

Table 1: Number (proportion) of the negative subset of the 500 most highly *COMT* – correlated genes in each brain area having three different consensus initiator element (Inr) sequences on either DNA strand

	Inr1 ^a	Inr2 ^b	Inr3 ^c	Number of Strands
Prefrontal Cortex	5 (.015)	3 (.009)	304 (.905)	336
Cerebellum	5 (.022)	5 (.022)	186 (.802)	232
Temporal Cortex	5 (.027)	2 (.011)	160 (.851)	188
Pons	4 (.029)	1 (.007)	120 (.882)	136

TSS +1 locus underlined

a. Inr: BBCABW

b. Inr: YYANWYY

c. Inr: YR