## **Supplementary materials**

**Supplementary Figure 1.** Purity of primary cultured microglia. (A) Phase-contrast image of primary cultured microglia. (B) Purity of microglia was identified by Iba1 staining. Scale bar =  $30 \mu m$ . (C) Quantification of the Iba1-positive microglia. Note that about 90% of the cells are Iba1 positive.

**Supplementary Figure 2.** Effects of OECs-Exo on expression of IRF3 and p-c-Jun upon LPS treatment. (**A**) The mRNA level of IRF3 in microglia under normal condition, LPS or LPS plus OECs-Exo treatment. (**B**) Quantification of expression level of p-c-Jun in microglia under control, LPS or LPS plus OECs-Exo treatment. N = 3, \*P < 0.05, \*\*P < 0.01. (**C**) Immunostaining of p-c-Jun and F4/80. Scale bar = 30  $\mu$ m. (**D**-**E**) Quantification of the IFI/area and the nuclei/cytosol fluorescence intensity ratio of p-c-Jun. Note that OECs-Exo significantly supressed the increased IFI and nulear translocation of p-c-Jun induced by LPS. N = 3, \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001.

**Supplementary Figure 3.** (A) Representative confocal microscopy images showing uptake of OECs-Exo (red) by microglia/macrophages (green) in spinal cord at 1dpi, 5dpi, 10dpi. Scale bar=50  $\mu$ m. (B) Quantification of the percentage of internalization of OECs-Exo by microglia/macrophages. Results are presented as mean  $\pm$  SEM. N = 3/group.

**Supplementary Figure 4.** Full gels of CD63, CD9, pNF- $\kappa$ B, NF200, p-c-Jun,  $\beta$ -actin and GAPDH.

Sup Table 1.	PCR primer sequences	
Gene	Forward primer (5'-3')	Reverse primer (5'-3')
iNOS	GCCTAGTCAACTACAAGCCCC	AGAAACTTCCAGGGGCAAGC
CD86	AGACATGTGTAACCTGCACCAT	ACTTTTTCCGGTCCTGCCAA
IL-12	GTGTCAATCACGCTACCTCCT	CTTGGCAGGTCCAGAGACTG
IL-18	CAGCTCTTCTACCAGCAAACAT	CTTCCAACTGAGAGGCTGTGC
CD206	GAGGACTGCGTGGTGATGAA	CATGCCGTTTCCAGCCTTTC
Arginase1	AAGATGTGCCCTCTGTCTTTTAG	CCCCTCCTCGAGGCTGTC
IL-4	CCTTGCTGTCACCCTGTTCT	CGGTGCATGGAGTCCCTTTT
IL-10	CGCTGTCATCGATTTCTCCC	TAGACACCTTTGTCTTGGAGCTTAT
TNFα	ATGGGCTCCCTCTCATCAGT	GCTTGGTGGTTTGCTACGAC
IRF-3	TGGCTGCGAGTCTCAACTAC	CAGTTCCTGAGCCAGGGGAG
GAPDH	GGCTCTCTGCTCCTCCCT	GTCTATGAGACGAGGCTGGC

Sup Table 1. PCR primer sequences

## Sup Figure 1



Sup Figure 2



Sup Figure 3



## Sup Figure 4













