Microglia extracellular vesicles: focus on molecular composition and biological function

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Keywords

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Abstract

Extracellular Vesicles (EVs) are a heterogeneous family of cell-derived lipid bounded vesicles comprising exosomes and microvesicles. They are potentially produced by all types of cells and are used as a cell-to-cell communication method that allows protein, lipid, and genetic material exchange. Microglia cells produce a large number of EVs both in resting and activated conditions, in the latter case changing their production and related biological effects. Several actions of microglia in the central nervous system are ascribed to EVs, but the molecular mechanisms by which each effect occurs are still largely unknown. Conflicting functions have been ascribed to microglia-derived EVs starting from the neuronal support and ending with the propagation of inflammation and neurodegeneration, confirming the crucial role of these organelles in tuning brain homeostasis. Despite the increasing number of studies reported on microglia EVs, there is also a lot of fragmentation in the knowledge on the mechanism at the basis of their production and modification of their cargo. In this review, a collection of literature data about the surface and cargo proteins and lipids as well as the miRNA content of EVs produced by microglial cells has been reported. A special highlight was given to the works in which the EV molecular composition is linked to a precise biological function.

Introduction

The term 'extracellular vesicles' defines a heterogeneous family of cell-derived membrane vesicles originating from membrane shedding, which constitutes a major component in cell-to-cell communication. The number of biological functions and cargos, either on the membrane and in the lumen, ascribed to EVs has witnessed an enormous increase in the last 20 years [1–3].

According to their size and biogenesis [4,5], EVs can be divided into Microvesicles (MVs) and exosomes [6–11]. MVs refer to vesicles released by either healthy or apoptotic cells originating from the outward budding of the plasma membrane [4], and typically have dimensions spanning from 150 to 800 nm in diameter. Exosomes have peculiar biogenesis, starting with the formation of a Multivesicular Body (MVB) by endocytosis and ending with exocytosis [6]. Exosomes are characterized by dimensions spanning from 30 to 250 nm in diameter. Exosome and MV isolation from a biological sample is tricky due to their partial overlap in size and biomarker composition. This is one of the reasons because now EVs are classified as small EVs with a size of 30–250 nm and large EVs with a size of 150–800 nm [5,7,11].

The International Society Of Extracellular Vesicles (ISEV) proposed the Minimal Information For Studies Of Extracellular Vesicles (MISEV) guidelines for the field in 2014 [12], which was later updated in 2018 [11], in which the minimal requirements to validate an EV sample are extensively described. Briefly, there is the need to test/confirm the presence of: i) at least one protein associated with the plasma membrane or endosomes, ii) one cytosolic protein associated with EVs, iii) one component of non-EVs co-isolated structures (e.g. lipoprotein), and iv) one component associated to the specific EV sub-type of interest [11]. Several studies name EVs as exosomes without clear evidence of the EV nature; however, different authors have recently begun to align to the aforementioned guidelines and provide concrete evidence of the exosome isolation [7,13–15]. To avoid confusion, we decided to adopt the MVs and exosome nomenclature.

The first study relative to EVs in the Central Nervous System (CNS) was performed on Drosophila about 20 years ago [16]. From there on, the work on EV role in the brain has dramatically increased, and EVs have been directly implicated in the modulation of neuronal development [17], as well as in the exchange of membrane proteins within the brain cells [18]. EVs have been proposed as markers for pathologies such as Traumatic Brain Injuries (TBI) and strokes, Alzheimer's disease, and generally for some types of cancers and CNS disorders [19–21]. Microglia cells, fundamental cells ensuring the immune response

in the CNS [22], have revealed to be both producers and recipient cells for EVs. While the role of microglia as recipient cells for EVs has been reviewed elsewhere [6], here we focus on microglia cells as producers of EVs, and in detail describe the correlation between their molecular composition and exerted biological functions. The state-of-the-art of the biogenesis mechanism of both small and large EVs from microglia has been recently exhaustively described by Aires et al. [23,24]. Notably, some processes in microglia EVs biogenesis are still unclear, but the use of methods for interfering with EVs secretion, either pharmacologically or genetically, will possibly allow to fill this gap [25]. An example of this approach is provided by Bianco et al., who modulated acid sphingomyelinase activity to block or trigger MVs release from microglia [26].

The molecular composition of microglia EVs

EVs secretion from microglia cells can occur in unstimulated conditions or can be enhanced by different stimuli (**Table 1**), which can be pro-inflammatory or pro-regenerative, but also not related to the M1 and M2 states (**Fig. 1**). The cellular models used to derive microglia-EVs are discussed elsewhere [22]. Microglia phenotype is highly affected by the microenvironment; accordingly, the EV molecular composition produced by resting or activated microglia is widely variable. The proteins, lipid, and miRNA present in the membrane or as cargo (**Fig. 2**) can vary in composition or abundance based on the state of donor cells [27,28] causing a wide spectrum of effects in recipient cells (**Table 2**).

PROTEIN COMPOSITION

The content of resting and stimulated microglia-derived EVs was investigated by Potolicchio et al., who reported an extensive proteomic analysis of the content of EV originated from the N9 murine cell line [29]. These vesicles include metabolic enzymes (e.g. glyceraldehyde 3-phosphate dehydrogenase, GAPDH, pyruvate kinase), chaperones, tetraspanins, and membrane receptors, very similar to the profile of exosomes originated by both B-cells and dendritic cells, together with specific microglia markers CD13 and MCT-1 [29]. Similar results were obtained using primary microglia extracted from SJL/J mice [29]. Moreover the identification of the lactate transporter on exosomes reveals their ability to deliver energy substrates to match the energy needs during increased neuronal activity, strengthening a model in which lactate flux may serve as support to neurons during synaptic activation [30].

ATP is a typical stimulus used to increase the secretion of EVs by cultured microglia [6] both in N9 cells and primary rat and mouse microglia [31,32]. The purinergic P2X7 receptor has been reported as the

major mechanism underlying the ATP-mediated EV shedding and secretion in N9 cells. It has been hypothesized that EV loaded with functional P2X7 can be a way for the cells to downregulate their plasma membrane pool, thereby reducing P2X7-mediated apoptosis [28]. This observation also suggests that EVs may serve as a carrier to transfer functional receptors out of the cell membrane modifying the responsiveness of cells to specific extracellular stimuli.

ATP stimulation is capable of modifying the content of EVs released from rat primary microglia, enriching them in proteins implicated in cell adhesion/extracellular matrix organization, metabolism, and autophagolysosomal pathway (**Table 1**) [33]. The increase of neuronal and microglial autophagy has been correlated with synaptic pruning [34], underling a possible correlation with the ATP-mediated release of EVs. Interestingly, in EVs released upon ATP stimulation, Drago et al. reveal the presence of the complement protein C1q involved in the pruning process [35], suggesting the role of microglia-derived EVs as carriers to tag aberrant synapses. ATP stimulus also increases metabolism-related proteins and enzymes, which are not present in constitutive EVs. Among these, there are enzymes of glycolysis, lactate production, pentose phosphate pathway, glutamine metabolism, and fatty acid synthesis (**Table 1**) [33]. All these proteins reflect the metabolic changes occurring after ATP stimulation that promote the generation of the fatty acid pool needed for microglia scanning process and phagocytic activity.

Capsaicin has been reported as another activation stimulus by Marrone et al. for microglia-derived shedding [36]. They found that the activation of the Transient Receptor Potential Vanilloid Type 1 (TRPV1) receptor by capsaicin on C57BL/6-derived murine microglial cells increases the production of MVs similarly to ATP. These MVs bear the capability of inducing miniature excitatory postsynaptic currents (mEPSCs) in neurons, confirming their role in the modulation of the synaptic activity of neurons [37].

Notably, the inflammatory status of microglia cells can affect the responsiveness to extracellular ATP. Takenouchi et al. describe how ATP may induce the release of EV enriched with GAPDH from MG6 mouse cells only under inflammatory conditions [32]. Extracellular GAPDH is involved in the regulation of neuritogenesis, and these results further support the role that microglia-EVs can play in the modulation of neuronal development.

Inflammation mimicked by lipopolysaccharides (LPS) treatment has demonstrated the importance of EVs in the CNS as evidenced by Asai et al. [38]. They have demonstrated that the ATP stimulation of primary cultured murine microglia, but not neurons or astrocytes, led to secretion of Tau in the exosome

fraction, albeit not phosphorylated. However, Tau is then transmitted from exosomes to cultured neurons where it is phosphorylated. As phosphorylated Tau is a pathological hallmark of different neurodegenerative pathologies [39], this work confirms the importance of microglia exosomes in the propagation of Tau pathology.

Using α -synuclein as a stimulus in EV production from BV2 cells, Chang et al. have observed an increased secretion of EVs bearing a high level of mTNF- α (EVs-T) and a higher surface amount of MCH-II receptor, with respect to EVs obtained by untreated microglia. The EVs-T are able to increase apoptosis in rat cortical neurons [40] and this effect can be counteracted by mTNF- α neutralizing antibodies supporting the active role of EVs-T in this process. Interestingly, only EVs can produce this specific effect on recipient cells, in fact, a concentration of soluble TNF- α equal to that contained in the α -synuclein-induced EVs does not show the same apoptotic effects on cortical neurons.

Recently, it has also been reported that α -synuclein can not only be a microglia stimulus to produce EV, but also a cargo [41]. Specifically, plasma exosomes derived by PD patients are internalized by BV2 cells where they caused cell autophagy dysregulation, leading to an increased accumulation of intracellular α -synuclein and its accelerated secretion into the extracellular medium, as exosome cargo. These data strengthen the role played by microglia in neuroinflammatory conditions through the release of EVs, carrying pathological proteins and inflammatory mediators.

Neuroinflammation is the primary response of microglia not only in case of infection but also upon TBI in which it can mediate neuronal damage [42]. Kumar et al. analyzed microparticles (MPs) extracted from the blood of C57BL/6 mice after surgical TBI, and showed that TBI increases the release of microglia-derived MPs. TBI-derived MPs can upregulate pro-inflammatory signaling in recipient cells. The same authors also evaluate the effects of LPS in the modulation of microglia-derived EV production. They show how the presence of IL-1 β and miR-155 is highly increased in LPS-derived EVs and sufficient to activate resting microglia and propagate the pro-inflammatory response [42–44]. In fact, the treatment of primary and immortalized microglia (BV2) cells with these MPs increases the expression of IL-1 β , TNF- α , CCL2, IL-6, NOS2, and miR-155. MPs extracted both from BV2 cells upon LPS treatment and from mice primary microglia isolated *ex vivo*, following TBI, can initiate neuroinflammation upon intracortical injection in naïve animals, confirming the pro-inflammatory capability of TBI-derived MPs [6,42,43].

The ischemic injury induced in rat primary microglia culture (Oxygen-Glucose Deprivation, OGD model) is also able to promote the release of exosomes carrying Nicotinamide Phosphoribosyltransferase

(NAMPT), a key enzyme of the nicotinamide adenine dinucleotide synthesis pathway which acts as a pro-inflammatory agent in the CNS [45]. The release of exosomes seems to be mediated by the increase of extracellular ATP in response to the ischemic injury.

It has been recently demonstrated that the strength of the inflammatory response mediated by microglia depends on the tissue microenvironment. Murgoci et al. report an extensive proteomic analysis on exosomes extracted from neonatal rat microglial cells derived from two different CNS locations, the Spinal Cord Microglia (SpC-M) and Cortex Microglia (Cx-M), in resting condition and upon stimulation with LPS. The results demonstrate that microglia functions and their inflammatory response depend on their origin and these characteristics are preserved if microglia cells are maintained in vitro [46]. Cx-M and SpC-M derived exosomes displayed some of the typical EVs proteins (including tetraspanin CD81, transmembrane protein Anxa2, cytosolic protein S100, C1q) and proteins related to development, (such as axonal growth promoters, neuroprotective factors, promoters of neurites outgrowth), and have been implicated in neurodevelopment (Table 1). By contrast, only Cx-M derived exosomes revealed an enrichment in neurite outgrowth, nerve regeneration and axonogenesis, whereas only SpC-M-derived exosomes showed an enrichment of inflammation and injury categories. Specifically, in control and LPSinduced SpC-M exosomes, several proteins involved in inflammatory processes have been identified. LPS-treated Cx-M exosomes present proteins involved in chemotaxis, NFkB pathway, and metabolic processes. Finally, exosomes purified from LPS-treated microglia revealed the presence of proinflammatory chemokines, as well as the presence of IL-6 and TNF- α [46]. Interestingly, both sources of microglia exosomes enhance the neurite outgrowth of Dorsal Root Ganglion Cells (DRGs). Conversely, only SpC-derived exosomes under LPS stimulation significantly attenuated glioma proliferation supporting the idea that the origin of microglia can ensure different biological functions [46].

Glebov et al. have provided first evidence of neurotransmitter-induced communication between neurons and microglia via released exosomes [47]. Serotonin (5-HT) can induce exosome release from BV2 microglial cells, by stimulation of 5-HTRs. The stimulation of 5-HTRs causes an increase of intracellular Ca^{2+} via PLC phosphorylation that seems to favor the fusion of MVBs with the plasma membrane and consequently increases the secretion of EVs. The vesicles have shown an increased concentration of Insulin Degrading Enzyme (IDE), flotillin-1, and actin. Glebov et al. also suggest an active communication of serotoninergic neurons with microglia mediated by the EVs; however, this suggestion is not supported by a direct evaluation.

Another peculiar stimulus promoting the EV release is the Wnt3a protein. Wnt3a is a morphogenic protein of the WNT family, which is emerging to play diverse roles in the adult CNS [48,49]. Hooper et al. study the effects of recombinant Wnt3a administration to primary rat microglial cells on EVs production and composition [50]. Wnt3a does not increase EVs secretion; however, a proteomic analysis reveals that Wnt3-induced EVs are enriched in proteins associated with cellular metabolism, architecture, and protein synthesis [50]. Wnt3a-induced EVs also contain Wnt3a, ubiquitin, and proteasome components such as proteasome subunit beta type-7 and alpha type-2 (**Table 1**), suggesting a possible role in the extracellular space clearance.

The leech is also a model to study microglia [22], and proteomic analysis of its microglia-derived EVs have been found to contain components involved in neuronal development, neurite growth, axon guidance, filopodium assembly, and dendrite development, all effects confirmed by *in vitro* administration to recipient neurons [51,52].

LIPID COMPOSITION

Hundreds of lipids populate the plasma membrane [53]. The lipid environment in the membrane is related to its protein content [54], and lipids can regulate receptor-dependent cell signaling [55,56]. Accordingly, a functional role for lipids is expected also for EVs. Indeed, the interest in lipid research on EVs is growing very rapidly [57]. Most of the lipidomic studies agree in identifying Cholesterol (CHOL), Sphingomyelin (SM), glycosphingolipids, and Phosphatidylserine (PS) as the most recurrent lipids in EVs [53]. The lipid composition is reported to be highly variable among different vesicles, with the highest content of CHOL and SM observed in exosomes rather than MVs [58]. It has to be noted though that most of the studies on EV lipid composition do not include microglia-derived EVs [59]. However, Osteikoetxea et al. described the protein/lipid ratio of exosomes released from resting BV2 cells [60], suggesting a similarity between microglia derived-EVs and EVs derived from other types of cells.

The relatively few studies performed on microglia-derived EVs stand out for providing compelling evidence of the functional role of EVs lipids. A striking example is the recent observation that the lipid fraction of microglia-derived EVs can drive oligodendrocytes precursor cell (OPC) maturation [61], almost to the same extent as the same EVs enriched with miR-219, a crucial component of promyelinating exosomes [61,62]. The authors have characterized the lipid content of EVs derived from primary rat microglia stimulated with ATP. They have used non-polarized microglia (NS), or microglia polarized by MSC co-culture (MSC-EVs), a Th1 cocktail of inflammatory cytokines (i-EVs), or IL-4, a pro-regenerative cytokine (IL-4-EVs). They detect the presence of sphingosine-1-phosphate (S1P) that crucially acts as an attractive guidance cue for OPC migration to the myelin lesion site *in vivo* due to S1P receptor expression on the OPC plasma membrane [61]. However, while MSC-EVs and IL-4-EVs can also promote OPC maturation and initiate remyelination, i-EVs stop OPC maturation [61]. Co-culture experiments revealed that the negative effect of i-EVs is mediated by the detrimental action of protein and/or RNA cargoes onto astrocytes, which are converted into the A1 harmful phenotype [63]. In another work, Prada et al. show that PS of MVs derived from microglia treated with ATP is involved in the recognition and contact with different recipient cells [64]. This is in line with the PS role previously reported for the targeting of monocyte/macrophage-derived MVs onto platelets [65]. However, they also found that MVs contact microglia and astrocytes in very different ways. MVs on microglia display a net drifted movement on the plasma membrane, possibly driven by cortical actin flow, before internalization. Instead, on astrocytes, they are static and almost do not internalize. Finally, they show that cloaking of MVs-related PS by annexin-V reduces MVs adhesion only in part, suggesting that other surface molecules likely contribute to their interaction with glial cells.

Microglia-derived EVs can also crucially influence the excitation/inhibition balance in the adult CNS. The endocannabinoid N-Arachidonoylethanolamine (AEA), a lipid crucially involved in the modulation of synaptic transmission, is enriched in EVs derived from ATP-stimulated N9 cells and primary rat microglia [66]. AEA is mainly present in the MVs sample, which, administrated to cultured hippocampal neurons, can modulate GABAergic transmission, due to the activation of presynaptic CB1. The same group has also shown that microglia-derived EVs positively affect spontaneous excitatory neurotransmission *in vitro* and *in vivo* by promoting neuronal production of ceramide and sphingosine [37]. These data may suggest that EVs are good vehicles of bioactive hydrophobic compounds like endocannabinoids, whose transport across the CNS would be otherwise difficult to explain [66].

MIRNA COMPOSITION

miRNAs are small non-coding RNAs, which play a crucial role in the post-transcriptional regulation of gene expression [67]. Their packaging into EVs has been widely recognized as a way to protect them from degradation and potentiate their response [68]. The observation that the highest expression of tissue-specific miRNAs is found in the brain [69] has motivated several studies about the role of EVs-associated miRNA in brain pathologies [70], as well as an in-depth characterization of the miRNA content of microglia-derived EVs [71,72].

The miRNA content of microglia-derived EVs was analyzed by RNA-seq by Lemaire et al. in resting leech primary microglia, finding an enrichment of 6 miRNAs: miR-1860, miR-1705, miR-2284y-6, miR-146a, miR-858, miR-7718. Most of the selected miRNA show at least 2-fold increased expression in microglia parental cells than in recipient neurons, qualifying them also as a possible signature for microglia-derived EVs [51]. Noteworthy, some changes caused by the administration of these EVs to recipient neurons can be ascribed to these miRNAs.

This work has partially confirmed Prada et al. study in which an extensive miRNA profiling of MVs, exosomes, and parental cells from ATP-stimulated primary rat microglia was performed [71]. 21 miRNAs have been found differentially abundant in M1-EVs versus M2-EVs. miR-146a-5p, miR-181a, and miR-223, all displaying conserved synaptic targets, are overexpressed in EVs released from M1 microglia, as well as in the Cerebro Spinal Fluid (CSF) of patients with multiple sclerosis. miR-146a-5p carried by microglia-derived EVs downregulated synaptic Syt1 and Nlg1 protein levels in recipient neurons, leading to loss of dendritic spines and synaptic density in vitro and in vivo [71]. By this mechanism, inflamed microglia can selectively transfer miRNA to neurons via MVs, directly affecting synaptic transmission. The miRNA signature of microglia-derived EVs has also been measured in N9 cells polarized with LPS and then stimulated with ATP. miR-146a was confirmed to be part of M1-EVs, together with miR-155 and, in line with their increase in the parental cells after the M0-M1 transition. EVs from M1-N9 cells also showed a reduction of miR-124, another characteristic typical of this transition [73], further confirming that the composition and biological effect of EVs strictly depends on the condition of the parental cell. Tian et al. analyzed miRNAs in EVs derived from BV2 cells polarized with LPS to induce the M1 shift and with IL-4 to induce the M2 one. They identified miR-26a as a distinguishing miRNA upregulated in M2 microglia-derived exosomes. As miR-26a was previously shown to promote angiogenesis in glioma [74], and miR-26a loaded EVs can promote angiogenesis in vitro and in vivo, they claim that exosomes released by IL-4-polarized microglia may have therapeutic potential in the treatment of stroke [75].

Interestingly, EVs secreted by LPS-treated primary rat microglia can have anticancer effects, on rat glioma. The molecular mechanism by which this occurs is unknown but it has been proposed to derive from the miRNA cargo of EVs, in particular the family of miR-16 [76].

Fernandes et al. used SH-SY5Y cells transfected with the Swedish mutant of APP695 (SH_{Swe}) as a source of APP- and A β -enriched exosomes and they administrated them to human microglia HMC3 cells [77]. While SH_{Swe} exosomes are enriched in miR-155, miR-146a, miR-124, miR-21, and miR-125b, exosomes

in turn released by HMC3 cells contain just one upregulated miRNA, miR-21, one of the most common regulators of the inflammatory response. This work thus suggests an important role for miR-21 in neuron-microglia communication during AD pathogenesis [78].

There is a wide interest in understanding the role of EVs-associated miRNAs in several pathologies of the brain such as TBI [79], neuroinflammation and depression [80], and neurodegeneration [70,81]. Understanding whether these miRNAs represent a pathology biomarker or also a target for therapeutic intervention will certainly constitute an exciting avenue for future investigations.

Conclusion

The composition of EVs can vary greatly depending on both the stimulus received from the parental cells and the cell model used making the overall picture stimulating but also puzzling. It is worth noticing that, even though there are a lot of studies regarding the biological effect of EVs in the CNS, including their interaction with tumors and neurodegenerative diseases, the majority of them still consider microglial cells as recipient cells for EVs produced by other components of the CNS [82–84]. Considering the role of microglia and its nature of restless immune cells, it will be important to identify all the possible effects of these cells on neighboring cells mediated by EVs.

Recently, EVs have been extensively manipulated and engineered by different approaches such as passive or active drug loading, active targeting, and pH-responsive EVs. These can be focused on modifying both the secreting cells (endogenous engineering) and the EVs after isolation (exogenous engineering). EVs can be produced with artificial features aimed to improve their translational potential. However, the lack of knowledge on the mechanisms of action and the specific biological effects of microglia-derived EVs in different pathological conditions has so far hindered their development and use in this field. Understanding deeper the functional roles of these structures would pave the way for the development of therapies that include microglia-EVs, as already done and consolidated for mesenchymal stem cells [85–87].

Perspective

1. EVs are a pivotal method of cell-to-cell communication in both physiological and pathological conditions. Although accumulating evidence demonstrates the crucial role of microglia-derived

EVs in the CNS, the molecular composition and the mechanisms underlying their release have not been fully understood yet.

- 2. Thanks to omics approaches, this field is accumulating knowledge about the EVs composition, but there is a lack of a unifying vision allowing the correlation of a precise EV component with one or more biological effects.
- 3. Most of the data in the field mostly derive from animal models, and the interspecies differences call for the need to look for reliable human models. Further studies will be necessary to deepen the composition-function relationship of human EVs and understand if they can be exploited for therapeutic applications.

Abbreviations

5-HT	Serotonin
5-HTRs	5-HT Receptors
AEA	N-Arachidonoylethanolamine
CHOL	Cholesterol
CNS	Central Nervous System
CSF	Cerebro Spinal Fluid
Cx-M	Cortex Microglia
DRGs	Dorsal Root Ganglion Cells
EVs	Extracellular Vesicles
IDE	Insulin Degrading Enzyme
ISEV	International Society of Extracellular Vesicles
LPS	Lipopolysaccharides
MCHII	Major Histocompatibility Complex Class II
mEPSCs	Miniature Excitatory Postsynaptic Currents
miRNA	MicroRNA
MISEV	Minimal Information for Studies of Extracellular Vesicles
MPs	Microparticles
MVB	Multivesicular Body
MVs	Microvesicles
NAMPT	Nicotinamide Phosphoribosyltransferase
OGDNAMPT	Oxygen-Glucose Deprivation Nicotinamide phosphoribosyltransferase
OPCOGD	Oligodendrocytes Precursor CellOxygen-Glucose Deprivation
PDOPC	Parkinson's DiseaseOligodendrocytes Precursor Cell
PSPD	PhosphatidylserineParkinson's Disease
S1PPS	Sphingosine-1-PhosphatePhosphatidylserine
SMS1P	SphingomyelinSphingosine-1-Phosphate
SpC-MSM	Spinal Cord MicrogliaSphingomyelin
TBISpC-M	Traumatic Brain InjuriesSpinal Cord Microglia
TRPV1TBI	Transient Receptor Potential Vanilloid Type 1 Traumatic Brain Injuries
TRPV1	Transient Receptor Potential Vanilloid Type 1

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Competing Interests

The authors declare that there are no competing interests associated with the manuscript.

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Author Contribution

L.C. and C.G. conceived the area of focus for this review. L.C., C.G. and L.M. wrote the main text. C.M. and L.M. edited.

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Tables and Figures

Model	Stimulus	EVs Component	
BV2	α-Syn [40]	MHCII, TNF-α [40]	
	LPS [42,50]	IL1-β [42]	
		TNF-α, IL1-β [50]	
	Serotonin [47]	IDE, Flotilin-1, Actin [47]	
	Wnt3 [50]	Actin, peroxidasin homolog, 40S ribosomal, ubiquitin, proteasome su	
		units [50]	
	IL-4 [75,88]	CD63, CD9, TSG101, miR-124 [88]	
		miR-466, miR-6360, miR-26a, miR-1940 [75]	
N9	ATP [31,89]	AEA, TSG101, flotilin-1 [89]	
		IL1-β, P2X7 [31]	
MG6	ATP [32]	GAPDH, IL1-β [32]	
	LPS [32]	GAPDH [32]	
Primary Rat Microglia	ATP [31,33]	GAPDH, IL1-β [31]	
		Glucose-6-phosphate isomerase, Gpi, Lactate dehydrogenase A, LDHA,	
		Malate dehydrogenase 2, Mdh2, Tranketolase, Glucose-6-phosphate 1-	
		dehydrogenase, G6PD, Glutamate dehydrogenase 1, Acetyl-CoA	
		carboxylase-β, Acacβ [33]	
	LPS [46]	Cfh, mrc2, C4a, Lgal3bp, Lgals1, C1S, Cfhr1	
		MCT-1, CD63, CD13, GAPDH, pyruvate kinase, CXCL1, CXCL2,	
		CXCL3, CCl2, CCL7, IL-6, TNF-α [46]	
	TBI [42,61]	G6PD, ACACβ [61]	
		P2Y12, CD45 [42]	
	Th1 Cytokines	C1q, TNF-α [61]	
	cocktail [61,71]	miR-181a, miR-223, miR-146a [71]	
	Resting [61]	Spp1, Spon1, Sned1, Srpx, Hspg2, Fstl1, Olfml3	
		Sphingomyelin, lactosylceramide, ceramide, GM3, glucosylceramide,	
		S1P [61]	
Primary Mouse Microglia	ATP [42]	miR-155, miR-146, P2Y12, CD45 [42]	
	LPS+ATP [38]	TSG101, MHCII, hTau [38]	
	TBI [32]	GAPDH [32]	
Primary Leech Microglia	Resting [51]	miR-1860, miR-1705, miR-2284y-6, miR-146a, miR-858, miR-7718 [51]	

Table 1 Main components of microglia-derived EVs, divided by cell model and production stimulus.

Table 2 Effects of EVs on recipient cells. The parental cell model, the production stimulus, the effect raised and the recipient cells are reported.

Parental	Stimulus	Effect on parental cells	Pathways influenced	Recipient Cells
Cells				
BV2	α-Syn	Inflammation [40]	Neurodegeneration,	Rat cortical neurons
			Neurotoxicity [40]	[40]
	LPS	Inflammation [42]	Neuroinflammation [42]	Primary mouse and
			Propagation of inflammation	immortalized rat
			Neurodegeneration [50]	microglia [42,50]
	Serotonin	Increased EVs	n/d	n/d
		production [47]		
	Wnt3	n/d	Cellular metabolism,	n/d
			maintaining of cellular	
			architecture, protein synthesis	
			and degradation,	
			neuroprotection and	
			extracellular space clearance	
			[50]	
	IL-4	M2-Shift [88]	Modulation of neuroprotection,	Rat neurons [88]
			upregulation of miR-124 [88]	
BV2	Resting	M2-Shift [90]	Anti-inflammatory pathways	in vitro mouse
expressing	6		[90]	microglia, in vivo
IL-4			L J	mouse phagocytes
				[90]
N9	ATP	Activation [31]	Neuroinflammation [31]	n/d
MG6	LPS	Inflammation [32]	n/d	n/d
Primary	ATP	Activation [31,33]	Cell adhesion, extracellular	Primary rat
Rat			matrix organization,	astrocytes and
microglia			autophagolysosomal pathways,	microglia [33]
			cellular metabolism [33]	
			Neuroinflammation [31]	
	Resting	n/d	Neurogenesis, proliferation and	Rat neurons [37,66]
			growth [46]	
			mEPSC induction, Sphingolipid	
			metabolism [37]	
			CB1 activation, mIPSC	
			inhibition [66]	
	LPS	n/d	Inflammation and immune	n/d
			response [46]	
	Polarization by MSC	Pro-regenerative shift	OPC maturation, migration and	OPC and DRG
	co-culture and by IL-4	[61]	remyelination [61]	neurons [61]
Primary	TBI	Inflammatory response	Inflammatory signaling [42]	Primary mouse and
Mouse		[42]		immortalized rat
Microglia				microglia [42]

	Capsaicin	Increased MVs	Induction of mEPSC in neurons	Mouse Neurons
		production [36]	[36]	[36]
	IL4/ATP	M2-Shift [91]	Inhibition of cell senescence	in vivo mouse
			[91]	myeloid cells [91]
Primary	Resting	n/d	Neuroprotection and neurite	n/d
Leech			development [51]	
microglia				

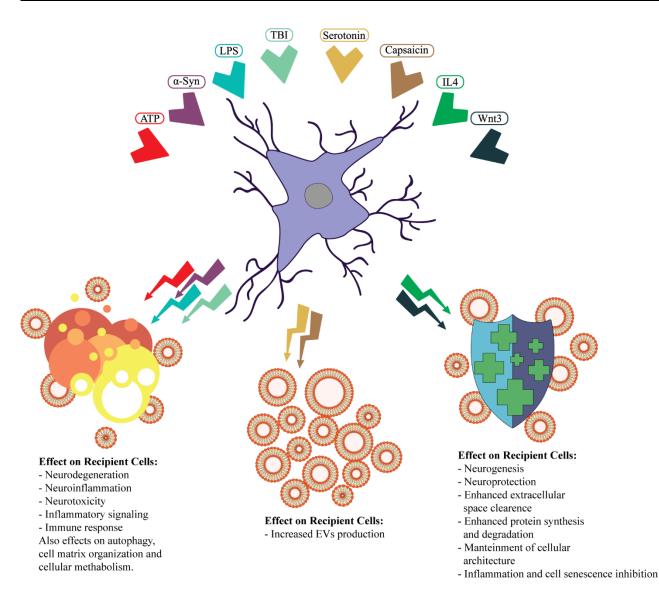


Figure 1 Schematic representation of the stimuli used to promote EV production by microglial cells. Stimuli are color-coded; arrows at the bottom in the same color-code indicate the main biological effect raised by each stimulated EVs in the CNS.

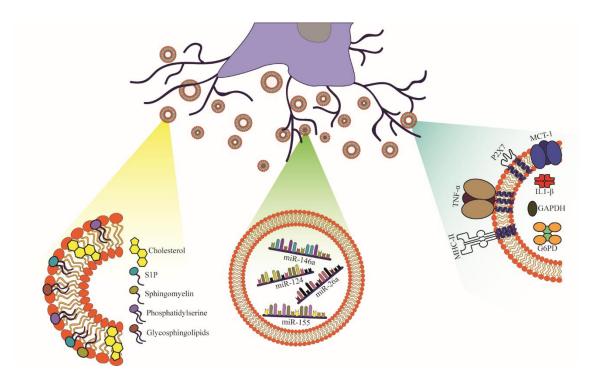


Figure 2 Main components of microglia-derived EVs. Molecules are grouped in Lipids (left), miRNA (center) and proteins (right). The most representative member of each category has been reported.