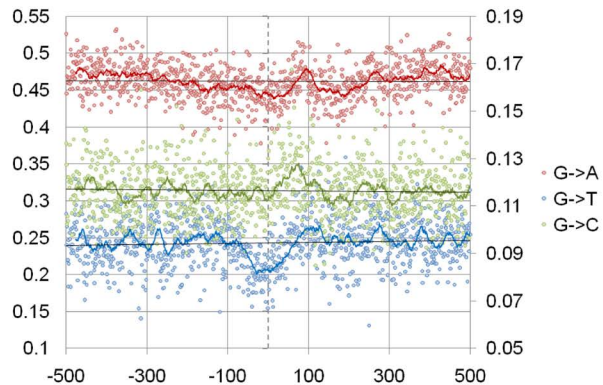
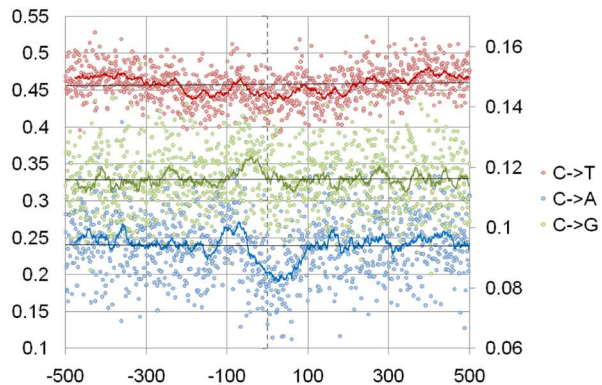
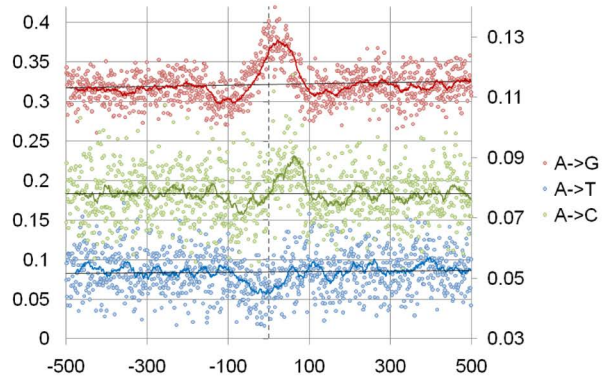
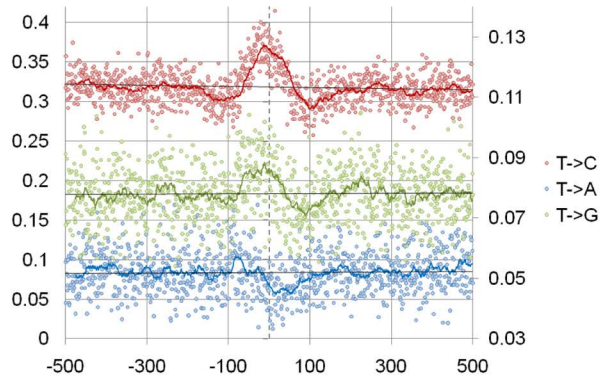
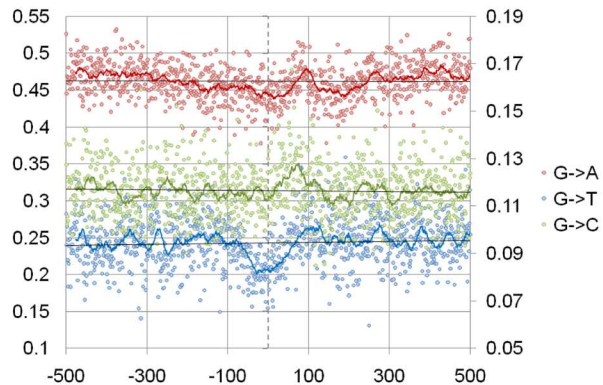
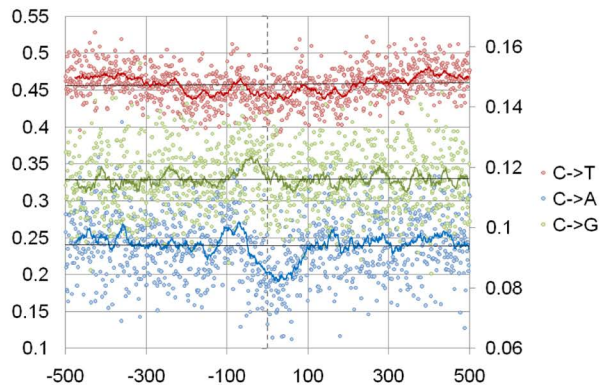
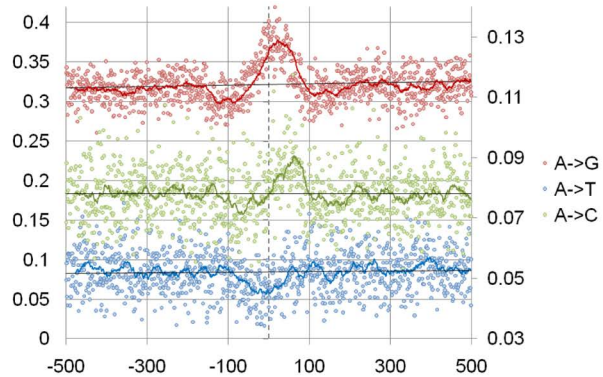
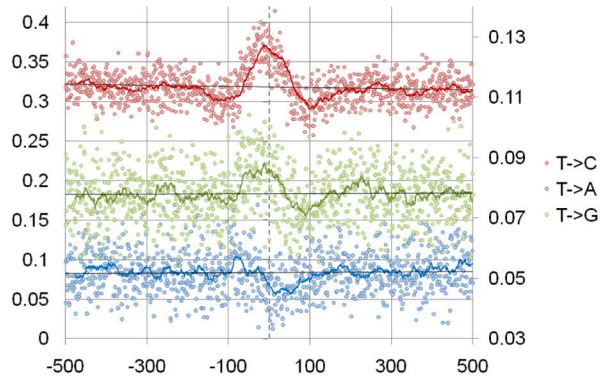


Human-chimpanzee divergence (% of ancestral bases)



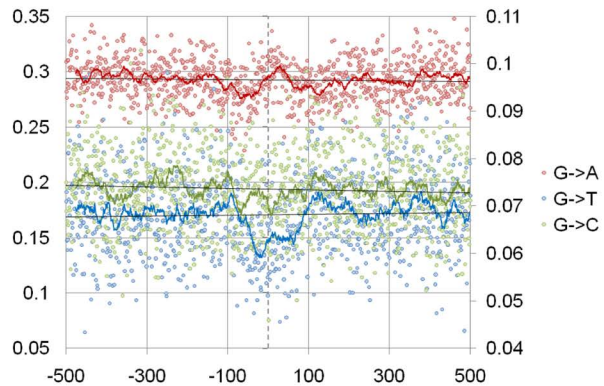
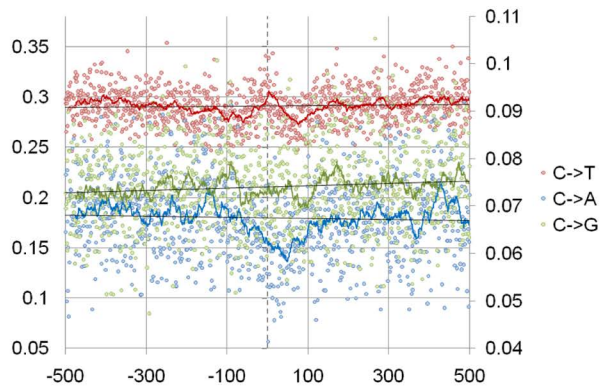
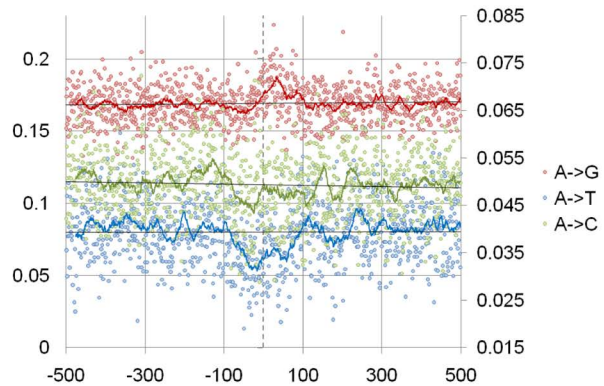
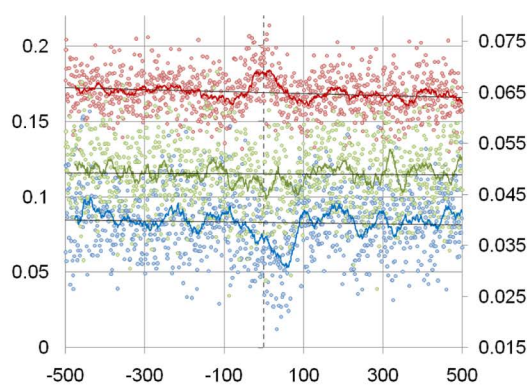
Position from nucleosome dyad (bp)

Human-chimpanzee divergence (% of ancestral bases)

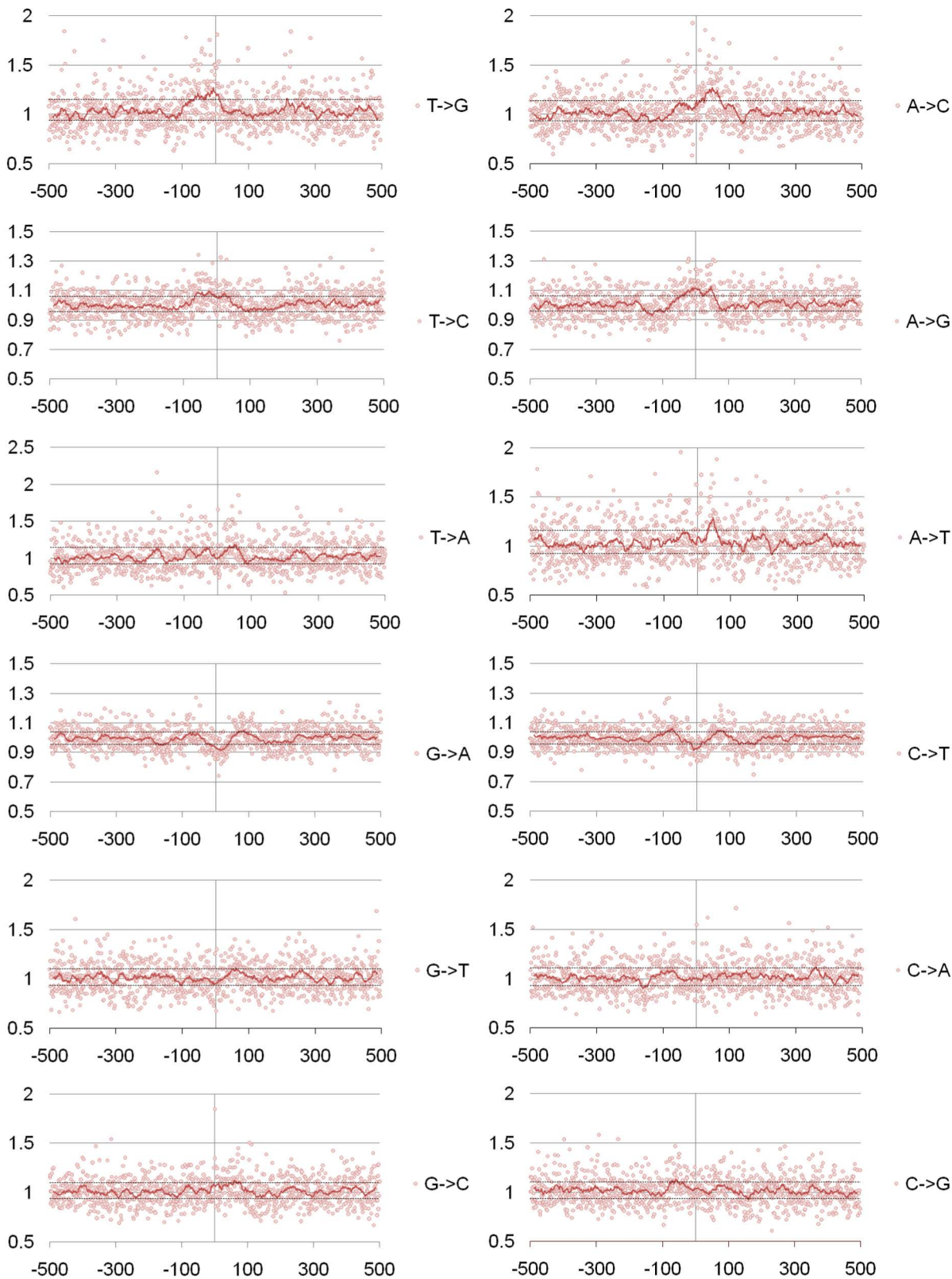


Position from nucleosome dyad (bp)

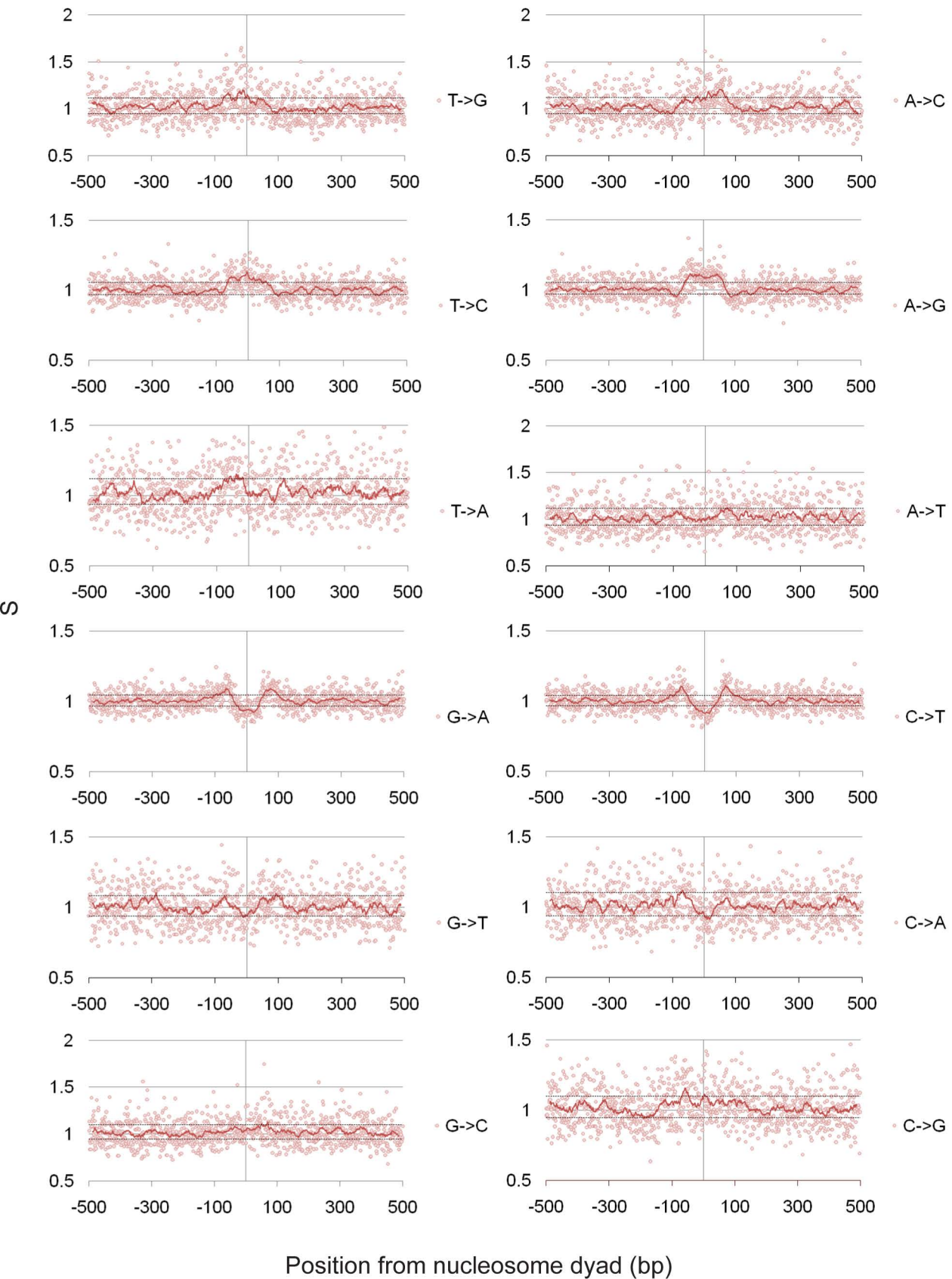
Rate of intraspecies divergence (% of ancestral bases)



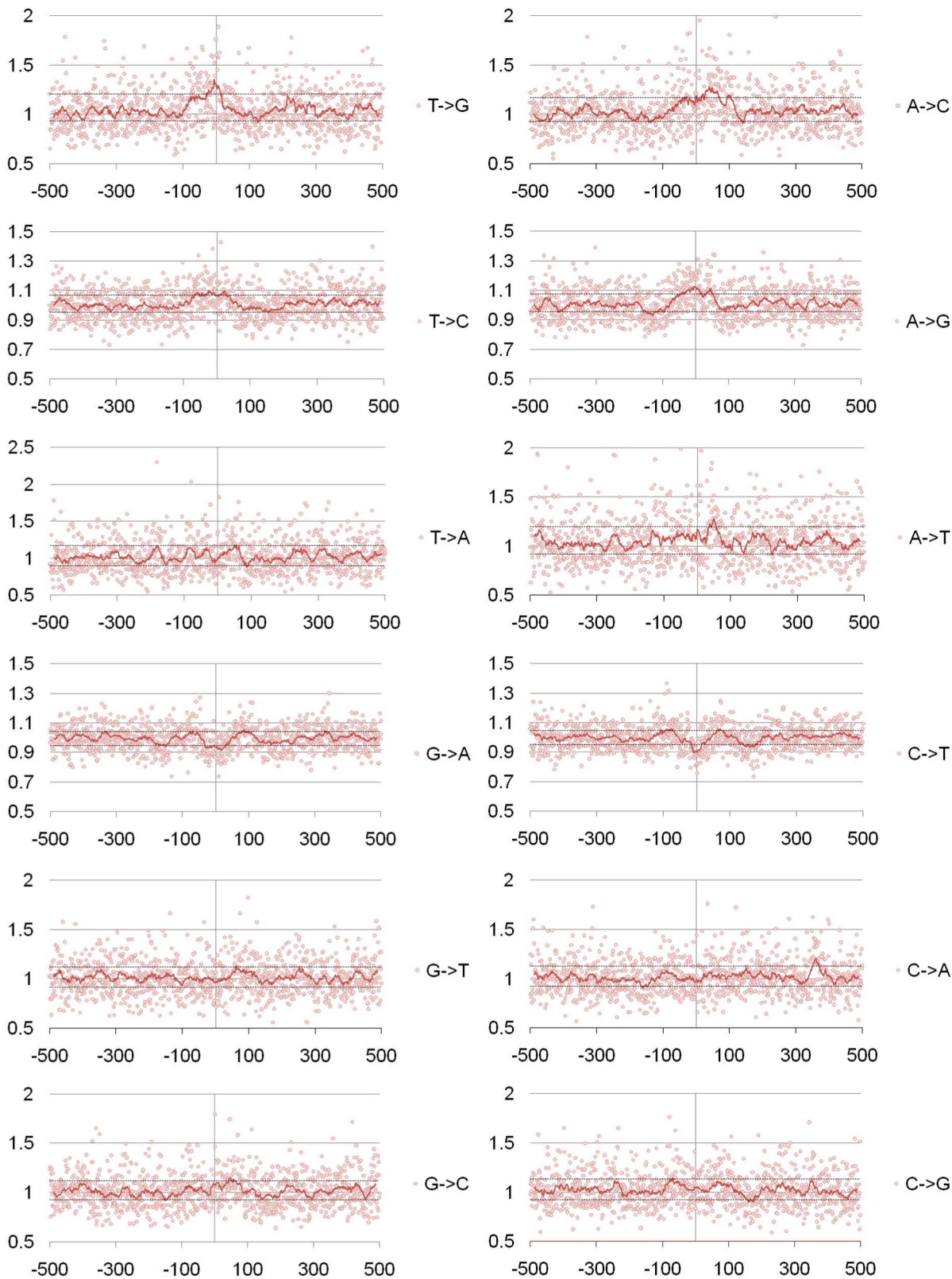
Position from nucleosome dyad (bp)

ω 

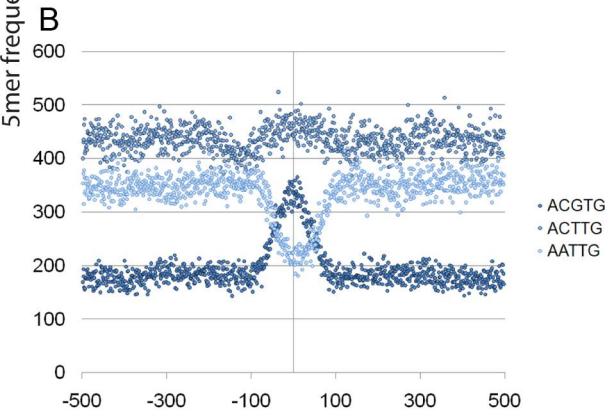
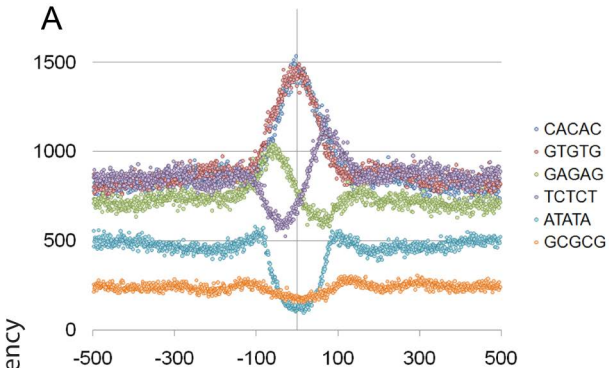
Position from nucleosome dyad



ω

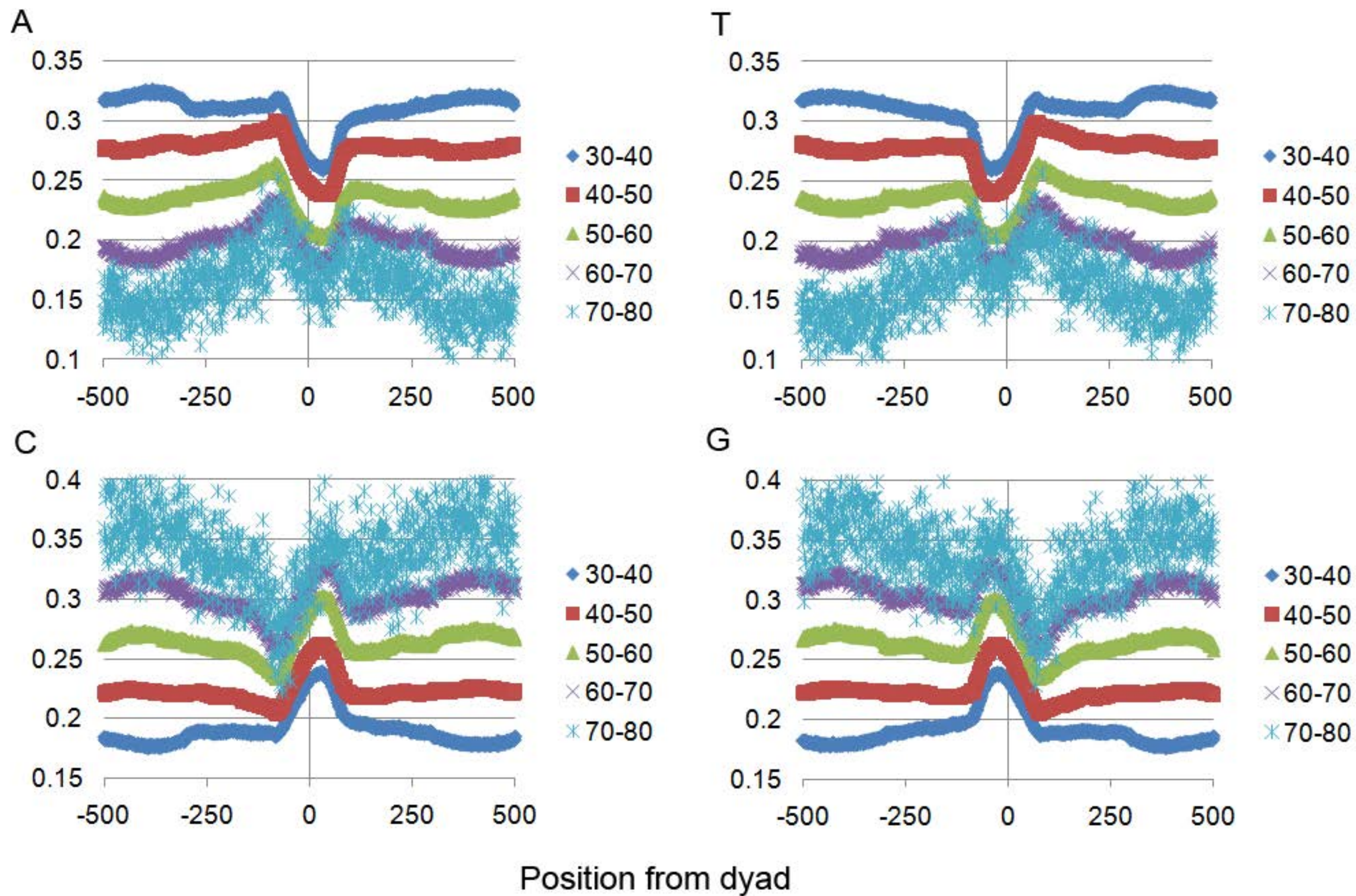


Position from nucleosome dyad

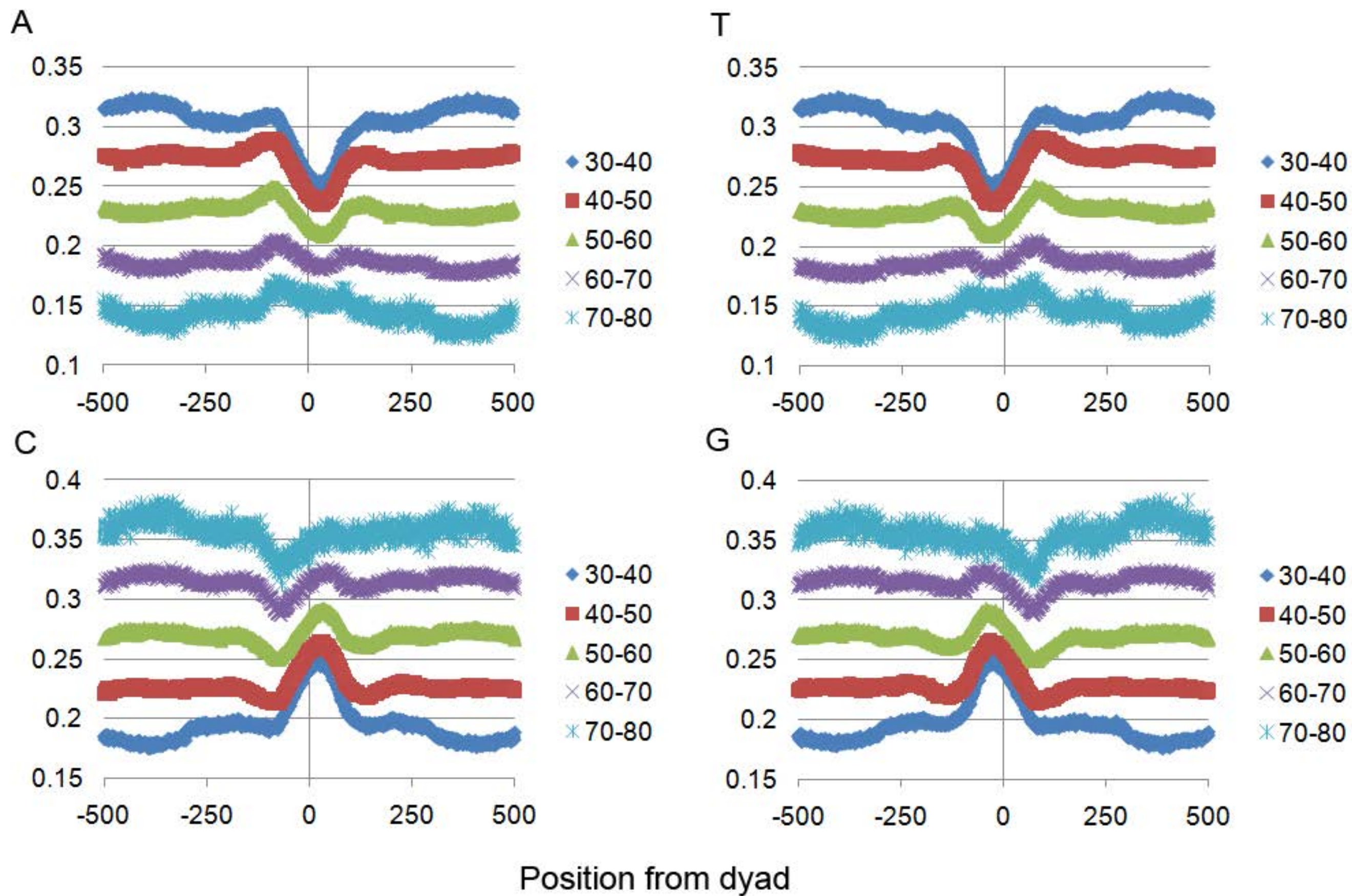


Position from nucleosome dyad

Mononucleotide frequencies



Mononucleotide frequencies



	-75 to +75 (Nucleosome core)			+/-75 to 105 (Linker region)			-105 to +105 (Core and linker)		
	GC < 45%	All	GC > 55%	GC < 45%	All	GC > 55%	GC < 45%	All	GC > 55%
C->T	-5.6 **	-1.6 *	3.0 *	0.4	2.2 *	3.3	-3.9 **	-0.5	3.2 *
G->A	-6.8 **	-2.7 **	2.4 *	1.2	2.5 *	5.6 *	-4.5 **	-1.3	3.1 *
C->A	1.0	-0.1	1.9	5.1	4.1 *	0.6	2.2	1.1	1.4
G->T	-1.0	0.1	0.2	0.3	0.7	-0.2	-0.3	0.3	-0.1
G->C	4.4	4.4 *	7.2 *	4.5	3.8 *	3.9	4.4 *	4.1 **	6.2 *
A->T	7.6 *	6.8 *	1.8	12.1 *	1.6	-1.7	9.1 *	5.2 *	0.7
C->G	3.9	3.1 *	3.3	3.4	2.4	0.0	3.8	3.0 *	2.6
T->A	6.2 *	5.8 *	3.8	0.1	-0.9	-5.8	4.0	3.6 *	1.3
T->C	10.4 **	4.8 **	-0.2	0.7	-1.6	-3.6	7.9 **	3.1 **	-0.9
A->G	11.9 **	6.5 **	-3.0	0.0	-3.2 *	-8.1 *	8.7 **	3.8 **	-4.4 *
T->G	13.3 **	10.0 **	3.2	2.1	1.8	-0.7	10.1 **	7.7 **	2.3
A->C	17.4 **	10.6 **	1.4	1.9	0.4	-4.0	12.8 **	7.6 **	-0.2