Table 1: Distribution of the top 200 negatively COMT-correlated genes by Chromosome: Temporal

Chromosome	Frequency	Expected	Enrichment	P value
1	20	20.9	0.96	0.84000
2	10	12.8	0.78	0.42000
3	17	10.8	1.58	0.05000
4	10	7.6	1.32	0.37000
5	21	8.9	2.35	0.00000
6	13	10.7	1.21	0.47000
7	4	8.9	0.45	0.09000
8	7	6.8	1.04	0.94000
9	8	8.1	0.99	0.97000
10	13	7.5	1.73	0.04000
11	6	13.2	0.45	0.04000
12	13	10.8	1.20	0.49000
13	4	3.3	1.22	0.70000
14	7	6.2	1.13	0.74000
15	7	5.8	1.21	0.61000
16	4	8.3	0.48	0.13000
17	7	11.7	0.60	0.16000
18	6	2.8	2.11	0.06000
19	4	13.8	0.29	0.01000
20	6	5.8	1.04	0.93000
21	4	2.7	1.51	0.42000
22	1	4.4	0.23	0.10000
X	8	8.3	0.97	0.92000