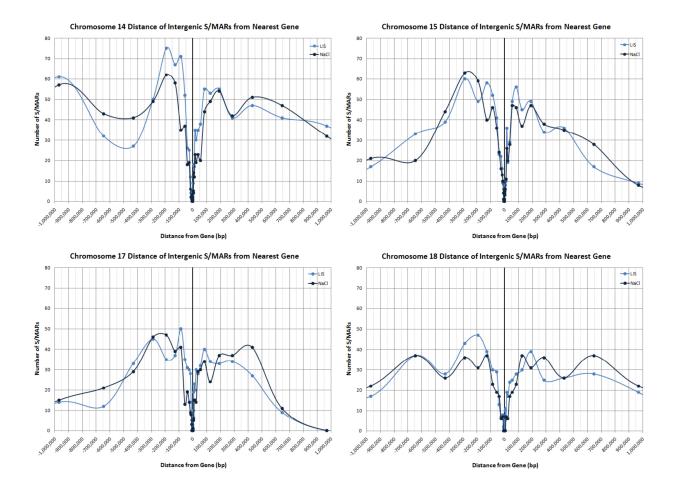


Supplemental Figure 1. Spacing of S/MARs along human chromosomes 14, 15, 17 and 18. The binned distance between S/MARs is shown for nuclear scaffolds/matrices isolated by LIS (light blue) or NaCl (dark blue) extraction with the peak distance frequency for each indicated and averages of each given.



Supplemental Figure 2. Intergenic nuclear scaffold/matrix attachment and expression. SARs (light blue) and MARs (dark blue) were mapped according to their distance from both the 5' and 3' ends of genes on chromosomes 14, 15, 17 and 18.