

3 **Rab8a/SNARE complex activation**
4 **promotes vesicle anchoring and transport**
5 **in spinal astrocytes to drive neuropathic**
6 **pain**

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23 **ABSTRACT**

24 Neuropathic pain (NPP) remains a clinically challenging condition, driven by the activation
25 of spinal astrocytes and the complex release of inflammatory mediators. This study aimed to
26 examine the roles of Rab8a and SNARE complex proteins in activated astrocytes to uncover
27 the underlying mechanisms of NPP. The research was conducted using a rat model with
28 chronic constriction injury (CCI) of the sciatic nerve and primary astrocytes treated with
29 lipopolysaccharide. Enhanced expression of Rab8a was noted specifically in spinal dorsal
30 horn astrocytes through immunofluorescence. Electron microscopy observations showed
31 increased vesicular transport and exocytic activity in activated astrocytes, which was
32 corroborated by elevated levels of inflammatory cytokines such as IL-1 β and TNF- α detected
33 through quantitative PCR. Western blot analyses confirmed significant upregulation of Rab8a,
34 VAMP2, and Syntaxin16 in these cells. Furthermore, the application of botulinum neurotoxin
35 type A (BONT/A) reduced the levels of vesicle transport-associated proteins, inhibiting
36 vesicular transport in activated astrocytes. These findings suggest that the Rab8a/SNARE
37 pathway in astrocytes enhances vesicle transport and anchoring, increasing the secretion of
38 bioactive molecules that may play a crucial role in the pathophysiology of NPP. Inhibiting
39 this pathway with BONT/A offers a novel therapeutic target for managing NPP, highlighting
40 its potential utility in clinical interventions.

41 **KEYWORDS:** Neuropathic pain, astrocytes, Rab8a, vesicular transport, SNARE proteins

42

43 INTRODUCTION

44 Neuropathic pain (NPP) represents a global therapeutic challenge characterized by complex
45 pathophysiological mechanisms and a lack of effective clinical analgesics [1, 2, 3, 4, 5]. The
46 functional specificity of cortical networks and their projection targets in the pain process
47 occurs at least on four interconnected levels: dynamic activity states within the cortical
48 network; functionally distinct subdomains; specific circuit connections that distinguish pain
49 from other functions; and co-active cell assemblies [6]. Among these, intercellular
50 communication and molecular signaling pathways within specific circuit connections play a
51 pivotal role in the sensitization and regulation of nociceptive pathways in the sensory nervous
52 system and the pathological process of NPP [7, 8, 9, 10].

53 Astrocytes, distinguishable by their expression of glial fibrillary acidic protein (GFAP) across
54 all major branches and processes, dynamically modulate in response to injury through gap
55 junction protein complexes that physically couple adjacent cells, allowing free exchange of
56 ions and cytoplasmic components [11]. Inhibition of astrocyte activation can significantly
57 alleviate pain caused by peripheral nerve damage in the early stages of NPP [12, 13, 14].

58 Astrocytes mediate intercellular communication within the nervous system through the
59 production and secretion of neuroactive substances [15, 16, 17]. Injury signals drive
60 phenotypic transformation of astrocytes and the release of inflammatory mediators, playing
61 roles in central and peripheral sensitization and participating in the progression of NPP. An
62 important characteristic of their activation is the increased release of bioactive molecules
63 such as inflammatory factors, ATP, and glutamate [18, 19, 20, 21, 22, 23, 24].

64 Furthermore, astrocytes contain vesicles that store and release bioactive molecules in an
65 activity-dependent manner, a principal mechanism in the pathophysiology of
66 neurodegenerative diseases [25, 26, 27, 28, 29, 30, 31]. However, the specific mechanisms by
67 which astrocytes in NPP increase the secretion of bioactive molecules remain unclear [32],

68 complicating the identification of targets for intervention.
69 Rab proteins, acting as molecular switches in vesicle transport, interact with upstream
70 regulators and downstream effectors, playing a critical role in vesicle movement, docking,
71 and fusion [33, 34]. In their active GTP-bound form, Rab proteins activate downstream
72 effector proteins, recruit cytoplasmic adhesion factors, and regulate vesicle dynamics [35, 36,
73 37, 38, 39, 40, 41, 42]. The fusion of vesicles with the cell membrane also relies on a set of
74 transmembrane proteins known as the SNARE complex, which provides the molecular basis
75 for directed vesicle transport, targeting, docking, and membrane fusion [43, 44, 45]. Currently,
76 the role of Rab8a in vesicle release processes in spinal astrocytes has not been reported. Thus,
77 this study aims to examine the modification of Rab8a in activated astrocytes using a rat
78 model with sciatic nerve ligation and lipopolysaccharide (LPS)-treated primary astrocytes to
79 investigate its role in SNARE complex formation and vesicle transport and to explore the
80 impact of the Rab8a/SNARE signaling pathway on NPP and its mechanisms. By revealing
81 the role of this signaling pathway in regulating astrocyte vesicle transport and secretion
82 functions, we aim to provide a new perspective on the molecular mechanisms of NPP and lay
83 the groundwork for developing targeted therapeutic strategies, which hold significant
84 scientific and clinical relevance.

85

86 **MATERIALS AND METHODS**

87 **Experimental animals**

88 Male Sprague-Dawley (SD) rats, aged 7-8 weeks (200-230 g), were obtained from the
89 Experimental Animal Research Institute of the Army Medical University. These rats were
90 housed in a controlled environment at 25°C with a 12-hour light/dark cycle, with free access
91 to food and water. The animal experimental processes were approved by the Ethnic
92 Committee of The First Affiliated Hospital of Chongqing Medical University

93 (AMUWEC20210719) and conducted in strict accordance with the standard of the Guide for
94 the Care and Use of Laboratory Animals published by the Ministry of Science and
95 Technology of the People's Republic of China in 2006.

96

97 **Induction of NPP through chronic constriction injury (CCI)**

98 Ten SD rats (aged 7-8 weeks, weighing 200-230 g) were utilized. The sample size calculation
99 was based on setting the range of acceptable degrees of freedom (DF) for ANOVA analysis
100 between 10 and 20. Let N represent the total number of subjects, k the number of groups, and
101 n the number of subjects per group, calculated as $n = DF/k + 1$. Hence, the minimum total
102 sample size N(min) was determined to be 6, and the maximum total sample size N(max) was
103 11 [46]. The ten rats were randomly divided into two groups: a normal group (control group,
104 n=5) and a CCI group (ligation group, n=5). Each group underwent specific procedures: the
105 normal group received a sham operation without ligation; the CCI group was subjected to a
106 procedure established in previous studies [47]. Briefly, a blunt dissection was performed in
107 the biceps femoris, exposing the common sciatic nerve at the mid-thigh level. Approximately
108 1 cm of the nerve was freed from surrounding connective tissue near its trifurcation, and three
109 loops of 4.0 non-absorbable surgical suture (Shanghai Fosun) were loosely tied around it at 1
110 mm intervals. Under 30x magnification, these ties did not significantly compress the nerve's
111 diameter but did induce slight and transient twitches in the muscles innervated by the sciatic
112 nerve. The test animals were subsequently maintained for 14 days.

113

114 **Astrocyte culture**

115 Primary astrocytes were prepared from one-day-old SD rats, following the procedure
116 described by Sebastian Schildge et al. [48]. These cells were isolated from the cerebral cortex
117 and subsequently cultured in 25 cm² flasks pre-coated with 50 µg/mL poly-D-lysine. The

118 culture medium used was DMEM (Gibco, New York, USA) supplemented with 10%
119 heat-inactivated fetal bovine serum and 1% penicillin-streptomycin (Beyotime, Shanghai,
120 China). The cultures were maintained under conditions of 5% CO₂ at 37°C. The medium was
121 replaced the day following the initial culture and thereafter every two days. On the seventh
122 day, the cultures were placed on a rotating shaker at 37°C for 6 hours (240 rpm) to detach
123 microglial and oligodendrocyte precursor cells. Following this, the medium was discarded,
124 and the astrocytes were cultured at a final density of 1.2×10^6 cells per well in 6-well plates
125 and 4×10^4 cells per well in 96-well plates for subsequent cell counting kit-8 (CCK8) assay.
126 In the drug treatment groups, astrocytes were co-incubated with LPS (100 ng/mL) and
127 botulinum toxin A (BONT/A) (0.1 U/mL) for one hour [49, 50, 51, 52]. In this study, LPS
128 was used as a cell activator and BONT/A as a vesicular secretion inhibitor.
129 Immunofluorescence (IF) staining with GFAP (an astrocyte marker, BM-0055, Bioss, Wuhan,
130 China) was performed to identify the astrocytes. A high-purity population of astrocytes (over
131 95% GFAP-positive) was obtained [53]. To ensure cell culture quality, high-quality fetal
132 bovine serum and culture medium, along with sterile plastic products designed specifically
133 for tissue culture, were used. To prevent microbial contamination, 100 U/mL
134 penicillin-streptomycin (Bi Yun Tian, C0222) was employed to protect against cellular
135 contamination. Mycoplasma testing was performed prior to experiments to exclude
136 mycoplasma infections.

137

138 **Cell viability assay**

139 Cell viability was assessed using CCK-8 (Bioss, Beijing, China). Astrocytes were cultured in
140 96-well plates for 24 hours. Following treatment with LPS (100 ng/mL) for 24 hours, CCK-8
141 solution was added to each well and incubated at 37°C for 2 hours. Absorbance was
142 measured at 450 nm using a microplate reader.

143 **Immunohistochemistry (IHC)**

144 Spinal cord tissues from CCI rats were collected on day 14 post-sciatic nerve ligation. Rats
145 were deeply anesthetized with isoflurane (2-2.5%, airflow 500-700 mL/min) and then
146 perfused intracardially with 4% paraformaldehyde (Sigma) pre-cooled to 4°C. The spinal
147 cord was quickly removed and immersed in 4% paraformaldehyde at 4°C overnight. After
148 fixation, the spinal cord was dehydrated, and the lumbar enlargement region was sectioned
149 into 16 µm thick slices. Endogenous peroxidase activity was blocked using 3% H₂O₂ for 20
150 minutes. Sections were incubated with 10% normal goat serum and anti-Rab8a antibody
151 (1:150; LifeSpan Biosciences) at 37°C for 1 hour, followed by overnight incubation at 4°C.
152 After PBS rinsing, sections were incubated at 37°C for 1 hour and visualized using an
153 enhanced nickel-DAB staining reagent for 5 minutes. IHC images were captured using a
154 microscope (Leica). Five random spinal cord sections were selected by two experienced
155 pathologists in a blinded manner, and the average optical density of all positively stained
156 astrocytes in the selected fields was measured and analyzed using Image-Pro Plus 6.0.

157

158 **Immunofluorescence (IF) staining**

159 For double-labeling IF experiments on spinal cord sections, prepared slices were treated with
160 3% H₂O₂ for 20 minutes to suppress endogenous peroxidase activity. The sections were then
161 incubated at 37°C for 1 hour, followed by overnight treatment at 4°C with 10% normal goat
162 serum. Subsequently, the slices were incubated with anti-GFAP monoclonal antibody (1:250,
163 Bioss) at 37°C for 1 hour, followed by co-incubation with anti-Rab8a polyclonal antibody
164 (1:150, LifeSpan Biosciences) at 37°C for 1 hour, and then overnight at 4°C.
165 FITC-conjugated goat anti-mouse antibody (1:500, Abcam, UK) and Cy3-conjugated goat
166 anti-rabbit antibody (1:600, Jackson ImmunoResearch, USA) were added and incubated at
167 37°C for 1 hour. Finally, nuclei were stained with 4',6-diamidino-2-phenylindole (DAPI)

168 (Sigma, USA) and analyzed by two experienced pathologists in a blinded manner using a
169 laser scanning confocal microscope (Olympus, Japan).

170 For IF staining experiments detecting GFAP in astrocytes, astrocytes grown on microscope
171 slides were fixed in 4% paraformaldehyde at 37°C for 30 minutes, followed by incubation in
172 5% BSA at 37°C for 1 hour and then overnight incubated with anti-GFAP monoclonal
173 antibody (1:250, Bioss) at 4°C. The cells were then incubated at 37°C for 1 hour with
174 FITC-conjugated goat anti-mouse antibody (1:500, Abcam, UK). Nuclei were visualized with
175 DAPI staining (Bioss, Beijing, China), and images were captured using a microscope (Leica).

176 In the double-labeling, IF experiments, astrocytes grown on microscope slides were fixed
177 with 4% paraformaldehyde at 37°C for 30 minutes, then incubated in 5% BSA at 37°C for 1
178 hour, followed by overnight incubation with anti-Rab8a polyclonal antibody (1:250, LifeSpan
179 Biosciences) at 4°C. FITC-conjugated goat anti-mouse antibody (1:500, Abcam, UK) and
180 Cy3-conjugated goat anti-rabbit antibody (1:600, Jackson ImmunoResearch, USA) were
181 added and incubated at 37°C for 1 hour. Nuclei were stained with DAPI (Cat# D9542-5MG,
182 Sigma, USA) and analyzed using a laser scanning confocal microscope (Leica).

183

184 **Western blot assay**

185 Cell lysates were collected from primary astrocyte cultures in RIPA buffer containing a
186 protease inhibitor cocktail for Western blot analysis 1-hour post-LPS stimulation. The
187 reaction mixtures were centrifuged at $12,000 \times g$ for 15 minutes at 4°C. Samples containing 2
188 μg of protein were heated at 100°C for 5 minutes in a loading buffer (5x Loading Buffer,
189 Beyotime, Shanghai, China). Separation was conducted using polyacrylamide gels (10-12.5%,
190 Epizyme, Beijing, China). Following membrane transfer, the membranes were incubated
191 overnight at 4°C with anti-GFAP monoclonal antibody (1:1000, Bioss), anti-Rab8a
192 polyclonal antibody (1:1000, LifeSpan), anti-VAMP2 polyclonal antibody (1:1000, Cell

193 Signaling), anti-Syntaxin16 polyclonal antibody (1:1000, Cell Signaling), and anti- β -actin
194 (1:1000, Proteintech). The membranes were then incubated for 1 hour with horseradish
195 peroxidase-conjugated secondary antibodies and visualized using ECL solution (Biosharp,
196 Shanghai, China). Immunocomplexes were detected using the Bio-Rad system, and relative
197 immunoreactivity levels were quantified using Image Lab software.

198

199 **Quantitative real-time polymerase chain reaction (qPCR)**

200 Total RNA from astrocytes was isolated using the RNAeasy™ animal RNA isolation kit with
201 the spin column, following the manufacturer's instructions (Beyotime, Shanghai, China).
202 RNA sample transcription was repeated using the PrimeScript™ RT reagent kit with gDNA
203 Eraser (Takara, Japan), according to the manufacturer's instructions. Real-time qPCR was
204 conducted using SYBR Premix Ex Taq II (Takara). The thermal cycling program included a
205 10-minute pre-incubation at 95°C, followed by 45 cycles of 10 seconds at 95°C, 30 seconds
206 at 60°C, and 60 seconds at 65°C. The specificity of the PCR products was verified through
207 melt curve analysis.

208

209 **Electron microscopy (EM)**

210 Astrocytes were co-incubated with LPS (100 ng/mL) or LPS (100 ng/mL) and BONT/A (0.1
211 U/mL) for 24 hours. Cells were then detached using a 0.025% trypsin-EDTA solution and
212 fixed with 2.5% glutaraldehyde at 4°C for 12 hours. The prepared cells were further fixed
213 with 1% osmium tetroxide at 4°C for 1 hour. After gradient dehydration, the cells were
214 embedded in resin. Embedded cell sections were then observed under a transmission EM.

215

216 **Ethical statement**

217 The animal experimental processes were approved by the Ethnic Committee of The First

218 Affiliated Hospital of Chongqing Medical University (AMUWEC20210719) and conducted
219 in strict accordance with the standard of the Guide for the Care and Use of Laboratory
220 Animals published by the Ministry of Science and Technology of the People's Republic of
221 China in 2006.

222

223 **Statistical analysis**

224 All statistical analyses were conducted using version 4.2.1 of R (R Foundation for Statistical
225 Computing). Quantitative data in this study were analyzed using GraphPad Prism version
226 9.5.0. Data were presented as mean \pm standard deviation. Initially, tests for normality and
227 homogeneity of variance were performed. If the data were normally distributed and the
228 variances were homogeneous, unpaired t-tests were used to compare differences between two
229 groups. One-way analysis of variance (ANOVA) was employed to compare differences
230 among multiple groups, followed by Tukey's post-hoc test for pairwise comparisons. A $P <$
231 0.05 was considered statistically significant, while a $P < 0.01$ was considered highly
232 significant.

233

234 **RESULTS**

235 **Activation of astrocytes and increased Rab8a expression in the spinal dorsal horn of** 236 **CCI rats**

237 In our study of the sciatic nerve ligation model in rats, we conducted IHC and IF staining to
238 investigate the potential mechanisms related to astrocytes in NPP. IHC analysis revealed a
239 significant increase in Rab8a expression in the spinal dorsal horn of rats subjected to sciatic
240 nerve ligation compared to controls (Figure 1A-B, $P < 0.01$).

241 IF staining further explored the distribution of Rab8a in the spinal dorsal horn. GFAP (green
242 fluorescence), a marker of astrocytes, showed a notable increase in the NPP model, indicating

243 the activation of astrocytes. Rab8a (red fluorescence) staining was observed in various cell
244 types within the spinal dorsal horn, but a significant increase in Rab8a expression was
245 evident in activated astrocytes (Figure 1C). These findings highlight the association between
246 Rab8a expression and astrocyte activation, suggesting its potential importance in the
247 pathophysiology of NPP.

248

249 **Upregulation of cytokine expression in LPS-induced activated astrocytes**

250 LPS is commonly utilized to simulate inflammatory responses, prompting a series of
251 experiments to investigate its effects on astrocytes. Initially, we assessed the viability of
252 astrocytes to gauge the activating effect of LPS. The results demonstrated a significant
253 increase in the survival rate of astrocytes cultured with LPS compared to the control group
254 (Figure 2A, $P < 0.05$).

255 IF and Western blot analyses revealed a significant increase in the expression of the GFAP
256 protein in cells treated with LPS (Figure 2B, $P < 0.05$). Moreover, compared to the control
257 group, cells in the LPS group exhibited increased cell volume and shorter, thicker processes
258 (Figure 2C). This phenomenon likely reflects the morphological changes of astrocytes under
259 LPS treatment, further supporting their activated state.

260 Further analysis through qPCR was conducted to measure the mRNA levels of
261 pro-inflammatory cytokines, including TNF- α and IL-1 β , in activated astrocytes. The results
262 showed significant upregulation of TNF- α and IL-1 β mRNA levels in activated astrocytes
263 (Figure 3A-B, $P < 0.001$).

264 In summary, our findings reveal the activating effects of LPS on astrocytes, including
265 increased cell viability, elevated expression of GFAP protein, morphological changes, and the
266 upregulation of pro-inflammatory cytokines TNF- α and IL-1 β mRNA levels in activated
267 astrocytes.

268 **Increased vesicular transport in LPS-induced activated astrocytes**

269 To investigate changes in vesicular transport within activated astrocytes, EM was employed
270 to examine vesicular transport. In the LPS-treated group, the Golgi apparatus was increased
271 and enlarged, with more Golgi vesicles around the trans-Golgi network (TGN).
272 Mitochondrial numbers were increased, showing swollen, spherical forms with reduced
273 cristae. The quantity of vesicles in activated astrocytes was higher in the LPS group, with
274 vesicles accumulating near the cell membrane (Figure 4A-B).

275 Given the molecular basis of vesicle and plasma membrane fusion established by the SNARE
276 complex [44], the co-expression of Rab8a and VAMP2 in activated astrocytes was further
277 investigated through IF experiments. It was observed that the positive IF staining of Rab8a
278 and VAMP2 was more aggregated in activated astrocytes. Additionally, astrocytes activated
279 by LPS also exhibited co-localization of Rab8a and VAMP2 expression, with these proteins
280 displaying a relatively uniform distribution across various cell types (Figure 5A).

281 Furthermore, co-localization of Rab8a and Syntaxin16 expression was also observed in
282 astrocytes activated by LPS. However, the distribution of the positive IF staining for Rab8a
283 and Syntaxin16 was not entirely consistent across different activated astrocytes (Figure 5B).

284 Western blot analysis further examined the expression levels of vesicular transport-related
285 proteins (Rab8a, VAMP2, and Syntaxin16). The results indicated that, following LPS
286 treatment, the levels of Rab8a, VAMP2, and Syntaxin16 proteins were significantly higher in
287 astrocytes compared to the control group (Figure 5C-E, $P < 0.05$).

288 These findings reveal increased vesicular transport in LPS-induced activated astrocytes,
289 accompanied by upregulation of protein levels of Rab8a and SNARE.

290

291 **BONT/A inhibits vesicular transport in LPS-induced activated astrocytes**

292 To assess the impact of the vesicular secretion inhibitor BONT/A on vesicular transport, we

293 conducted Western blot experiments to detect changes in proteins related to vesicular
294 transport (Rab8a, VAMP2, and Syntaxin16) in astrocytes activated by LPS treatment
295 following BONT/A administration. The results indicated a significant reduction in the
296 expression levels of Rab8a, VAMP2, and Syntaxin16 proteins in astrocytes treated with
297 BONT/A (BTX group) compared to those treated with LPS alone (LPS group) (Figure
298 6A-D).

299 To gain a comprehensive understanding of BONT/A's effect, EM was used to evaluate
300 changes in vesicular transport. The findings demonstrated that vesicular transport within
301 astrocytes activated by LPS was significantly inhibited following BONT/A treatment,
302 evidenced by a decrease in the number of intracellular vesicles and a marked reduction in
303 vesicle accumulation near the cell membrane (Figure 7A-B).

304 These results reveal the inhibitory effect of BONT/A on vesicular transport in astrocytes
305 activated by LPS induction.

306

307 **DISCUSSION**

308 Our research demonstrates that injury signals drive the transformation and activation of
309 astrocytes, leading to increased release of pain-associated bioactive molecules such as
310 inflammatory factors, ATP, and glutamate. These molecules play roles in central and
311 peripheral sensitization and contribute to the progression of NPP [18, 19]. Astrocyte
312 activation is a heterogeneous process involving multiple molecular, cellular, and functional
313 changes, including alterations in vesicular secretion [25, 54, 55, 56]. However, the specific
314 mechanisms underlying vesicle and inflammatory mediator release remain unclear [32].
315 Rab8a protein and the SNARE complex are involved in vesicle-directed transport, targeting
316 docking, and fusion with the cell membrane [43, 44, 45], but their mechanisms in NPP have
317 yet to be confirmed.

318 In our study using a CCI rat model, Rab8a was highly expressed in astrocytes within the
319 spinal dorsal horn following neural injury, suggesting increased vesicle docking and transport
320 activity, a possible manifestation of astrocyte activation. EM revealed a significant increase in
321 internal vesicle number and transport activity toward the plasma membrane, resulting in
322 heightened exocytic activity. Quantitative PCR, IF, and Western blot results showed
323 significant increases in the expression of cytokines such as TNF- α and IL-1 β , as well as
324 Rab8a, VAMP2, and Syntaxin16 in activated astrocytes. Treatment with BONT/A
325 significantly reduced the levels of Rab8a, VAMP2, and Syntaxin16 proteins in astrocytes.
326 Collectively, these findings suggest that the activation of the Rab8a/SNARE complex
327 pathway is crucial for vesicular transport and bioactive molecule release in astrocytes and
328 represents an important component in the pathogenesis of NPP.

329 Rab8a, a small GTPase, is essential for vesicle transport in various cell types and is involved
330 in cilia formation [57, 58]. Rab8a can interact with effectors or directly with SNARE to
331 recognize t-SNARE on target membranes, promoting v-SNARE and t-SNARE pairing, thus
332 guiding vesicle-directed transport and targeted docking [59, 60, 61, 62, 63, 64]. The control
333 of vesicle transport by Rab8a may facilitate the formation of different membrane protrusions,
334 while VAMP2 and Syntaxin16, components of the SNARE complex, are critical proteins in
335 vesicle docking. Our results, combined with previous studies [65, 66], suggest that
336 Rab8a-mediated enhanced transport of vesicles from the trans-Golgi network (TGN) to the
337 plasma membrane may underpin the molecular basis for astrocyte release of bioactive
338 molecules involved in the onset and maintenance of NPP. Enhanced vesicular transport in
339 LPS-activated astrocyte models likely represents a crucial mechanism for the secretion of
340 bioactive molecules by activated astrocytes, with activated pathways for cytokine synthesis
341 and secretion contributing to disease progression.

342 Furthermore, the application of BONT/A suggests that targeting components of the SNARE

343 complex can effectively reduce vesicular transport in astrocytes. Preclinical and clinical
344 studies have reported the efficacy of BONT/A in treating central NPP. BONT/A inhibits the
345 secretion of substance P and CGRP in DRG, suppresses the expression of TRPV1 and P2X3,
346 and exerts central effects through retrograde axonal transport [67, 68, 69, 70]. BONT/A not
347 only cleaves SNAP-25 at presynaptic terminals but also cleaves SNARE proteins
348 retrogradely in growth cones and the central brain, inhibiting the exocytosis of vesicles
349 containing norepinephrine, glutamate, substance P, and calcitonin gene-related peptide
350 (CGRP), as well as the expression of vanilloid receptors.

351

352 **CONCLUSION**

353 In summary, our study reveals that the activation of the Rab8a/SNARE complex pathway and
354 subsequent enhanced vesicle transport activity in the spinal dorsal horn following neural
355 injury are likely critical components in the cytokine cascade reaction mechanisms of NPP
356 (Figure 8). By elucidating the role of the Rab8a/SNARE complex in the development of NPP,
357 this study provides important insights for understanding the molecular basis of NPP and
358 developing new therapeutic strategies. Given the persistent activation of astrocytes under
359 chronic pain conditions and their recognized role in NPP, directing therapeutic interventions
360 towards reactive astrocytes holds significant potential. Our research demonstrates the critical
361 role of these proteins in astrocytes and emphasizes the importance of vesicle transport in
362 regulating NPP, offering new potential targets for NPP treatment. Targeting the
363 Rab8a/SNARE complex pathway could be an effective strategy for alleviating or treating
364 NPP. Based on our current understanding of astrocyte-mediated NPP, considering targeting
365 related signaling pathways, hemichannels, or purinergic receptors to inhibit the release of
366 neuroglial mediators, such as by inhibiting the expression or function of Rab8a to reduce the
367 release of inflammatory mediators, could provide valuable directions for developing novel

368 NPP therapeutic drugs. Additionally, targeting downstream mediators released by astrocytes,
369 such as chemokines and cytokine signaling, is a viable treatment strategy. Given that
370 astrocyte dysregulation is a common feature of nearly all chronic pain pathologies, and the
371 activation of astrocytes remains strong throughout persistent pain conditions, whether
372 targeting the activation of astrocytes or preventing their transition to a pro-inflammatory state
373 without affecting their normal homeostatic functions remains a significant challenge.

374 Although this study provides important insights into the role of the Rab8a/SNARE complex
375 in NPP, it has limitations. Firstly, the study is primarily based on animal models and cell
376 experiments, and its results need further validation in humans. Secondly, although BONT/A
377 can inhibit vesicular transport in astrocytes, its specific mechanisms of action and long-term
378 effects require further investigation. Additionally, this study did not fully resolve all potential
379 molecular mechanisms of the Rab8a/SNARE complex pathway in the pathogenesis of NPP,
380 necessitating further research to elucidate these mechanisms. Future research should focus on
381 several key areas. Firstly, the findings of this study need to be validated in a broader range of
382 biological models and explored through clinical studies to assess their potential application in
383 human NPP treatment. Secondly, specific intervention methods targeting the Rab8a/SNARE
384 complex pathway, including small molecule inhibitors, and RNA interference techniques,
385 should be explored to develop new treatment strategies. Additionally, investigating the role of
386 the Rab8a/SNARE complex in other cell types beyond astrocytes, such as neurons and
387 microglia, may reveal more complex pathological mechanisms of NPP.

388

389 **Author contributions**

390 YQX, GYW and YS designed the study. GQH and WXQ collated the data, carried out data
391 analyses and produced the initial draft of the manuscript. YQX, GYW and YS contributed to
392 drafting the manuscript. All authors have read and approved the final submitted manuscript.

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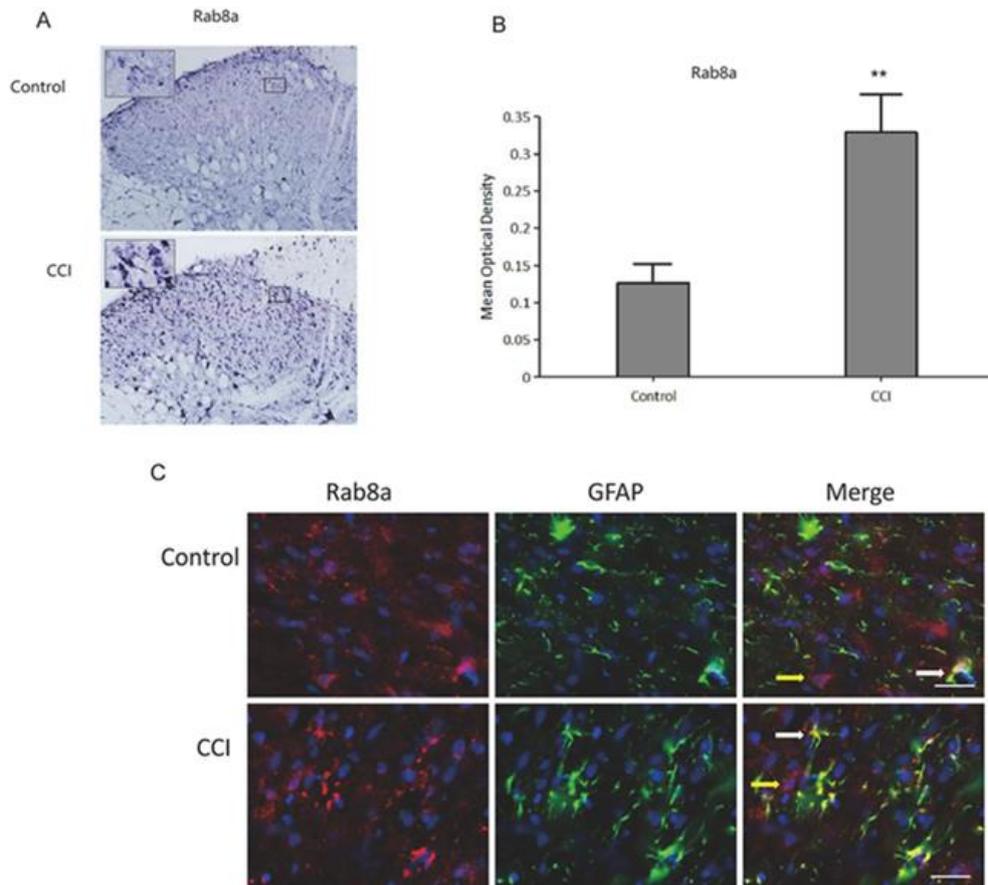
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645

646 **TABLES AND FIGURES WITH LEGENDS**



647

648 **Figure 1. Activation of astrocytes and Rab8a expression in the spinal dorsal horn of CCI**

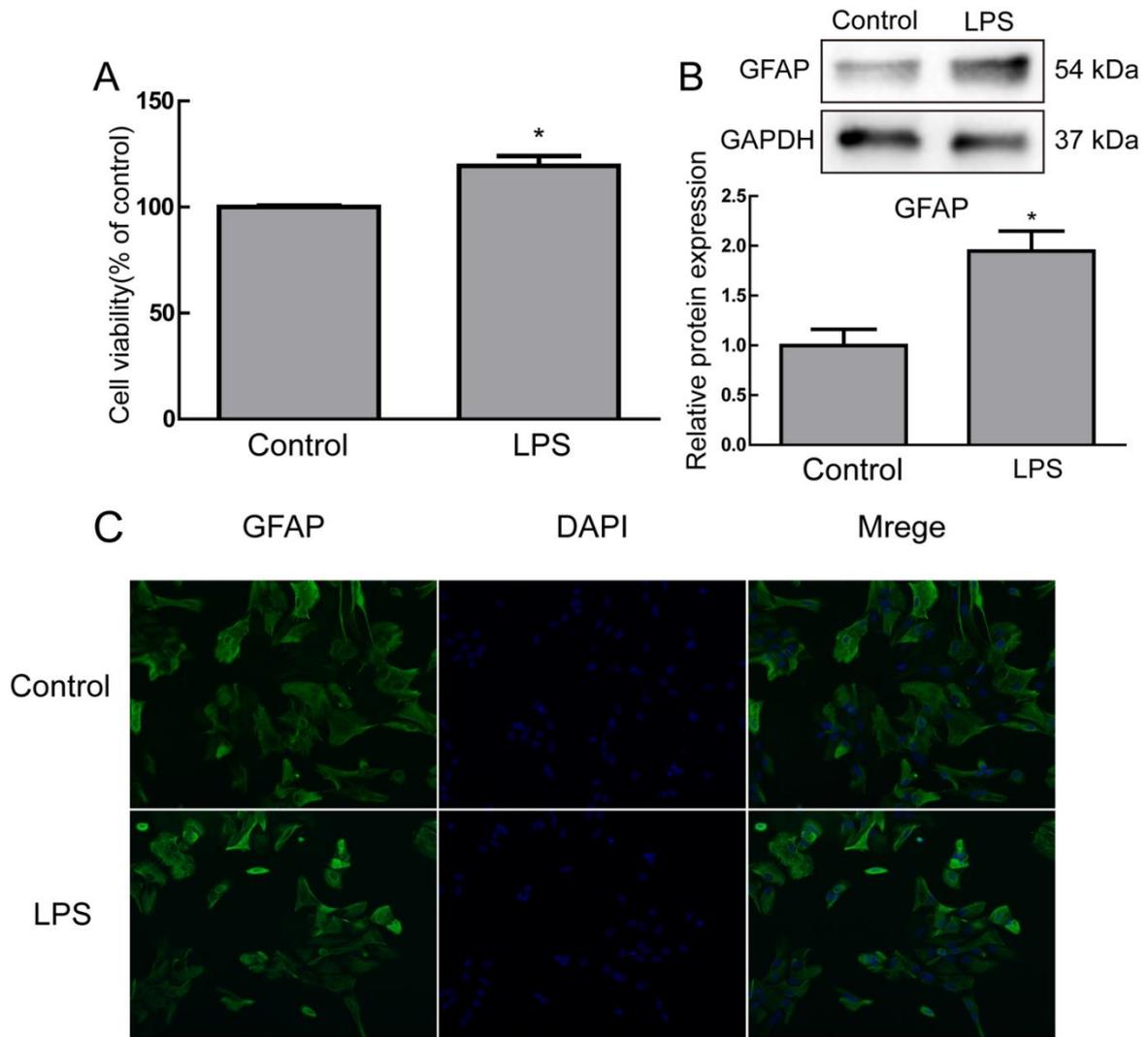
649 **rats.** (A) IHC staining of Rab8a in the spinal dorsal horn of CCI rats; (B) Quantitative

650 analysis of Rab8a protein in the spinal dorsal horn of CCI rats; (C) IF staining in the spinal

651 dorsal horn of CCI rats, showing GFAP-positive cells (green fluorescence) and the

652 distribution of Rab8a (red fluorescence). ** $P < 0.01$. CCI: Chronic constriction injury; IHC:

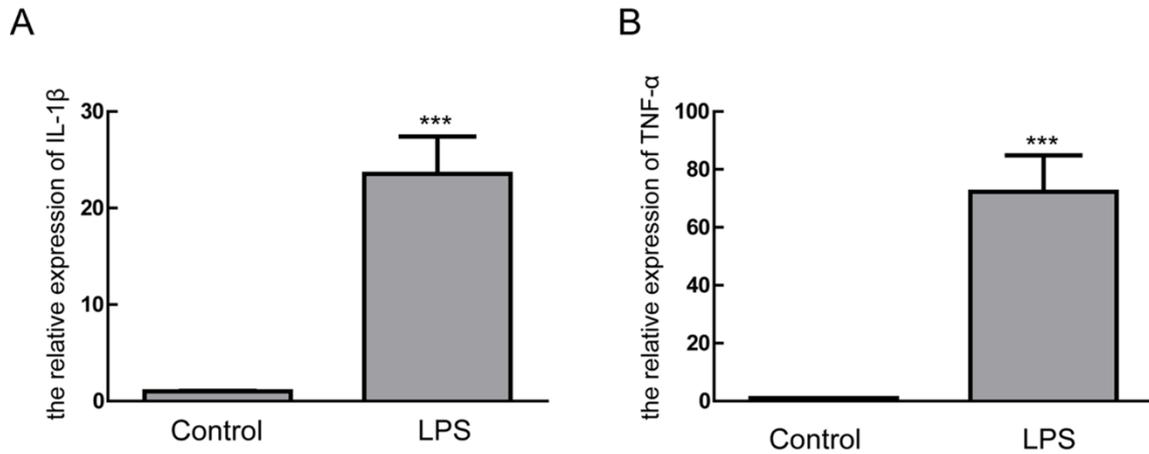
653 Immunohistochemistry; GFAP: Glial fibrillary acidic protein.



654

655 **Figure 2. The effects of LPS on astrocytes.** (A) The survival rate of astrocytes post-LPS
 656 treatment; (B) GFAP protein expression in astrocytes post-LPS treatment detected by Western
 657 blot; (C) IF staining of astrocytes post-LPS treatment, showing GFAP-positive cells (green
 658 fluorescence) and morphological changes. * $P < 0.05$. GFAP: Glial fibrillary acidic protein;
 659 LPS: Lipopolysaccharide; IF: Immunofluorescence.

660



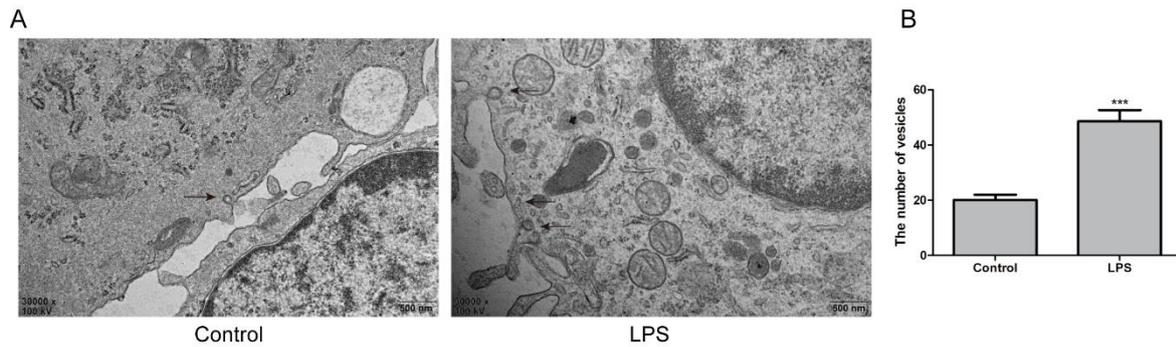
661

662 **Figure 3. Expression of cytokines in activated astrocytes.** (A) qPCR detection of TNF-α

663 mRNA levels in astrocytes post-LPS treatment; (B) qPCR detection of IL-1β mRNA levels in

664 astrocytes post-LPS treatment. *** $P < 0.001$. LPS: Lipopolysaccharide.

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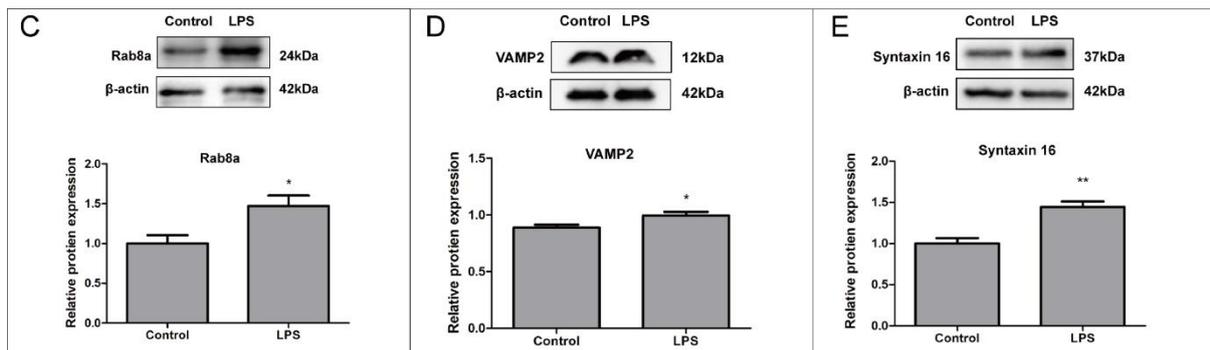
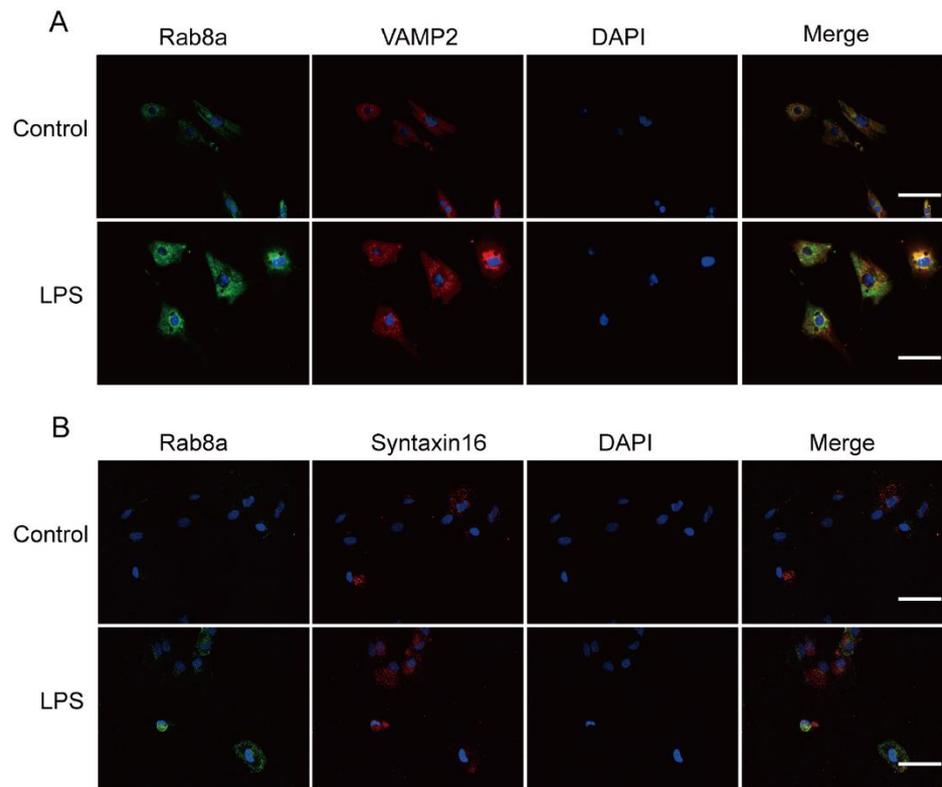
667 **Figure 4. Changes in vesicular transport in activated astrocytes.** (A) EM observation of

668 vesicular transport in astrocytes post-LPS treatment; (B) Statistical graph of vesicular

669 transport quantity in astrocytes post-LPS treatment. *** $P < 0.001$. EM: Electron microscopy;

670 LPS: Lipopolysaccharide.

671



672

673 **Figure 5. Levels of vesicular transport-related proteins in activated astrocytes.** (A-B) IF

674 staining of VAMP2 and Syntaxin16 in astrocytes post-LPS treatment; (C-E) Western blot

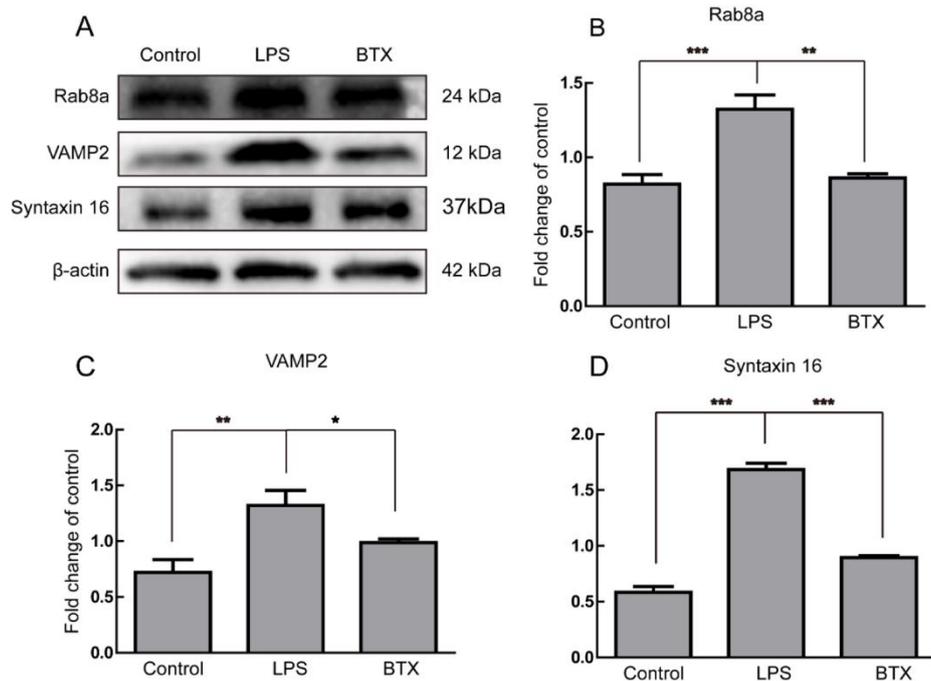
675 detection of Rab8a and SNARE proteins (VAMP2 and Syntaxin16) levels in astrocytes

676 post-LPS treatment. * $P < 0.05$, ** $P < 0.01$. IF: Immunofluorescence; VAMP2:

677 Vesicle-associated membrane protein; SNARE: Soluble N-ethylmaleimide-sensitive factor

678 attachment protein receptor.

679



680

681 **Figure 6. Effects of BONT/A on vesicular transport-related proteins in activated**

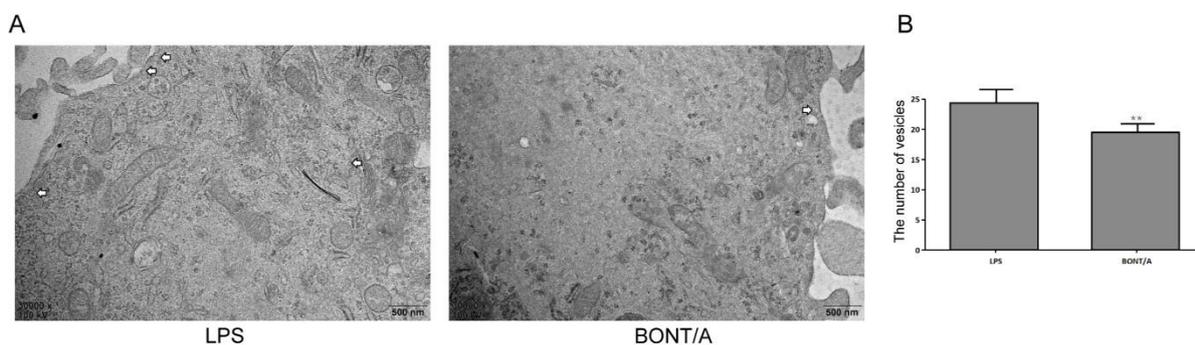
682 **astrocytes. (A)** Western blot detection of Rab8a, VAMP2, and Syntaxin16 protein expression

683 in activated astrocytes post-BONT/A treatment; (B-D) Quantitative analysis of Rab8a,

684 VAMP2, and Syntaxin16 proteins in activated astrocytes post-BONT/A treatment. * $P < 0.05$,

685 ** $P < 0.01$, *** $P < 0.001$. BONT/A: Botulinum neurotoxin type A.

686



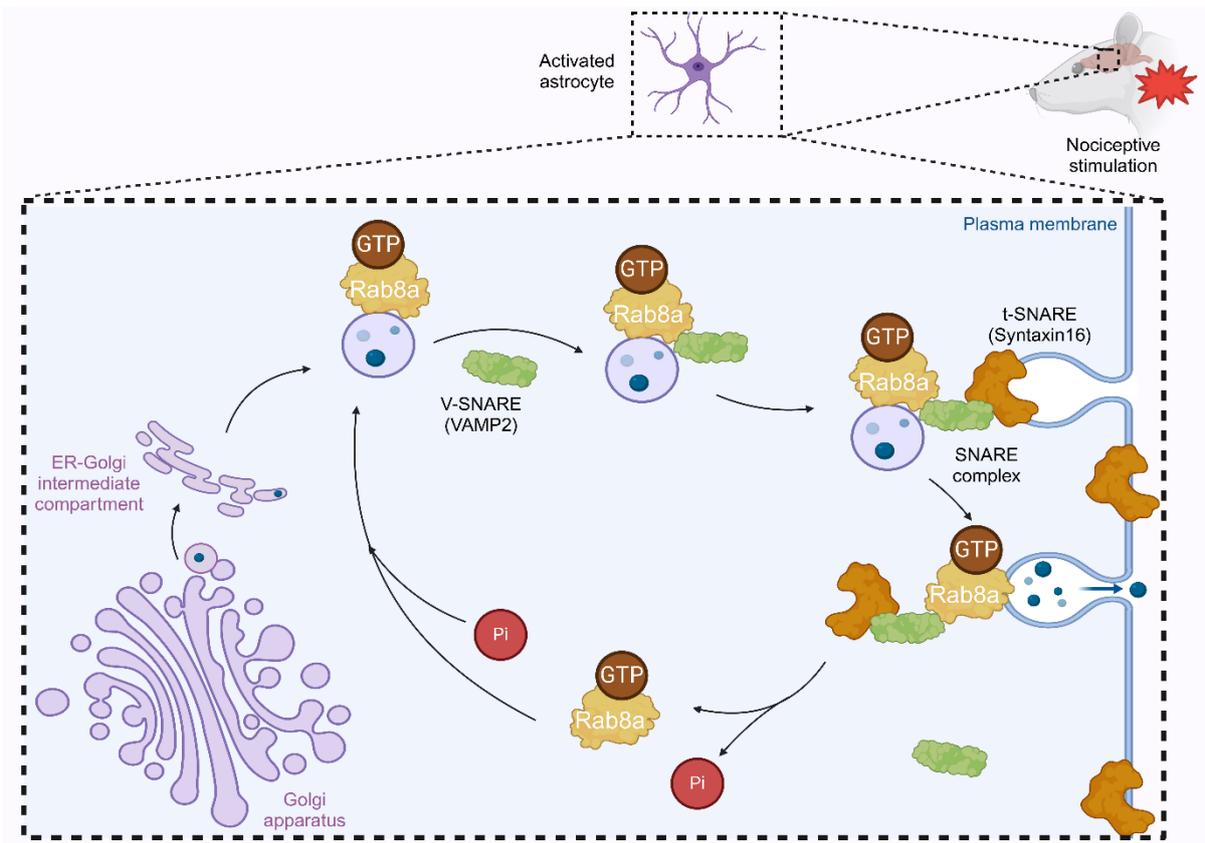
687

688 **Figure 7. Effects of BONT/A on vesicular transport in activated astrocytes. (A)** EM

689 observation of vesicular transport in activated astrocytes post-BONT/A treatment; (B)

690 Statistical analysis of vesicular transport quantity in activated astrocytes post-BONT/A

691 treatment. ** $P < 0.01$. BONT/A: Botulinum neurotoxin type A; EM: Electron microscopy.



692

693 **Figure 8. Activated Rab8a/SNARE complex drives the molecular mechanism of NPP by**
 694 **promoting vesicle anchoring and transportation in spinal astrocytes. NPP: Neuropathic**
 695 **pain.**