A Regulates Interphase Chromosome

Structure through Oligomerization

with Chromatin-Associated RNAs

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Table S1. List of fosmids, Related to Figures 1, S1, 6, S6.

Position	Clone names	Sequence name	Co-ordinates (hg38)	Properties
11p15.5	CI 11p15-49		chr11:736369	gene rich
11p15.5	H19-IGF2		chr11:2070000	gene rich
11p15.1	WI2-2224K21	G248P87004F11	chr11:17,488,939-17,528,052	gene rich
11p15.1	WI2-3168N23	G248P8017G12	chr11:19,034,283-19,072,361	gene rich
11p14.1	WI2-906E12	G248P8190C6	chr11:29,684,660-29,731,106	gene poor
11p14.1	WI2-1673H3	G248P83109D2	chr11:31,195,720-31,235,826	gene poor
Xq13.1	WI2-2977E1	G248P89772C1	chrX:69862989-69904969	gene rich
Xq13.1	WI2-1660A18	G248P87866A9	chrX:71,851,120-71,887,633	gene rich
Xq24	WI2-2726C8	G248P88902B4	chrX:121,055,336-121,094,297	gene poor
Xq25	WI2-465K3	G248P8676F2	chrX:123,014,421-123,054,105	gene poor
1p31.2	WI2-2013D5	G248P86197B3	chr1:68350693-68390943	gene poor
1p31.2	WI2-2446E1	G248P83504C1	chr1:69,563,715-69,605,049	gene poor
2p25.1	WI2-1914D7	G248P86693B4	chr2:11,338,760-11,377,104	gene rich
2p25.1	WI2-2004M11	G248P85784G6	chr2:11,800,555-11,838,935	gene rich
21q22.3	WI2-2110F20	G248P86155C10	chr21:42,316,026-42,353,544	gene rich
21q22.3	WI2-2736C8	G248P88986B4	chr21:42,398,031-42,434,239	gene rich

Fosmids were obtained from BacPac resources, DNA co-ordinates are hg38 and gene density was calculated from Ensembl 87.