# **Supplementary Information**

#### Role of miRNAs shuttled by mesenchymal stem cell-derived small extracellular vesicles in

#### modulating neuroinflammation

Debora Giunti, Chiara Marini, Benedetta Parodi, Cesare Usai, Marco Milanese, Giambattista Bonanno, Nicole Kerlero de Rosbo, Antonio Uccelli

#### Supplementary Data S1 online (.pdf)

# In-depth data analysis of miRNA expressed in IFN-γ-primed and unprimed MSCs identified by microarray analysis

Microarray analysis showed that miR-466q, miR-467g, miR-669c-3p, miR-466m-5p, miR-467f, miR-3082-5p, miR-466i-5p, miR-466i-3p and miR-5126 are significantly dysregulated in IFN-γ-primed MSCs compared to unprimed MSCs.

## Supplementary data S2 online (.xls)

List of the 1718 target genes predicted for miR-467f through miRWalk database

#### Supplementary data S3 online (.xls)

List of the 1157 target genes predicted for miR-466q through miRWalk database

# Supplementary Figure S1 online



## **Supplementary Figure S1 online**

# Validation of the significantly dysregulated miRNAs in immunomodulatory MSCs by RT-PCR

RT-PCR analysis on MSCs primed or not with IFN- $\gamma$  confirmed the significant dysregulation of the nine specific miRNAs upon IFN- $\gamma$  stimulation of MSCs. \**P* < 0.05, \*\**P* < 0.01. Data are presented as mean ± SEM of 3 independent experiments.



# **Supplementary Figure S2 online**

#### Supplementary Figure S2 online

# Characterization of s-EV isolated from IFN- $\gamma$ -primed MSCs through electron microscopy and Western blot

The presence of purified s-EV in the supernatant of IFN- $\gamma$ -primed MSCs was confirmed by electron microscopy which identified the presence of nanovesicles with a diameter range of 50-100 nm (representative image in panel **a**) and by western blot analysis showing the expression of the s-EV markers Alix and CD9 (one representative blot is shown in panel **b**)

# **Supplementary Figure S3 online**



#### Supplementary Figure S3 online

Positive control miRNA, which significantly decreases the mRNA expression of GAPDH, validates efficient transfection

Validation of transfection efficiency was assessed by RT-PCR analysis using a positive control (Cpos), which targets GAPDH expression. **a** cycle threshold and **b** relative quantification over HPRT, as reference gene. \*\*P < 0.01. Data are presented as mean  $\pm$  SEM of 3 independent experiments

# **Supplementary Figure S4 online**



## **Supplementary Figure S4 online**

# Expression of Tnf, II1b and Cx3cr1 by LPS-activated N9 cells transfected with miRNAs that are dysregulated in IFN- $\gamma$ -primed MSCs but not in their derived s-EV

RT-PCR analysis shows that miR-466i-5p, miR-467g, miR-3082-5p, and miR-669c-3p did not affect the expression of Tnf and Il1b in LPS-activated microglia, and that miR-3082-5p induced a significant increase in Cx3cr1 expression in activated N9 cells. ##P < 0.01, untreated (N9) vs LPS-activated N9 cells (N9+LPS); ###P < 0.001, N9 vs N9+LPS; \*\*P < 0.01 N9+LPS vs N9+LPS transfected with specific miRNA (N9+LPS+miRNA). Data are presented as mean ± SEM of 3 independent experiments.

# **Supplementary Figure S5 online**



#### Supplementary Figure S5 online Expression of miR-467f and miR-466q in microglia cell line activated or not with LPS

RT-PCR results demonstrated that N9 microglia line cells express both miR-467f and miR-466q at basal level, and that miRNA expression did not increase upon LPS-activation of the cells. Data are presented as mean  $\pm$  SEM of 3 independent experiments



# **Supplementary Figure S6 online**

## **Supplementary Figure S6 online**

#### LPS is a stronger more consistent activator of N9 cells that IFN- $\gamma$

**a** Analysis of Tnf and II1b mRNA expression by RT-PCR and **b** quantification of pp38 by Western blotting performed in N9 cells stimulated or not with IFN- $\gamma$  (100 ng/ml) or LPS (1 µg/ml) for 24h showed a stronger activation of N9 cell upon LPS activation in comparison to IFN- $\gamma$ . \**P* < 0.05, \*\**P* < 0.01, \*\*\* *P* < 0.001. Data are presented as mean ± SEM of 3 independent experiments (a) and as fold induction of 1 experiment (b).

# Supplementary Figure S7 online





Supplementary Figure S7 online Unedited gels of Supplementary Figure S6 online