

Supplemental materials

Transcriptomic and metabolomic analysis provide insights into the attenuation of neuroinflammation by nervonic acid in MPTP-stimulated PD model mice

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Table S1. The primer sequences of RT-PCR

β -actin	5'-GTCAGGTCACTACATCGGCAAT-3'(forward)	5'-AGAGGTCTTACGGATGTCAACGT-3'(reverse);
Mpp4	5'-CTCAGCCGTACCAACCTCAC-3'(forward)	5'-GCGGAACCCAGCGATAAA-3'(reverse);
Mpp3	5'-CAGAGTCCAACCCCAGTCCT-3'(forward)	5'-TCTTCTGGCAACCGTGTC-3'(reverse);
Mpp5	5'-TTGTTTCTGGCAAGCATT-3'(forward)	5'-CGGAGGGCTTCAAGGACTG-3'(reverse);
Mpp7	5'-CAAGGCAATCCGTGTAAAG-3'(forward)	5'-TCAGAGCCAGTCTCCGTTCC-3'(reverse);
Zdhhc23	5'-CAATGCCTGTCCCTGGTCTA-3'(forward)	5'-TTTCTGCCACATACGCTGT-3'(reverse);
Cd300a	5'-GAGCAGAATGAGTGCCAGTATGT-3'(forward)	5'-GTTGAATGCCACGGATGAATA-3'(reverse);
Rasd1	5'-ATGTGCCAAGCGACTCTG-3'(forward)	5'-CTCGATGGTAGGGGTGTAAGC-3'(reverse);
Rasd2	5'-ACTCAATGTGCCTGCTAACGAACTC-3'(forward)	5'-TCCTCGATAGTGGCGTGTA-3'(reverse);
Ptgs2	5'-TCTCAATGAGTACCGCAAACG-3(forward)	5'-TGGTCTCCCCAAAGATAGCA-3'(reverse);
Alox5ap	5'-ATACTTGTGGCTATCTGGG-3(forward)	5'-CGTCGTGCTTACCGTTCTG-3'(reverse);
Ryr1	5'-GTGCTGCTTACCAACCTGTC-3(forward)	5'-AAATAGATCCGTTCTATGCGAC-3'(reverse);
Slc18a2	5'-CCTCATGGACCACAACTGC-3(forward)	5'-CAATGGATGGCGTGACTAAG-3'(reverse);
Penk	5'-GGGGCTTATGCGGTTCT-3(forward)	5'-CACAGCTTCAGGCAGTGTAGT-3'(reverse);
Adcy5	5'-CTTCAGGGAGCCGACTTA-3(forward)	5'-GCAGGAAACACGACAGGTAGA-3'(reverse);
Pde1c	5'-GAAGCCCAGGTTCAAGAGC-3(forward)	5'-CGAAGGACCACGTATCCAC-3'(reverse);
Gbp2	5'-AAGGGTGACAACCAGAACATGAC-3(forward)	5'-CACAAAGTTAGCGGAATCGT-3'(reverse);
Ddc	5'-CACATTGACGCTGCTTACG-3(forward)	5'-TCAGTCCTCCTCTTCACCCA-3'(reverse);
Th	5'-GGTCTACTGTCTGCCGTGAT-3(forward)	5'-TGTGCGGTCAGCCAACAT-3'(reverse);
Ak7	5'-CCTAAGTATCCTCCGTGACTGG-3(forward)	5'-CGTAATCGAAGCGTTGTTGA-3'(reverse);
Pla2g4a	5'-TCCCTTGATTCTCGACCTC-3(forward)	5'-TCATTTGCCACTTCTC-3'(reverse).

Fig. S1 The mice from head up to full turn down was recorded as turn time in pole climbing tests

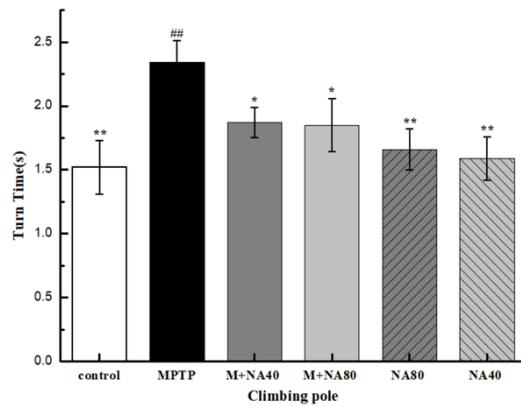


Fig. S2 The Y-Maze analysis the total number of entering arms of mice

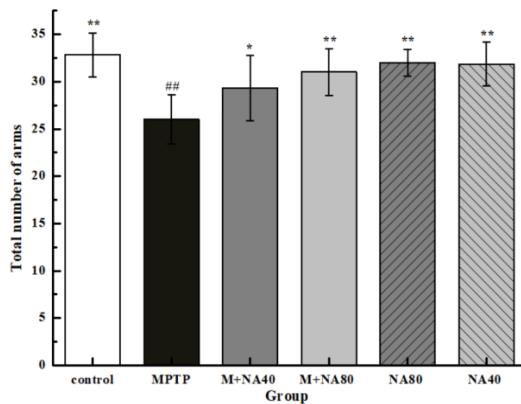


Fig. S3 Average escape distance of reaching the hidden platform during training (5 days); Average escape latency of reaching the hidden platform during training (5 days).

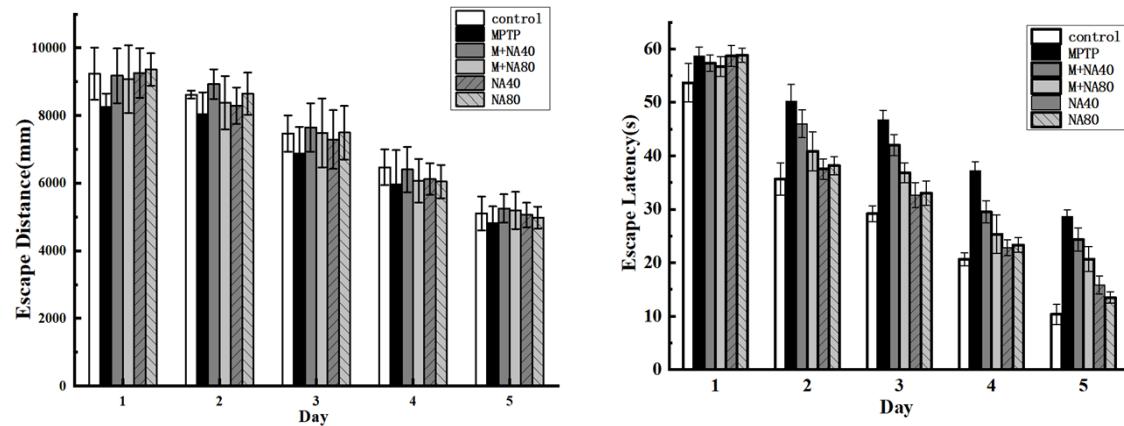


Fig. S4 On the fifth day, the average escape latency of mice reaching the hidden platform.

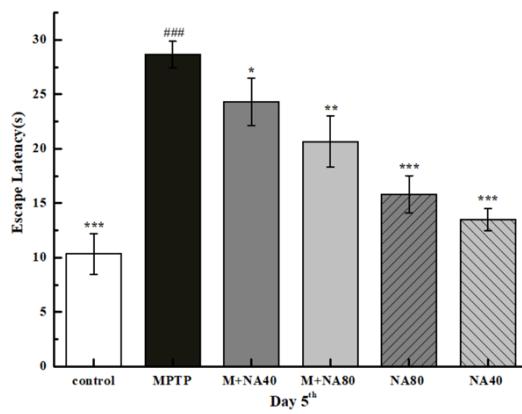


Fig. S5 On the 6th day, the time spent in the target quadrant where the hidden platform;

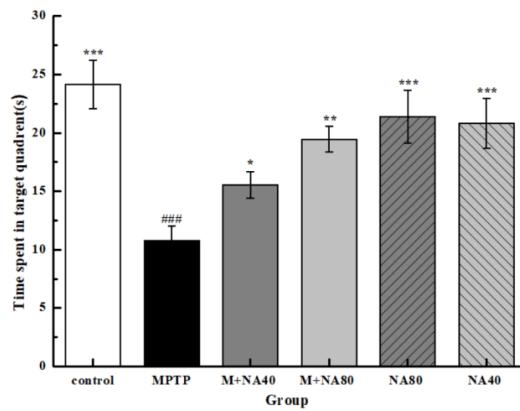


Fig. S6 Correlation coefficient (R^2) between samples, PCA diagram and statistical map of differential genes. (A) Heat map of correlation coefficient between samples and two dimensional principal component analysis (PCA) diagram. (B) Volcano map and statistical map of the number of up-regulated and down-regulated genes.

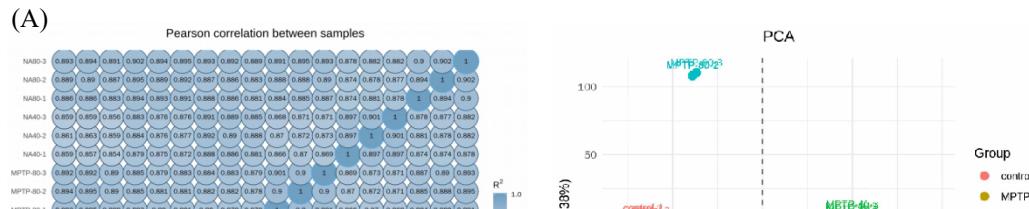


Fig. S7 Verification of Gene Expression Levels. All the experiments were repeated at least three times and similar results were observed. Values are mean \pm SEM, ($n = 4$ samples/group). p# < 0.05, p## < 0.01 vs control group, p* < 0.05, p** < 0.01 vs MPTP-induced group.

