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Supplemental Material

Evaluation of Neurotoxicity in BALB/c Mice following Chronic Exposure to Polystyrene Microplastics

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Figure S2. BBB integrity was detected by biotin tracer experiments. The sections were stained with biotin (red) and DAPI (blue). The existence of biotin in brain tissues was examined by immunofluorescence microscopy. The biotin signal of the liver parenchyma was used as a positive control (scale bar = $20 \mu m$).

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Figure S7. The total distance moved (A) and average movement speed (B) of mice in testing day (day 3) of the novel object recognition test are shown. The results are expressed as means \pm SD (n = 10 mice/per group). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05 vs. control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.

Figure S8. The numbers of neurons in the hippocampal cornu ammonis 1 (CA1), cornu ammonis 3 (CA3), and dentate gyrus (DG) sections were counted (n = 3 mice/group, n = 3 slides/mice). Data are shown as mean \pm SD. The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05 compared with control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.

Figure S9. The mRNA levels of *caspase 3* (A), the *Bax/Bcl-2* mRNA ratio (B) in the hippocampus were tested with quantitative real-time PCR (qRT-PCR) by normalizing to *Gapdh*. The results are expressed as means \pm SD (n = 3, N = 3 mice/group). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05 and ***P* < 0.01 compared with the control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.

Figure S10. The expression of synapsin 1, synaptophysin, and PSD 95 protein in the hippocampus was measured via western blotting. The western blotting results were quantified and statistically analyzed, as shown in (Figure 7C) as mean \pm SD (n = 3, N = 3 mice/group). The mean and SD summary data for quantification of western blotting are shown in Table S3. *P*-Values for all comparisons are reported in Table S4.

Figure S11. The mRNA expression levels of *Gap43*(A), *Syt 4* (B), and *Ncam*(C) were detected by qRT-PCR in the hippocampus of mice exposed to PS-MPs and control mice. Data are shown as mean \pm SD (n = 3). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. ***P* < 0.01 compared with control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.

Figure S12. (A) A confocal fluorescence microscope was used to detect colocalization of Alexa Fluor 594-labled MCP-1 (red)/TNF- α (red) and DAPI (blue) (scale bar = 20 µm, n = 3) in the hippocampus of mice exposed to PS-MPs and control mice. (B) Percent of positivity was calculated based on the percentage of MCP-1 positive cells out of the total number of cells in an image. (C) Percent of positivity was calculated based on the percentage of TNF- α positive cells out of the total number of cells in an image. Data are shown as mean ± SD (n = 3). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05, ***P* < 0.01 compared with control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.

Accession number	Gene	Primer	Sequence $5' \rightarrow 3'$	Product size (base pairs)
NIM 001280726 1	Candh	Forward	AGGTCGGTGTGAACGGATTTG	179
NM_001289720.1	Gapan	Reverse	TGTAGACCATGTAGTTGAGTCA	1/8
ND4 001040120 1	Dduf	Forward	TCATACTTCGGTTGCATGAAGG	127
INM_001048139.1	Бапј	Reverse	AGACCTCTCGAACCTGCCC	157
NIM 001252241 1	C. 4 1	Forward	CTGTCACCACTGTTGCGAC	120
NM_001232341.1	Syl I	Reverse	GGCAATGGGATTTTATGCAGTTC	150
NIM 000209 2	Sut 1	Forward	TGCTTTTGGCCTCGTCTTCA	102
NM_009308.3	<i>Syl</i> 4	Reverse	GCGGTTTTACCCTTCACTTCAC	192
NNA 001001445 1	Mague	Forward	GACAGAACCCGAAAAGGGC	04
NM_001081445.1	ncam	Reverse	GTTGGGGACCGTCTTGACTT	94
NIM 000002 2	$C_{am}/2$	Forward	TGGTGTCAAGCCGGAAGATAA	114
INM_008085.2	Gap45	Reverse	GCTGGTGCATCACCCTTCT	114
NIM 001279601 1	Tnf-α	Forward	TCCCAGGTTCTCTTCAAGGGA	51
NWI_001278001.1		Reverse	GGTGAGGAGCACGTAGTCGG	51
NIM 011222.2	Mcp-1	Forward	TTAAAAACCTGGATCGGAACCAA	101
INIM_011555.5		Reverse	GCATTAGCTTCAGATTTACGGGT	121
NIM 021274 2	Cual10	Forward	CCAAGTGCTGCCGTCATTTTC	157
INIWI_021274.2	CACITO	Reverse	GGCTCGCAGGGATGATTTCAA	137
NM 031168 2	11.6	Forward	TAGTCCTTCCTACCCCAATTTCC	76
NM_031108.2	11-0	Reverse	TTGGTCCTTAGCCACTCCTTC	70
NIM 008261 /	11 1 R	Forward	GCAACTGTTCCTGAACTCAACT	80
NWI_000301.4	11-1 <i>p</i>	Reverse	ATCTTTTGGGGTCCGTCAACT	09
NM 000810 3	Camara 3	Forward	ATGGAGAACAACAAAACCTCAGT	74
NM_009810.3	Cuspuse 5	Reverse	TTGCTCCCATGTATGGTCTTTAC	/4
NIM 007527 2	Dav	Forward	TGAAGACAGGGGCCTTTTTG	140
INIMI_00/527.3	DUλ	Reverse	AATTCGCCGGAGACACTCG	140
NIM 000741 5	Rol 2	Forward	ATGCCTTTGTGGAACTATATGGC	120
11111_007/41.3	DCI-2	Reverse	GGTATGCACCCAGAGTGATGC	120

Table S1. Primer sequences used for qRT-PCR.

 Table S2. Detailed information of primary antibodies.

antibody	species	company	catalog	dilution
Anti-GAPDH	mouse mAb	Proteintech	60004-1-lg	1:1000 (WB)
Anti-PSD95	rabbit polyclonal Ab	Proteintech	20665-1-AP	1:1000 (WB)
Anti-synaptophysin	rabbit polyclonal Ab	Proteintech	17785-1-AP	1:1000 (WB)
Anti-synapsin 1	rabbit polyclonal Ab	Proteintech	20258-1-AP	1:1000 (WB)

Figure 1C: zeta potential	mean	SD		
(mV)				
0.5 μm	-31.10	0.64		
4 μm	-37.53	0.75		
10 µm	-29.97	0.50		
Figure 4B: Time for mice (e	exposed or not t	to PS-MPs) to f	ind the platfor	m during
learning trials (seconds)				
	mean	SD	mean	SD
	Da	y 1	Day2	
Control	78.53	19.19	52.01	30.15
0.5 μm 100 μg/L	71.82	28.44	52.08	30.15
0.5 μm 1000 μg/L	65.94	32.08	63.7	25.88
4 μm 100 μg/L	71.16	20.79	63.585	25.60
4 μm 1000 μg/L	82.21	13.72	69.28	21.20
10 μm 100 μg/L	75.25	27.71	64.13	28.17
10 µm 1000 µg/L	67.46	25.12	66.57	31.26
	Day 3		Da	ny 4
Control	55.31	31.10	39.63	29.75
0.5 μm 100 μg/L	50.02	31.28	50.37	24.42
0.5 μm 1000 μg/L	54.72	30.66	51.54	27.95
4 μm 100 μg/L	61.33	24.31	48.52	28.35

 Table S3. The mean and SD summary data for quantification.

4 μm 1000 μg/L	65.79	29.00	60.03	28.57
10 μm 100 μg/L	47.87	28.94	48.52	31.71
10 μm 1000 μg/L	70.06	26.77	67.43	29.61
	Da	y 5		
Control	25.40	13.85		
0.5 μm 100 μg/L	33.04	24.52		
0.5 μm 1000 μg/L	40.34	30.64		
4 μm 100 μg/L	37.59	24.74		
4 μm 1000 μg/L	46.81	21.80		
10 μm 100 μg/L	38.39	32.72		
10 μm 1000 μg/L	54.72	26.76		

Figure 4D: Number of times mice (exposed or not to PS-MPs) crossed the target

platform on sixth day

	mean	SD	
Control	3.75	1.85	
0.5 μm 100 μg/L	1.25	0.97	
0.5 μm 1000 μg/L	0.875	1.05	
4 μm 100 μg/L	1.00	1.00	
4 μm 1000 μg/L	0.625	0.70	
10 μm 100 μg/L	1.25	0.83	
10 μm 1000 μg/L	0.75	0.83	

Figure 4E: Percentage of time mice (exposed or not to PS-MPs) took in the quadrant

	mean	SD	
Control	44.37	12.80	
0.5 μm 100 μg/L	25.72	17.17	
0.5 μm 1000 μg/L	24.03	19.16	
4 μm 100 μg/L	32.40	15.68	
4 μm 1000 μg/L	16.87	15.47	
10 μm 100 μg/L	22.44	12.93	
10 μm 1000 μg/L	20.06	9.77	

of the target platform (sixth day)

Figure 5C: Novelty scores (time spent (s) investigating novel object/time spent (s)

investigating both objects in total for mice exposed or not to PS-MPs

	mean	SD	
Control	0.46	0.07	
0.5 μm 100 μg/L	0.26	0.13	
0.5 μm 1000 μg/L	0.24	0.13	
4 μm 100 μg/L	0.27	0.13	
4 μm 1000 μg/L	0.26	0.07	
10 μm 100 μg/L	0.26	0.13	
10 μm 1000 μg/L	0.22	0.09	

Figure 7B: Number of dendritic spines in hippocampus of mice exposed or not to PS-MPs

	mean	SD	
Control	12.25	0.83	
0.5 μm 1000 μg/L	6.50	0.50	
4 μm 1000 μg/L	4.50	0.50	
10 μm 1000 μg/L	5.25	1.09	

Figure 7C: Protein expression of synapsin 1 in the hippocampus of mice exposed or

not to PS-MPs

Synapsin/GAPDH ratio	mean	SD	
Control	1.0222	0.0299	
0.5 μm 100 μg/L	0.7478	0.0935	
0.5 μm 1000 μg/L	0.4472	0.0523	
4 μm 100 μg/L	0.7969	0.0932	
4 μm 1000 μg/L	0.5002	0.0585	
10 μm 100 μg/L	0.8760	0.1095	
10 μm 1000 μg/L	0.6659	0.0778	

Figure 7C: Protein expression of synaptophysin in the hippocampus of mice exposed

or not to PS-MPs

Synaptophysin/GAPDH	mean	SD	
ratio			
Control	1.0039	0.0236	

0.5 μm 100 μg/L	0.7508	0.0576	
0.5 μm 1000 μg/L	0.3228	0.1531	
4 μm 100 μg/L	0.5738	0.0705	
4 μm 1000 μg/L	0.3609	0.0616	
10 μm 100 μg/L	0.5631	0.0672	
10 μm 1000 μg/L	0.2965	0.1265	

Figure 7C: Protein expression of PSD95 in the hippocampus of mice exposed or not to

PS-MPs

PSD95/GAPDH ratio	mean	SD	
Control	1.0097	0.0159	
0.5 μm 100 μg/L	0.7614	0.0743	
0.5 μm 1000 μg/L	0.4557	0.0828	
4 μm 100 μg/L	0.7880	0.0754	
4 μm 1000 μg/L	0.4572	0.0446	
10 μm 100 μg/L	0.8130	0.1168	
10 μm 1000 μg/L	0.8368	0.0817	

Figure 7D: mRNA expression of *Bdnf* in the hippocampus of mice exposed or not to

PS-MPs

Bdnf/Gapdh ratio	mean	SD	
Control	1.0015	0.0397	
0.5 μm 100 μg/L	0.6347	0.0382	
0.5 μm 1000 μg/L	0.7628	0.0121	

4 μm 100 μg/L	0.7141	0.0634	
4 μm 1000 μg/L	0.7258	0.0457	
10 μm 100 μg/L	0.6733	0.0411	
10 μm 1000 μg/L	0.6889	0.0312	

Figure 7E: mRNA expression of *syt1* in the hippocampus of mice exposed or not to PS-

MPs

syt1/Gapdh ratio	mean	SD	
Control	1.0009	0.0275	
0.5 μm 100 μg/L	0.7317	0.0592	
0.5 μm 1000 μg/L	0.8920	0.0336	
4 μm 100 μg/L	0.9568	0.0192	
4 μm 1000 μg/L	0.8523	0.0023	
10 μm 100 μg/L	0.8485	0.0230	
10 μm 1000 μg/L	0.8965	0.0141	

Figure 8: Expression various inflammation-related mRNAs in the hippocampus of

mice exposed or not to PS-MPs

<i>Tnf-α/Gapdh</i> ratio	mean	SD	
Control	1.0004	0.0344	
0.5 μm 100 μg/L	2.2895	0.3995	
0.5 μm 1000 μg/L	4.5516	0.5359	
4 μm 100 μg/L	5.4554	0.7650	
4 μm 1000 μg/L	14.0082	1.4922	

3.3178	0.5503				
4.5896	0.0068				
mean	SD				
1.0162	0.1777				
2.1331	0.3062				
7.2351	2.2340				
4.5054	0.8103				
10.8836	2.9398				
7.2899	1.8559				
12.4737	1.7237				
mean	SD				
1.0261	0.2181				
1.1366	0.0387				
1.2323	0.0274				
1.3007	0.0333				
1.1565	0.0153				
1.0271	0.0076				
1.0865	0.0667				
mean	SD				
	3.3178 4.5896 mean 1.0162 2.1331 7.2351 4.5054 10.8836 7.2899 12.4737 mean 1.0261 1.1366 1.2323 1.3007 1.1565 1.0271 1.0865	3.3178 0.5503 4.5896 0.0068 mean SD 1.0162 0.1777 2.1331 0.3062 7.2351 2.2340 4.5054 0.8103 10.8836 2.9398 7.2899 1.8559 12.4737 1.7237 mean SD 1.0261 0.2181 1.1366 0.0387 1.2323 0.0274 1.3007 0.0333 1.1565 0.0153 1.0271 0.0076 1.0865 0.0667	3.3178 0.5503 4.5896 0.0068 mean SD 1.0162 0.1777 2.1331 0.3062 7.2351 2.2340 4.5054 0.8103 10.8836 2.9398 7.2899 1.8559 12.4737 1.7237 mean SD mean SD 1.0261 0.2181 1.1366 0.0387 1.3007 0.0333 1.1565 0.0153 1.0271 0.0076 1.0865 0.0667		

Control	1.0066	0.1162	
0.5 μm 100 μg/L	1.6457	0.0567	
0.5 μm 1000 μg/L	1.8675	0.3591	
4 μm 100 μg/L	1.9543	0.3027	
4 μm 1000 μg/L	3.1457	0.6149	
10 μm 100 μg/L	2.8568	0.6061	
10 μm 1000 μg/L	3.8213	0.5092	
Cxcl10/Gapdh ratio	mean	SD	
Control	1.0086	0.1404	
0.5 μm 100 μg/L	3.9309	0.5711	
0.5 μm 1000 μg/L	3.1314	0.7554	
4 μm 100 μg/L	5.3048	0.1642	
4 μm 1000 μg/L	4.4966	0.8317	
10 μm 100 μg/L	4.1451	0.5905	
10 μm 1000 μg/L	6.1071	1.5465	

Figure S4: Average swim speed (mm/s) on days 1-5 in the Morris water maze test for

mice exposed or not to PS-MPs

	mean	SD	mean	SD
	Day 1		y 1 Day 2	
Control	7.23	3.47	9.07	3.06
0.5 μm 100 μg/L	10.63	3.39	10.63	3.37

0.5 μm 1000 μg/L	10.51	2.13	9.93	2.43
4 μm 100 μg/L	9.70	1.63	11.41	3.14
4 μm 1000 μg/L	9.72	2.89	11.57	2.61
10 μm 100 μg/L	10.69	2.78	10.20	3.69
10 µm 1000 µg/L	7.00	3.39	8.01	3.08
	Da	y 3	Da	ny 4
Control	9.60	2.40	8.94	4.48
0.5 μm 100 μg/L	6.15	4.31	8.90	3.38
0.5 μm 1000 μg/L	8.13	4.14	6.99	4.12
4 μm 100 μg/L	10.37	2.76	8.19	2.65
4 μm 1000 μg/L	10.64	3.59	6.70	3.85
10 μm 100 μg/L	12.04	3.15	8.47	1.91
10 μm 1000 μg/L	7.19	3.74	5.65	3.53
	Da	y 5		
Control	6.99	3.41		
0.5 μm 100 μg/L	8.95	4.36		
0.5 μm 1000 μg/L	7.53	2.92		
4 μm 100 μg/L	8.46	3.27		
4 μm 1000 μg/L	8.70	3.32		
10 μm 100 μg/L	10.11	2.88		
10 μm 1000 μg/L	5.79	3.98		

Figure S5: Total path length to find the hidden platform (mm) of mice exposed or not to PS-MPs on days 1-5 in Morris water maze test

	mean	SD	mean	SD
	Da	y 1	Da	ny 2
Control	605.28	302.50	484.55	328.86
0.5 μm 100 μg/L	751.98	403.53	462.85	177.71
0.5 μm 1000 μg/L	661.03	326.06	607.90	270.06
4 μm 100 μg/L	699.20	218.69	668.47	382.04
4 μm 1000 μg/L	807.78	297.77	817.31	330.70
10 μm 100 μg/L	770.91	268.20	601.62	293.47
10 μm 1000 μg/L	561.54	291.52	551.15	306.82
	Day 3		Day 4	
Control	546.82	376.23	347.48	290.25
0.5 μm 100 μg/L	259.47	339.36	487.42	313.87
0.5 μm 1000 μg/L	440.49	419.87	395.86	296.70
4 μm 100 μg/L	608.05	258.77	382.22	245.75
4 μm 1000 μg/L	643.65	318.67	407.91	311.49
10 μm 100 μg/L	747.16	339.14	394.88	220.00
10 μm 1000 μg/L	440.00	313.13	348.95	308.49
	Day 5			
Control	184.70	129.15		
0.5 μm 100 μg/L	318.53	313.21		

0.5 μm 1000 μg/L	251.68	234.49	
4 μm 100 μg/L	324.23	248.26	
4 μm 1000 μg/L	377.74	187.09	
10 μm 100 μg/L	442.81	414.12	
10 μm 1000 μg/L	230.22	115.40	

Figure S6A: The recognition scores (time spent (s) investigating one object/time spent

(s) investigating both objects in total) of mice exposed or not to PS-MPs on day 1

	mean	SD	
Control	0.50	0.01	
0.5 μm 100 μg/L	0.50	0.01	
0.5 μm 1000 μg/L	0.50	0.01	
4 μm 100 μg/L	0.50	0.01	
4 μm 1000 μg/L	0.49	0.01	
10 μm 100 μg/L	0.49	0.01	
10 μm 1000 μg/L	0.50	0.01	

Figure S6B: The recognition scores (time spent (s) investigating one object/time spent

(s) investigating both objects in total) of mice exposed or not to PS-MPs on day 2

	mean	SD	
Control	0.50	0.01	
0.5 μm 100 μg/L	0.50	0.01	
0.5 μm 1000 μg/L	0.51	0.01	
4 μm 100 μg/L	0.50	0.01	

4 μm 1000 μg/L	0.50	0.01	
10 μm 100 μg/L	0.50	0.01	
10 μm 1000 μg/L	0.49	0.01	

Figure S7A: The total distance moved (m) for mice (exposed or not to PS-MPs) in the

novel object recognition testing day (day 3)

	mean	SD	
Control	4.10	0.80	
0.5 μm 100 μg/L	4.04	1.05	
0.5 μm 1000 μg/L	3.63	0.97	
4 μm 100 μg/L	3.95	1.32	
4 μm 1000 μg/L	3.37	1.29	
10 μm 100 μg/L	3.63	1.21	
10 μm 1000 μg/L	2.73	1.11	

Figure S7B: The average movement speed (mm/s) for mice (exposed or not to PS-

MPs) in the novel object recognition testing day (day 3)

	mean	SD	
Control	34.19	6.63	
0.5 μm 100 μg/L	33.66	8.75	
0.5 μm 1000 μg/L	30.23	8.06	
4 μm 100 μg/L	32.90	11.03	
4 μm 1000 μg/L	28.06	10.79	
10 μm 100 μg/L	30.23	10.08	

10 μm 1000 μg/L	22.73	9.29		
Figure S8A: Number of neurons in the hippocampal CA1 for mice exposed or not to				
PS-MPs				
	mean	SD		
Control	1.01	0.02		
0.5 μm 100 μg/L	1.00	0.03		
0.5 μm 1000 μg/L	0.98	0.04		
4 μm 100 μg/L	0.97	0.04		
4 μm 1000 μg/L	0.97	0.04		
10 μm 100 μg/L	0.99	0.03		
10 μm 1000 μg/L	0.99	0.05		
Figure S8B: Number of neu	rons in the hip	pocampal CA3	for mice expos	sed or not to

Figure S8B: Number of neurons in the hippocampal CA3 for mice exposed or no

PS-MPs

	mean	SD	
Control	1.02	0.03	
0.5 μm 100 μg/L	0.98	0.03	
0.5 μm 1000 μg/L	0.98	0.03	
4 μm 100 μg/L	0.95	0.06	
4 μm 1000 μg/L	0.98	0.06	
10 μm 100 μg/L	0.98	0.05	
10 μm 1000 μg/L	0.96	0.06	

Figure S8C: Number of neurons in the dentate gyrus for mice exposed or not to PS-MPs

	mean	SD	
Control	1.01	0.02	
0.5 μm 100 μg/L	0.96	0.04	
0.5 μm 1000 μg/L	0.97	0.03	
4 μm 100 μg/L	0.93	0.05	
4 μm 1000 μg/L	0.94	0.06	
10 μm 100 μg/L	0.97	0.04	
10 μm 1000 μg/L	0.99	0.05	

Figure S9A: mRNA level of *caspase-3* in the hippocampus of mice exposed or not to

PS-MPs

Caspase-3/Gapdh ratio	mean	SD	
Control	0.9999	0.0171	
0.5 μm 100 μg/L	1.1327	0.0559	
0.5 µm 1000 µg/L	1.2987	0.1153	
4 μm 100 μg/L	1.1491	0.0267	
4 μm 1000 μg/L	1.3089	0.0838	
10 μm 100 μg/L	1.1281	0.0304	
10 μm 1000 μg/L	1.1773	0.0354	

Figure S9B: Bax/Bcl-2 mRNA ratio in the hippocampus of mice exposed or not to PS-

MPs

Bax/Bcl-2 ratio	mean	SD	
Control	1.0003	0.0372	
0.5 μm 100 μg/L	1.8476	0.1304	
0.5 μm 1000 μg/L	1.6885	0.1353	
4 μm 100 μg/L	1.8125	0.2710	
4 μm 1000 μg/L	1.6351	0.0880	
10 μm 100 μg/L	1.6239	0.0967	
10 μm 1000 μg/L	1.6136	0.0880	

Figure S11A: Expression of *Gap43* mRNA in hippocampus of mice exposed or not to

PS-MPs

Gap43/Gapdh ratio	mean	SD	
Control	1.0013	0.0408	
0.5 μm 100 μg/L	1.2930	0.2807	
0.5 μm 1000 μg/L	1.1059	0.0664	
4 μm 100 μg/L	1.3914	0.2779	
4 μm 1000 μg/L	1.2130	0.2703	
10 μm 100 μg/L	1.3225	0.3283	
10 μm 1000 μg/L	1.3995	0.2376	

Figure S11B: Expression of syt4 mRNA in hippocampus of mice exposed or not to PS-

MPs

syt4/Gapdh ratio	mean	SD	
Control	1.0013	0.0361	

0.5 μm 100 μg/L	1.3104	0.1811	
0.5 μm 1000 μg/L	1.1961	0.1022	
4 μm 100 μg/L	1.1193	0.0650	
4 μm 1000 μg/L	1.1186	0.0825	
10 μm 100 μg/L	1.0513	0.0765	
10 μm 1000 μg/L	1.0575	0.0550	

Figure S11C: Expression of *Ncam* mRNA in hippocampus of mice exposed or not to

PS-MPs

Ncam/Gapdh ratio	mean	SD	
Control	1.0007	0.0156	
0.5 μm 100 μg/L	1.2054	0.1273	
0.5 μm 1000 μg/L	1.0183	0.0461	
4 μm 100 μg/L	1.1795	0.1093	
4 μm 1000 μg/L	0.8776	0.1129	
10 μm 100 μg/L	0.9086	0.0136	
10 μm 1000 μg/L	0.9664	0.0340	

Figure S12B: Percent of total cells in view that were *Mcp-1* positive in the

hippocampus of mice exposed or not to PS-MPs

	mean	SD	
Control	5.50	1.08	
0.5 μm 100 μg/L	14.67	2.75	
0.5 μm 1000 μg/L	15.30	2.79	

4 μm 100 μg/L	15.33	1.45	
4 μm 1000 μg/L	13.43	2.66	
10 μm 100 μg/L	14.57	2.65	
10 μm 1000 μg/L	18.43	1.72	

Figure S12C: Percent of total cells in view that were TNF- α positive in the hippocampus

of mice exposed or not to PS-MPs

Figure S12C	mean	SD	
Control	4.90	1.14	
0.5 μm 100 μg/L	14.83	2.13	
0.5 μm 1000 μg/L	15.70	1.80	
4 μm 100 μg/L	21.20	3.24	
4 μm 1000 μg/L	20.03	2.42	
10 μm 100 μg/L	20.87	2.27	
10 μm 1000 μg/L	15.40	1.61	

Table S4. *P*-values for non-statistically and statistically significant results.

Figure 4B: Time for mice (exposed or not to PS-MPs) to find the platform during				
learning trials (seconds)				
	P value	Summary	P value	Summary
	Day1		Day 2	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.6127	ns	0.9986	ns
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.3880	ns	0.4518	ns

Control vs 4 µm 100 µg/L PS-MPs group	0.5606	ns	0.4541	ns
Control vs 4 µm 1000 µg/L PS-MPs group	0.6864	ns	0.2371	ns
Control vs 10 µm 100 µg/L PS-MPs group	0.8004	ns	0.4523	ns
Control vs 10 µm 1000 µg/L PS-MPs group	0.3697	ns	0.3920	ns
	Da	y 3	Da	ay 4
Control vs 0.5 µm 100 µg/L PS-MPs group	0.7560	ns	0.4728	ns
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.9719	ns	0.4530	ns
Control vs 4 µm 100 µg/L PS-MPs group	0.6928	ns	0.5764	ns
Control vs 4 µm 1000 µg/L PS-MPs group	0.5247	ns	0.2119	ns
Control vs 10 µm 100 µg/L PS-MPs group	0.6502	ns	0.5973	ns
Control vs 10 µm 1000 µg/L PS-MPs group	0.3577	ns	0.1016	ns
	Da	y 5		
Control vs 0.5 µm 100 µg/L PS-MPs group	0.4846	ns		
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.2593	ns		
Control vs 4 µm 100 µg/L PS-MPs group	0.2743	ns		
Control vs 4 µm 1000 µg/L PS-MPs group	0.0457	*		
Control vs 10 µm 100 µg/L PS-MPs group	0.3500	ns		
Control vs 10 µm 1000 µg/L PS-MPs group	0.0254	*		

Figure 4D: Number of times mice (exposed or not to PS-MPs) crossed the target platform

on sixth day

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0069	**	

Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0031	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0039	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0009	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0057	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0016	**	

Figure 4E: Percentage of time mice (exposed or not to PS-MPs) took in the quadrant of

the target platform (sixth day)

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0071	**	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0035	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0352	*	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0028	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0066	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0013	**	

Figure 5C: Novelty scores (time spent (s) investigating novel object/time spent (s)

investigating both objects in total for mice exposed or not to PS-MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0026	**	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0011	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0046	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0000	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0021	**	

Control vs 10 µm 1000 µg/L PS-MPs group	0.0001	**			
Figure 7B: Number of dendritic spines in h	ippocampus	of mice exp	osed or not	to PS-MPs	
	P value	Summary			
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0001	**			
Control vs 4 µm 1000 µg/L PS-MPs group	0.0000	**			
Control vs 10 µm 1000 µg/L PS-MPs group	0.0001	**			
Figure 7C: Protein expression of synapsin 1 in the hippocampus of mice exposed or not					
to PS-MPs					
	P value	Summary			
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0168	*			
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0002	**			
Control vs 4 µm 100 µg/L PS-MPs group	0.0312	*			
Control vs 4 µm 1000 µg/L PS-MPs group	0.0004	**			
Control vs 10 µm 100 µg/L PS-MPs group	0.1427	ns			
Control vs 10 µm 1000 µg/L PS-MPs group	0.0038	**			

Figure 7C: Protein expression of synaptophysin in the hippocampus of mice exposed or

not to PS-MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0045	**	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0034	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0012	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0002	**	

Control vs 10 µm 100 µg/L PS-MPs group	0.0009	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0015	**	

Figure 7C: Protein expression of PSD95 in the hippocampus of mice exposed or not to

PS-MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0099	**	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0007	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0152	*	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0001	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0777	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0425	*	

Figure 7D: mRNA expression of *Bdnf* in the hippocampus of mice exposed or not to PS-

MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0007	**	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0012	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0056	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0030	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0013	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0009	**	

Figure 7E: mRNA expression of syt1 in the hippocampus of mice exposed or not to PS-

MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0043	**	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0238	*	
Control vs 4 µm 100 µg/L PS-MPs group	0.1363	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0016	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0038	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0087	**	

Figure 8: Expression various inflammation-related mRNAs in the hippocampus of mice

exposed or not to PS-MPs

Tnf-a	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0104	*	
Control vs 0.5 μm 1000 μg/L PS-MPs group	0.0007	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0012	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0002	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0040	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0000	**	
Mcp-1	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0111	*	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0172	*	
Control vs 4 µm 100 µg/L PS-MPs group	0.0040	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0091	**	

Control vs 10 µm 100 µg/L PS-MPs group	0.0089	**		
Control vs 10 µm 1000 µg/L PS-MPs group	0.0007	**		
<i>Il-6</i>	P value	Summary		
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0179	*		
Control vs 0.5 μm 1000 μg/L PS-MPs group	0.0015	**		
Control vs 4 µm 100 µg/L PS-MPs group	0.0012	**		
Control vs 4 µm 1000 µg/L PS-MPs group	0.0008	**		
Control vs 10 µm 100 µg/L PS-MPs group	0.0173	*		
Control vs 10 µm 1000 µg/L PS-MPs group	0.1801	ns		
ΙΙ-1β	P value	Summary		
<i>Il-1β</i> Control vs 0.5 μm 100 μg/L PS-MPs group	P value 0.0022	Summary **		
<i>Il-1β</i> Control vs 0.5 μm 100 μg/L PS-MPs group Control vs 0.5 μm 1000 μg/L PS-MPs group	P value 0.0022 0.0321	Summary ** *		
<i>Il-1β</i> Control vs 0.5 μm 100 μg/L PS-MPs group Control vs 0.5 μm 1000 μg/L PS-MPs group Control vs 4 μm 100 μg/L PS-MPs group	P value 0.0022 0.0321 0.0144	Summary ** * *		
<i>Il-1β</i> Control vs 0.5 μm 100 μg/L PS-MPs group Control vs 0.5 μm 1000 μg/L PS-MPs group Control vs 4 μm 100 μg/L PS-MPs group Control vs 4 μm 1000 μg/L PS-MPs group	P value 0.0022 0.0321 0.0144 0.0084	Summary ** * * * *		
Il-1βControl vs 0.5 μm 100 μg/L PS-MPs groupControl vs 0.5 μm 1000 μg/L PS-MPs groupControl vs 4 μm 100 μg/L PS-MPs groupControl vs 4 μm 1000 μg/L PS-MPs groupControl vs 10 μm 100 μg/L PS-MPs group	P value 0.0022 0.0321 0.0144 0.0084 0.0133	Summary ** * * * * * * *		
Il-1βControl vs 0.5 µm 100 µg/L PS-MPs groupControl vs 0.5 µm 1000 µg/L PS-MPs groupControl vs 4 µm 100 µg/L PS-MPs groupControl vs 4 µm 1000 µg/L PS-MPs groupControl vs 10 µm 100 µg/L PS-MPs groupControl vs 10 µm 1000 µg/L PS-MPs group	P value 0.0022 0.0321 0.0144 0.0084 0.0133 0.0016	Summary ** * * * * * * * * *		
II-1βControl vs 0.5 μm 100 μg/L PS-MPs groupControl vs 0.5 μm 1000 μg/L PS-MPs groupControl vs 4 μm 100 μg/L PS-MPs groupControl vs 4 μm 1000 μg/L PS-MPs groupControl vs 10 μm 1000 μg/L PS-MPs groupControl vs 10 μm 1000 μg/L PS-MPs group	P value 0.0022 0.0321 0.0144 0.0084 0.0133 0.0016	Summary ** * * * * * * * *		
<i>II-1β</i> Control vs 0.5 μm 100 μg/L PS-MPs group Control vs 0.5 μm 1000 μg/L PS-MPs group Control vs 4 μm 100 μg/L PS-MPs group Control vs 10 μm 100 μg/L PS-MPs group Control vs 10 μm 1000 μg/L PS-MPs group	P value 0.0022 0.0321 0.0144 0.0084 0.0133 0.0016 P value	Summary ** * * * * Summary		
II-1β Control vs 0.5 μm 100 μg/L PS-MPs group Control vs 0.5 μm 1000 μg/L PS-MPs group Control vs 4 μm 100 μg/L PS-MPs group Control vs 4 μm 1000 μg/L PS-MPs group Control vs 10 μm 1000 μg/L PS-MPs group	P value 0.0022 0.0321 0.0144 0.0084 0.0133 0.0016 P value 0.0022	Summary ** * * * Summary **		

Control vs 4 µm 100 µg/L PS-MPs group	0.0000	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0043	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0019	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0097	**	

Figure S4: Average swim speed (mm/s) on days 1-5 in the Morris water maze test for mice

exposed or not to PS-MPs

	P value	Summary	P value	Summary
	Day 1		Day 2	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0853	ns	0.3790	ns
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0512	ns	0.5696	ns
Control vs 4 µm 100 µg/L PS-MPs group	0.1107	ns	0.1967	ns
Control vs 4 µm 1000 µg/L PS-MPs group	0.1674	ns	0.1223	ns
Control vs 10 µm 100 µg/L PS-MPs group	0.0584	ns	0.5439	ns
Control vs 10 µm 1000 µg/L PS-MPs group	0.9004	ns	0.5270	ns
	Day 3		Day 4	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0858	ns	0.9848	ns
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.4306	ns	0.4103	ns
Control vs 4 µm 100 µg/L PS-MPs group	0.5860	ns	0.7105	ns
Control vs 4 µm 1000 µg/L PS-MPs group	0.5346	ns	0.3320	ns
Control vs 10 µm 100 µg/L PS-MPs group	0.1255	ns	0.8036	ns
Control vs 10 µm 1000 µg/L PS-MPs group	0.1734	ns	0.1491	ns
	Day 5			

Control vs 0.5 µm 100 µg/L PS-MPs group	0.3632	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.7552	ns	
Control vs 4 µm 100 µg/L PS-MPs group	0.4222	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.3570	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.0851	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.5563	ns	

Figure S5: Total path length to find the hidden platform (mm) of mice exposed or not to

PS-MPs on days 1-5 in Morris water maze test

	P value	Summary	P value	Summary
Figure S5	Da	у 1	Day 2	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.4543	ns	0.8801	ns
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.7451	ns	0.4559	ns
Control vs 4 µm 100 µg/L PS-MPs group	0.5164	ns	0.3508	ns
Control vs 4 µm 1000 µg/L PS-MPs group	0.2275	ns	0.0780	ns
Control vs 10 µm 100 µg/L PS-MPs group	0.2967	ns	0.4938	ns
Control vs 10 µm 1000 µg/L PS-MPs group	0.7869	ns	0.7011	ns
	Day 3		Day 4	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.1557	ns	0.4011	ns
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.6255	ns	0.7623	ns
Control vs 4 µm 100 µg/L PS-MPs group	0.7281	ns	0.8126	ns
Control vs 4 µm 1000 µg/L PS-MPs group	0.6115	ns	0.7129	ns
Control vs 10 µm 100 µg/L PS-MPs group	0.3131	ns	0.7357	ns

Control vs 10 µm 1000 µg/L PS-MPs group	0.5729	ns	0.9928	ns
	Da	Day 5		
Control vs 0.5 µm 100 µg/L PS-MPs group	0.3137	ns		
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.5187	ns		
Control vs 4 µm 100 µg/L PS-MPs group	0.2083	ns		
Control vs 4 µm 1000 µg/L PS-MPs group	0.0413	*		
Control vs 10 µm 100 µg/L PS-MPs group	0.1378	ns		
Control vs 10 µm 1000 µg/L PS-MPs group	0.4982	ns		

Figure S6A: The recognition scores (time spent (s) investigating one object/time spent (s)

investigating both objects in total) of mice exposed or not to PS-MPs on day 1

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.3473	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.9779	ns	
Control vs 4 µm 100 µg/L PS-MPs group	0.5071	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.2017	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.1485	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.4350	ns	

Figure S6B: The recognition scores (time spent (s) investigating one object/time spent (s)

investigating both objects in total) of mice exposed or not to PS-MPs on day 2

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.5867	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.4449	ns	

Control vs 4 µm 100 µg/L PS-MPs group	0.5676	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.9157	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.9893	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.3454	ns	

Figure S7A: The total distance moved (m) for mice (exposed or not to PS-MPs) in the

novel object recognition testing day (day 3)

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.6502	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.8094	ns	
Control vs 4 µm 100 µg/L PS-MPs group	0.8052	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.2210	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.4001	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0188	*	

Figure S7B: The average movement speed (mm/s) for mice (exposed or not to PS-MPs) in

the novel object recognition testing day (day 3)

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.6505	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.8090	ns	
Control vs 4 µm 100 µg/L PS-MPs group	0.9811	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.8047	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.8165	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0188	*	

Figure S8A: Number of neurons in the hippocampal CA1 for mice exposed or not to PS-

MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.5243	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.1837	ns	
Control vs 4 µm 100 µg/L PS-MPs group	0.0810	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.1404	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.3345	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.5133	ns	

Figure S8B: Number of neurons in the hippocampal CA3 for mice exposed or not to PS-

MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.1137	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.1109	ns	
Control vs 4 µm 100 µg/L PS-MPs group	0.0887	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.3072	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.1756	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.1308	ns	

Figure S8C: Number of neurons in the dentate gyrus for mice exposed or not to PS-MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0527	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0461	*	

Control vs 4 µm 100 µg/L PS-MPs group	0.0313	*	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0819	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.1050	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.5691	ns	

Figure S9A: mRNA level of *caspase-3* in the hippocampus of mice exposed or not to PS-

MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0326	*	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0223	*	
Control vs 4 µm 100 µg/L PS-MPs group	0.0027	**	
Control vs 4 μ m 1000 μ g/L PS-MPs group	0.0069	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0065	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0031	**	

Figure S9B: *Bax/Bcl-2* mRNA ratio in the hippocampus of mice exposed or not to PS-MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0009	**	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0023	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0137	*	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0007	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0010	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0008	**	

Figure S11A: Expression of *Gap43* mRNA in hippocampus of mice exposed or not to PS-

MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.2195	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.1306	ns	
Control vs 4 µm 100 µg/L PS-MPs group	0.1210	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.3349	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.2416	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0798	ns	

Figure S11B: Expression of syt4 mRNA in hippocampus of mice exposed or not to PS-MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0770	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0638	ns	
Control vs 4 µm 100 µg/L PS-MPs group	0.0882	ns	
Control vs 4 μ m 1000 μ g/L PS-MPs group	0.1392	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.4501	ns	
Control vs 10 µm 1000 µg/L PS-MPs group	0.2932	ns	

Figure S11C: Expression of Ncam mRNA in hippocampus of mice exposed or not to PS-

MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0869	ns	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.6351	ns	

Control vs 4 µm 100 µg/L PS-MPs group	0.0839	ns	
Control vs 4 µm 1000 µg/L PS-MPs group	0.2013	ns	
Control vs 10 µm 100 µg/L PS-MPs group	0.0033	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.2646	ns	

Figure S12B: Percent of total cells in view that were *Mcp-1* positive in the hippocampus

of mice exposed or not to PS-MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0118	*	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0098	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0015	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0174	*	
Control vs 10 µm 100 µg/L PS-MPs group	0.0110	*	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0008	**	

Figure S12C: Percent of total cells in view that were TNF-a positive in the hippocampus

of mice exposed or not to PS-MPs

	P value	Summary	
Control vs 0.5 µm 100 µg/L PS-MPs group	0.0044	**	
Control vs 0.5 µm 1000 µg/L PS-MPs group	0.0020	**	
Control vs 4 µm 100 µg/L PS-MPs group	0.0026	**	
Control vs 4 µm 1000 µg/L PS-MPs group	0.0013	**	
Control vs 10 µm 100 µg/L PS-MPs group	0.0009	**	
Control vs 10 µm 1000 µg/L PS-MPs group	0.0017	**	

Table S5. Thigmotaxis of mice in Morris water maze test. Thigmotaxis is defined as mice were swimming in the outer 10% close to walls, which means mice swim almost exclusively in the periphery. The rate of thigmotaxis was expressed as the ratio of the number of mice showing thigmotaxis to the total number of mice according to the swim path trajectories.

	Trials	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6
Control	40	9 (22.5%)	8 (20%)	7 (17.5%)	5 (12.5%)	4 (10%)	4 (10%)
100 μg/L 0.5 μm	40	8 (20%)	7 (17.5%)	7 (17.5%)	6 (15%)	4 (10%)	4 (10%)
1000 μg/L 0.5 μm	40	9 (22.5%)	8 (20%)	8 (20%)	6 (15%)	5 (12.5%)	3 (7.5%)
100 μg/L 4 μm	40	8 (20%)	8 (20%)	7 (17.5%)	5 (12.5%)	5 (12.5%)	4 (10%)
1000 μg/L 4 μm	40	8 (20%)	8 (20%)	6 (15%)	6 (15%)	4 (10%)	4 (10%)
100 μg/L 10 μm	40	8 (20%)	8 (20%)	7 (17.5%)	7 (17.5%)	6 (15%)	3 (7.5%)
1000 μg/L 10 μm	40	9 (22.5%)	8 (20%)	7 (17.5%)	7 (17.5%)	6 (15%)	4 (10%)



10 μm (1000 μg/L)



Figure S1. Biodistribution of various diameters of polystyrene MPs (PS-MPs) in different tissues of mice. Mice were provided drinking water containing three sizes of fluorescent PS-MPs for 180 consecutive days. Fluorescence pictures of excised brain tissues and gastric tissues were detected. The colors in pictures indicated that fluorescence PS-MPs in tissues.

	Liver	Hippocampus	Hypothalamus	Cortex
Control			finder and	
0.5 μm (100 μg/L)				
0.5 μm (1000 μg/L)				
4 μm (100 μg/L)				
4 μm (1000 μg/L)				
10 μm (100 μg/L)				
10 μm (1000 μg/L)			Sel	20 µm



Figure S2. BBB integrity was detected by biotin tracer experiments. The sections were stained with biotin (red) and DAPI (blue). The existence of biotin in brain tissues was examined by immunofluorescence microscopy. The biotin signal of the liver parenchyma was used as a positive control (scale bar = $20 \mu m$).



Figure S3. Ultrastructure of BBB in PS-MP-exposed and unexposed mice as detected by an electron microscope. Upper images displayed the ultrastructure of BBB. Lower images referred to magnified boxed areas.



Figure S4. The average swim speed of mice on days 1-5 in Morris water maze test is shown. Circle (Control), square (100 μ g/L), triangle (1000 μ g/L). (A) Control and 0.5 μ m PS-MPs exposure groups. (B) Control and 4 μ m PS-MPs exposure groups. (C) Control and 10 μ m PS-

MPs exposure groups. The results are expressed as means \pm SD (n = 10 mice/per group). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. Data were detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.



Figure S5. The total path length to find the hidden platform of mice on days 1-5 in Morris water maze test are shown. Circle (Control), square (100 µg/L), triangle (1000 µg/L). (A) Control and 0.5 µm PS-MPs exposure groups. (B) Control and 4 µm PS-MPs exposure groups. (C) Control and 10 µm PS-MPs exposure groups. The results are expressed as means \pm SD (n = 10 mice/per group). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05 vs. control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.



Figure S6. The recognition scores (time spent (s) investigating one object/time spent (s) investigating both objects in total) of mice on day 1 (A) and day 2 (B) of novel object recognition (NOR) experiment were examined (n = 10 mice/per group). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4.



Figure S7. The total distance moved (A) and average movement speed (B) of mice in testing day (day 3) of the novel object recognition test are shown. The results are expressed as means \pm SD (n = 10 mice/per group). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05 vs. control, as

detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.



Figure S8. The numbers of neurons in the hippocampal cornu ammonis 1 (CA1), cornu ammonis 3 (CA3), and dentate gyrus (DG) sections were counted (n = 3 mice/group, n = 3 slides/mice). Data are shown as mean \pm SD. The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05 compared with control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.



Figure S9. The mRNA levels of *caspase 3* (A), the *Bax/Bcl-2* mRNA ratio (B) in the hippocampus were tested with quantitative real-time PCR (qRT-PCR) by normalizing to *Gapdh*. The results are expressed as means \pm SD (n = 3, N = 3 mice/group). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05 and ***P* < 0.01 compared with the control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.



Figure S10. The expression of synapsin 1, synaptophysin, and PSD 95 protein in the hippocampus was measured via western blotting. The western blotting results were quantified and statistically analyzed, as shown in (Figure 7C) as mean \pm SD (n = 3, N = 3 mice/group). The mean and SD summary data for quantification of western blotting are shown in Table S3.

P-Values for all comparisons are reported in Table S4.



Figure S11. The mRNA expression levels of *Gap43*(A), *Syt 4* (B), and *Ncam*(C) were detected by qRT-PCR in the hippocampus of mice exposed to PS-MPs and control mice. Data are shown as mean \pm SD (n = 3). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. ***P* < 0.01 compared with control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.



Figure S12. (A) A confocal fluorescence microscope was used to detect colocalization of Alexa Fluor 594-labled MCP-1 (red)/TNF- α (red) and DAPI (blue) (scale bar = 20 μ m, n = 3) in the hippocampus of mice exposed to PS-MPs and control mice. (B) Percent of positivity was calculated based on the percentage of MCP-1 positive cells out of the total number of cells in an image. (C) Percent of positivity was calculated based on the percentage of TNF- α positive cells out of the total number of cells in an image. Data are shown as mean \pm SD (n = 3). The mean and SD summary data for quantification are shown in Table S3. *P*-Values for all comparisons are reported in Table S4. **P* < 0.05, ***P* < 0.01 compared with control, as detected by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparison tests.