Effects of MDMA on GFAP immunoreactivity

Four-way ANOVA of GFAP immunoreactivity in the CTX revealed a significant effect of treatment ($F_{1,48} = 6.64$, p < 0.05) and a significant gender × treatment interaction ($F_{1,48} = 4.26$, p < 0.05). Moreover, in the CPu a significant effect of gender ($F_{1,95} = 12.75$, p < 0.001), genotype ($F_{1,95} = 6.55$, p < 0.05) and treatment ($F_{1,95} = 358.44$, p < 0.0001) was observed. Furthermore, in the NAc a significant effect of gender ($F_{1,48} = 8.97$, p < 0.005) and a significant gender × genotype × age interaction ($F_{1,48} = 5.18$, p < 0.05) was observed. Finally, in the SNc, a significance of gender ($F_{1,97} = 17.22$, p < 0.001), genotype ($F_{1,97} = 11.60$, p < 0.001), age ($F_{1,97} = 30.34$, p < 0.0001), treatment ($F_{1,97} = 22.65$, p < 0.001) and a significant gender × genotype ($F_{1,97} = 8.50$, p < 0.005) and gender × age ($F_{1,97} = 96.79$, p < 0.0001) interactions were also observed, whereas in VTA a significant effect of gender ($F_{1,48} = 42.81$, p < 0.0005), a significant effect of age ($F_{1,48} = 5.86$, p < 0.05) and a significant gender × age interaction ($F_{1,48} = 8.59$, p < 0.05) was observed.

Effects of MDMA on CD11b immunoreactivity

Four-way ANOVA of CD11b immunoreactivity in the CTX revealed a significant effect of gender $(F_{1,48} = 13.60, p < 0.005)$, genotype $(F_{1,48} = 17.47, p < 0.0005)$, age $(F_{1,48} = 26.66, p < 0.0005)$, treatment $(F_{1,48} = 169.81, p < 0.0005)$, and significant gender × genotype $(F_{1,48} = 8.70, p < 0.005)$, gender × age $(F_{1,48} = 111.78, p < 0.0005)$, genotype × age $(F_{1,48} = 17.47, p < 0.005)$ interactions. Moreover, in the CPu a significant effect of gender $(F_{1,90} = 33.24, p < 0.0001)$, genotype $(F_{1,90} = 24.78, p < 0.001)$, treatment $(F_{1,90} = 50.68, p < 0.0001)$, and a significant gender × genotype $(F_{1,90} = 7.195, p < 0.01)$, age × treatment $(F_{1,90} = 5.51, p < 0.05)$ and gender × genotype × age $(F_{1,90} = 18.10, p < 0.005)$, age $(F_{1,48} = 60.66, p < 0.0005)$, treatment $(F_{1,48} = 47.49, p < 0.0005)$, and a significant gender × age interaction $(F_{1,48} = 11.23, p < 0.005)$, treatment $(F_{1,94} = 30.55, p < 0.0001)$, genotype $(F_{1,94} = 9.98, p < 0.005)$, treatment $(F_{1,94} = 80.58, p < 0.0001)$, treatment $(F_{1,94} = 80.58, p < 0.001)$, treatment $(F_{1,94} = 80.58, p < 0.001)$, the SNC a significant effect of gender $(F_{1,94} = 80.58, p < 0.001)$, the SNC a significant effect of gender $(F_{1,94} = 80.58, p < 0.001)$, the SNC a significant effect of gender $(F_{1,94} = 80.58, p < 0.001)$, the SNC a significant effect of gender $(F_{1,94} = 80.58, p < 0.001)$, the SNC a significant effect of gender $(F_{1,94} = 80.58, p < 0.001)$, the SNC a significant effect of gender $(F_{1,94} = 80.58, p < 0.001)$, the SNC a significant effect of gender $(F_{1,94} = 80.58, p < 0.001)$, the SNC a significant effect of gender $(F_{1,94} = 80.58, p < 0.001)$, the solution $(F_{1,94} = 80.58, p < 0.001)$, the solution $(F_{1,94} = 80.58, p < 0.005)$, the solution $(F_{1,94} = 80.58, p < 0.005)$.

p<0.0001), and gender × treatment ($F_{1,94} = 13.36$, p<0.001) and gender × genotype × age ($F_{1,94} = 5.58$, p<0.05) interactions were observed, whereas in VTA a significant effect of genotype ($F_{1,48} = 40.95$, p<0.0005), age ($F_{1,48} = 12.94$, p<0.005) and treatment ($F_{1,48} = 116.42$, p<0.0005) were observed.

Effects of MDMA on TH immunoreactivity

Four-way ANOVA of TH immunoreactivity in the CTX revealed a significant effect of gender ($F_{1,48} = 73.61$, p < 0.0005) and age ($F_{1,48} = 16.27$, p < 0.0005). Moreover, in CPu a significant effect of gender ($F_{1,96} = 112.47$, p < 0.0001), genotype ($F_{1,96} = 57.80$, p < 0.0001), age ($F_{1,96} = 259.84$, p < 0.0001), treatment ($F_{1,96} = 69.36$, p < 0.0001), and a significant gender × genotype ($F_{1,96} = 24.35$, p < 0.001), gender × age ($F_{1,96} = 37.21$, p < 0.0001) and gender × treatment ($F_{1,96} = 19$, p < 0.001) interactions were also observed in CPu. Furthermore, in the NAc a significant effect of gender ($F_{1,48} = 17.34$, p < 0.0005) and age ($F_{1,48} = 292.17$, p < 0.0005) was observed. Finally, in SNc a significant effect of gender ($F_{1,108} = 67.67$, p < 0.0001), genotype ($F_{1,108} = 51.96$, p < 0.0001), age ($F_{1,108} = 71.94$, p < 0.0001) and treatment ($F_{1,108} = 77.70$, p < 0.0001), and a significant gender × age ($F_{1,108} = 144.47$, p < 0.0001), gender × treatment ($F_{1,108} = 77.70$, p < 0.0001), genotype × treatment ($F_{1,108} = 5.79$, p < 0.005), age × treatment ($F_{1,108} = 7.75$, p < 0.01) interactions were observed, whereas in VTA a significant effect of gender ($F_{1,60} = 10.65$, p < 0.005) and age ($F_{1,60} = 45.16$, p < 0.0005) and a significant gender × age interaction ($F_{1,60} = 56.98$, p < 0.0005) was observed.