

Table 1: Number of positively COMT-correlated genes in the full sample (gene as unit of analysis) (Column 1), in the 500 most strongly correlated (ranked by p-values) (Column 3), and in the genes among the top 500 only found in the corresponding brain area but not among the top 500 genes in any of the other three areas (Column 5).

	<i>Pos. r full sample</i>	<i>Full Sample</i>	<i>Pos. r top 500 genes</i>	<i>Top 500 genes</i>	<i>Pos. r unique</i>	<i>Unique genes</i>
Prefrontal	6329	16670	330	501	75	199
Cerebellum	6158	16685	382	501	144	236
Temporal	5761	16636	407	501	149	209
Pons	6600	16616	433	501	190	235