

1 **Pannexin-1 promotes NLRP3 activation during apoptosis but is**
2 **dispensable for canonical or non-canonical inflammasome activation**

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16 apoptosis

17

18 **Abstract**

19 Inflammasomes are multimeric protein complex that assemble in the cytosol
20 upon microbial infection or cellular stress. Upon activation, inflammasomes
21 drive the maturation of proinflammatory cytokines interleukin (IL)-1 β and IL-18,
22 and also activate the pore-forming protein gasdermin D to initiate a form of lytic
23 cell death known as 'pyroptosis'. Pannexin-1 is channel-forming glycoprotein
24 that promotes membrane permeability and ATP release during apoptosis; and
25 was implicated in canonical NLRP3 or non-canonical inflammasome activation.
26 Here, by utilising three different pannexin-1 channel inhibitors and two lines of
27 *Panx1*^{-/-} macrophages, we provide genetic and pharmacological evidence that
28 pannexin-1 is dispensable for canonical or non-canonical inflammasome
29 activation. Instead, we demonstrate that pannexin-1 cleavage and resulting
30 channel activity during apoptosis promotes NLRP3 inflammasome activation.
31

32 **Introduction**

33

34 The innate immune system provides the first line of defence against microbial
35 pathogens but also drives inflammatory diseases. Inflammasomes, multimeric
36 protein complexes that assemble in the cytosol in response to infection or
37 cellular stress, are now emerging as key molecular drivers of both processes
38 [1]. The NLRP3 inflammasome comprises the NLRP3 sensor protein, the
39 adaptor protein ASC and the protease caspase-1. NLRP3 is an unusual sensor
40 protein because it has a unique ability to sense a wide variety of structurally-
41 unrelated molecules ranging from whole pathogens, bacterial toxins, ATP
42 released from damaged cells, particulate crystals or misfolded protein
43 aggregates such as α -synuclein generated during neurodegenerative disease
44 [2]. The precise mechanism by which NLRP3 senses these structurally diverse
45 agonists remains unclear, however, it appears that the vast majority of NLRP3
46 agonists elicit a common potassium efflux stress pathway that is sensed by
47 NLRP3 [3]. Upon activation, NLRP3 oligomerises and recruits the adaptor
48 protein ASC into large filamentous structure called the 'ASC speck', which
49 creates a multitude of binding sites for the caspase-1 zymogen [4, 5]. Caspase-
50 1 recruited within the inflammasome undergoes proximity-induced activation
51 and autoprocessing to generate an active p33/p10 fragment, and is further
52 processed to an inactive p20/p10 fragment that is released from the
53 inflammasome [6]. Active caspase-1 cleaves the pore-forming protein
54 gasdermin D (GSDMD) to induce a form of lytic cell death known as pyroptosis
55 [7, 8]. In parallel, caspase-1 cleaves interleukin (IL)-1 family members such as
56 pro-IL-1 β and triggers IL-1 β relocation to the plasma membrane to enable
57 GSDMD-dependent and independent secretion [9-11].

58

59 The non-canonical inflammasome pathway is activated by cytoplasmic
60 lipopolysaccharide (LPS) from Gram-negative bacteria [12]. Unexpectedly,
61 LPS sensing by the non-canonical inflammasome does not require a classical
62 pattern recognition receptor. Instead, cytoplasmic LPS appears to directly bind
63 the caspase recruitment domain (CARD) of caspase-11, triggering caspase-11
64 oligomerization and activation [13]. In macrophages, active caspase-11 cleaves

65 its substrate, GSDMD to trigger pyroptosis [7, 8], while in neutrophils, active
66 GSDMD promotes plasma and nuclear membrane damage and the extrusion
67 of neutrophil extracellular traps [14]. Of note, caspase-11 is unable to directly
68 process pro-IL-1 β , but does so indirectly by triggering GSDMD pores,
69 potassium efflux and NLRP3 inflammasome activation in a cell-intrinsic manner
70 [7, 8, 12, 15].

71

72 Pannexin-1 is a channel-forming transmembrane protein with that is expressed
73 in most cell types, including macrophages. Under resting state, the pannexin-1
74 channel is autoinhibited by its cytoplasmic C-terminal tail. During apoptosis,
75 effector caspase-3 and -7 cleave pannexin-1 at its C-terminus to promote
76 pannexin-1 channel opening and drive membrane permeability [16, 17]. In
77 agreement with this model, we recently demonstrate that pannexin-1 channels
78 promote NLRP3 inflammasome assembly during apoptosis [18]. Interestingly,
79 exposure of macrophages to pannexin-1 inhibitor carbenoxolone or the
80 pannexin-1 inhibitory peptide ¹⁰PANX were reported to suppress canonical
81 NLRP3 inflammasome activation [19-21]. More recently, pannexin-1 channel
82 was also implicated to promote pyroptosis and NLRP3 activation following the
83 assembly of the non-canonical inflammasome [22]. However, this finding is at
84 odds with the observation that caspase-11 drives NLRP3 inflammasome
85 activation through GSDMD pores [7, 8]. Whether pannexin-1 is required for
86 optimal GSDMD processing and NLRP3 inflammasome activation after
87 cytoplasmic LPS recognition is unclear and a focus of this study. Here, by using
88 two independent lines of *Panx1*-deficient cells, we provide genetic evidence
89 that pannexin-1 is only required for NLRP3 assembly during apoptosis but is
90 dispensable for canonical NLRP3 or non-canonical inflammasome activation.

91

92 **Results and discussion**

93

94 *Pannexin-1 channel activity is required for NLRP3 inflammasome activation*
95 *during apoptosis*

96

97 The requirement for pannexin-1 in canonical and non-canonical inflammasome
98 activation is controversial [19-21, 23]. To examine the function of pannexin-1 in
99 inflammasome activation, we first tested probenecid, trovafloxacin and
100 spironolactone, three well-established pharmacological inhibitors of pannexin-
101 1 channel activity [24-26]. We recently demonstrated that pannexin-1 promotes
102 potassium efflux and NLRP3 inflammasome activation during intrinsic and
103 extrinsic apoptosis in macrophages [18]. In agreement with that, we observed
104 robust processing of full length pannexin-1 into the cleaved p19 fragment when
105 bone marrow-derived macrophages (BMDMs) were exposed to the BH3-
106 mimetic ABT-737 and MCL1 inhibitor S63845 to induce intrinsic apoptosis (**Fig.**
107 **1A**). Probenecid, trovafloxacin and spironolactone had little to no effect on
108 pannexin-1 cleavage, but completely abrogated caspase-1 processing in
109 apoptotic BMDMs, indicating that pannexin-1 channel activity and resultant
110 cellular permeability drives NLRP3 inflammasome assembly during apoptosis
111 (**Fig. 1A**). Next, we investigated whether pannexin-1 inhibitors similarly block
112 IL-1 β secretion during intrinsic apoptosis. For this, we primed BMDMs with LPS
113 for 3 h to induce pro-IL-1 β expression, and stimulated the cells for another 16
114 h with the ABT-737 and S63845 for to induce intrinsic apoptosis in the presence
115 of absence of pannexin-1 inhibitors. Pannexin-1 inhibitors had little to no effect
116 on macrophage cytotoxicity (**Fig. 1B**). Consistent with our recent finding [18],
117 we observed that IL-1 β secretion was significantly reduced in apoptotic *Panx1*^{-/-}
118 ^{-/-} macrophages compared to WT cells. In agreement with this, the pannexin-1
119 inhibitor, spironolactone similarly reduced IL-1 β secretion in apoptotic wild type
120 macrophages (**Fig. 1C**). Of note, spironolactone also further reduced IL-1 β
121 secretion in *Panx1*^{-/-} BMDMs, indicating that spironolactone additionally
122 inhibits IL-1 β secretion through pannexin-1-independent mechanisms.
123 Surprisingly, while probenecid and trovafloxacin blocked caspase-1 processing
124 in unprimed apoptotic macrophages over a 5 h period (**Fig. 1A**), they were
125 ineffective in blocking IL-1 β secretion in LPS-primed apoptotic macrophages
126 over 16 h (**Fig. 1C**). Next, we investigated if pannexin-1 channel activity
127 similarly promotes NLRP3 activation during extrinsic apoptosis. Indeed,
128 probenecid, trovafloxacin and spironolactone similarly reduced caspase-1
129 cleavage but not pannexin-1 processing when BMDMs were treated with TNF

130 and the SMAC-mimetic AZD5582 to trigger extrinsic apoptosis (**Fig. 1D**). Taken
131 together, these data indicate that during apoptosis, pannexin-1 channel
132 opening promotes NLRP3 inflammasome activation, most likely by triggering
133 membrane permeability.

134

135 *Pannexin-1 is not required for canonical NLRP3 inflammasome activation*

136

137 Having established that pannexin-1 channel inhibitors suppress NLRP3
138 activation during apoptosis (**Fig. 1A, D**), we next investigated the function of
139 pannexin-1 in canonical NLRP3 inflammasome signalling. For this, we primed
140 wild type (WT) or *Panx1*^{-/-} BMDMs with LPS for 4 h to induce NLRP3 and pro-
141 IL-1 β expression, and stimulated the cells with the soluble NLRP3 agonists
142 nigericin and ATP, or the insoluble particulate agonist monosodium urate
143 (MSU). In contrast to apoptotic triggers (**Fig. 1A, D**), probenecid, trovafloxacin
144 and spironolactone had no effect on caspase-1 processing in nigericin-
145 stimulated macrophages (**Fig. 2A**). In agreement with that, cleavage of the
146 caspase-1 substrate GSDMD was unaffected by pannexin-1 inhibitors,
147 although a minor reduction in pro-IL-1 β processing was consistently observed
148 (**Fig. 2A**). Importantly, caspase-1 processing and cleavage of the caspase-1
149 substrates, pro-IL-1 β and GSDMD were comparable between WT and *Panx1*^{-/-}
150 *Panx1*^{-/-} BMDMs (**Fig. 2A, Supplementary Fig. 1A**). IL-1 β secretion and macrophage
151 pyroptosis, as measured by lactate dehydrogenase (LDH) release, were similar
152 between WT and *Panx1*^{-/-} BMDMs (**Fig. 2B-C, Supplementary Fig. 1B**). In
153 contrast, nigericin-induced IL-1 β secretion and pyroptosis were completely
154 dependent on GSDMD (**Fig. 2D-E**), consistent with previous reports (ref). Next,
155 we investigated whether pannexin-1 is required for ATP-induced NLRP3
156 inflammasome activation. Consistent with before, we observed comparable
157 levels of caspase-1, pro-IL-1 β and GSDMD processing between WT and
158 *Panx1*^{-/-} BMDMs upon ATP stimulation (**Fig. 2D, Supplementary Fig. 1A**). As
159 expected, IL-1 β secretion and pyroptosis were similar between ATP-stimulated
160 WT and *Panx1*^{-/-} BMDMs, but completely abrogated in *Gsdmd*^{-/-} cells (**Fig. 2G-**
161 **J, Supplementary Fig. 1B**). Interestingly, probenecid but not trovafloxacin or
162 spironolactone reduced caspase-1 processing and cleavage of caspase-1

163 substrates in both WT and *Panx1*^{-/-} BMDM, indicating that probenecid inhibits
164 ATP-induced inflammasome activation in a pannexin-1-independent manner.
165 In support of this, a recent study revealed that probenecid directly blocks the
166 ATP-gated ion channel P2X7 [27]. Next, we investigated whether pannexin-1
167 is required for NLRP3 inflammasome upon exposure to the particulate agonist,
168 MSU. In agreement with our findings with soluble NLRP3 agonists, we found
169 that pharmacological or genetic blockade of pannexin-1 had no impact on
170 caspase-1, pro-IL-1 β and GSDMD processing in BMDMs following MSU
171 stimulation (**Fig. 2K**). In line with that, IL-1 β secretion and pyroptosis were
172 similarly unaffected by *Panx1*-deficiency (**Fig. 2L-M**). Consistent with a recent
173 report [28], we observed that MSU-induced cell lysis and IL-1 β secretion
174 occurs independently of the pyroptotic effector GSDMD (**Fig. 2N-O**). Lastly, we
175 examined pannexin-1 cleavage following canonical NLRP3 inflammasome
176 activation. In line with our earlier observations (**Fig. 2A-C, F-H, K-M**), we found
177 no evidence of pannexin-1 cleavage during canonical inflammasome activation,
178 although treatment with ABT-737/S63845 to trigger apoptosis induced robust
179 pannexin-1 cleavage in macrophages as expected (**Fig. 2P**). Taken together,
180 these data indicate that while pannexin-1 is required for driving NLRP3
181 inflammasome assembly during apoptosis, it is dispensable for canonical
182 NLRP3 inflammasome activation in macrophages.

183

184 *GSDMD but not pannexin-1 promotes cell death and NLRP3 assembly*
185 *following non-canonical inflammasome activation*

186

187 Two recent landmark studies revealed that caspase-11 cleaves its substrate
188 GSDMD to drive pyroptosis, potassium efflux and NLRP3 activation [7, 8]. By
189 contrast, another study proposed that caspase-11-dependent pyroptosis and
190 NLRP3 assembly requires pannexin-1 channel activity [22], raising the
191 possibility that pannexin-1 channels potentiate GSDMD pore activity. To
192 investigate whether GSDMD processing, cell death or NLRP3 inflammasome
193 activation require pannexin-1 channel activity, we prepared WT and *Gsdmd*^{-/-}
194 BMDMs and activated the non-canonical inflammasome by LPS transfection in
195 the presence or absence of the pannexin-1 inhibitors probenecid, trovafloxacin

196 and spironolactone. In agreement with previous reports [8, 12], we observed
197 that cytoplasmic LPS induced cleavage of the caspase-11 substrate GSDMD
198 into the active p30 fragment and resulted in pyroptosis (**Fig. 3A-B**). In addition,
199 GSDMD was required to drive caspase-1 processing, pro-IL-1 β maturation and
200 secretion (**Fig. 3A, C**). Exposure of LPS-transfected BMDMs to probenecid,
201 trovafloxacin and spironolactone had no effect on GSDMD, caspase-1 and pro-
202 IL-1 β processing (**Fig. 3A**), while all three pannexin-1 inhibitors suppressed
203 caspase-1 processing in apoptotic macrophages (**Fig. 1A, D**). Next, we
204 compared LPS transfection in WT and *Panx1*^{-/-} BMDMs. In line with our
205 inhibitor data, we observed comparable levels of GSDMD, caspase-1 and pro-
206 IL-1 β processing in WT and *Panx1*^{-/-} BMDMs (**Fig. 3C, Supplementary Fig.**
207 **1C**). And as expected, IL-1 β secretion and pyroptosis were unaffected by
208 *Panx1*-deficiency compared to WT BMDM (**Fig. 3D-E, Supplementary Fig.**
209 **1D-E**). However, all three pannexin-1 inhibitors triggered a mild, but non-
210 significant reduction in IL-1 β secretion in both WT and *Panx1*^{-/-} macrophages
211 (**Fig. 3F**). Lastly, we investigated the status of pannexin-1 cleavage upon non-
212 canonical inflammasome activation. In agreement with our observation that
213 pannexin-1 is dispensable for non-canonical inflammasome signalling (**Fig. 3D-**
214 **F**), we did not find any evidence of pannexin-1 cleavage during non-canonical
215 inflammasome activation; in contrast, full length pannexin-1 was completely
216 converted into the cleaved p19 fragment in apoptotic macrophages (**Fig. 3H**).
217 Taken together, our data support the conclusion that non-canonical
218 inflammasome drives pyroptosis, NLRP3 activation and IL-1 β secretion through
219 GSDMD pores but not pannexin-1 channels.

220

221 **Concluding remarks**

222

223 Pannexin-1 is a channel-forming transmembrane protein with that is widely
224 expressed in most cell types, including in transformed cells [29]. Our finding
225 that pannexin-1 channel opening during apoptosis promotes NLRP3
226 inflammasome activation has major clinical relevance, especially under
227 conditions where IL-1 β signalling has been implicated to impair tumour
228 clearance [30]. While our study only examined apoptosis induced by SMAC-

229 mimetic, BH3-mimetic and MCL1 inhibitor, it is likely that other commonly used
230 apoptosis-inducing chemotherapies such as cisplatin and doxorubicin would
231 similarly activate the NLRP3 inflammasome through pannexin-1 channels. The
232 pannexin-1 channel inhibitor spironolactone is an inexpensive, orally-available
233 drug that is already approved for human use [26]. Therefore, spironolactone
234 could potentially offer rapid clinical translation and be administered in
235 combination with standard chemotherapeutics to reduce NLRP3 and IL-
236 1 β signalling to promote tumour clearance.

237

238 Pharmacological pannexin-1 inhibitor or siRNA knockdown of pannexin-1 were
239 reported to suppress canonical NLRP3 inflammasome activation [19-21].
240 Based on these observations, it was proposed that pannexin-1 channels
241 promote NLRP3 inflammasome assembly. However, this concept was
242 subsequently challenged as NLRP3 inflammasome signalling were found to be
243 similar between WT and *Panx1*^{-/-} BMDMs [23]. By using two independent lines
244 of *Panx1*^{-/-} BMDMs and three different pannexin-1 channel inhibitors, we
245 confirmed previous reports and conclude that pannexin-1 is indeed dispensable
246 for canonical NLRP3 inflammasome activation in murine macrophages.

247

248 Recognition of cytosolic Gram-negative bacteria by caspase-11 promotes
249 pyroptosis and NLRP3 inflammasome activation, and this pathway confers
250 susceptibility to murine models of endotoxin shock (ref). However, the
251 mechanism by which caspase-11 drives cell death and NLRP3 activation
252 remains debated. While two landmark studies identified that caspase-11
253 cleaves GSDMD to trigger pyroptosis and NLRP3 activation [7, 8], another
254 study proposed that caspase-11-dependent pyroptosis and NLRP3 activation
255 requires the membrane glycoprotein pannexin-1 [22]. In this model, the authors
256 proposed that caspase-11 cleaves pannexin-1 to drive membrane permeability
257 and NLRP3 inflammasome activation, while pannexin-1-dependent ATP
258 release promotes pyroptosis via engagement of the purinergic receptor P2X7.
259 Here, by using three different pannexin-1 channel inhibitors and 2 lines of
260 *Panx1*^{-/-} BMDMs, we provide evidence that pannexin-1 is not processed upon
261 non-canonical inflammasome activation and demonstrate that pannexin-1 is

262 dispensable for non-canonical inflammasome signalling in BMDM. In contrast,
263 our study supports the conclusion that active caspase-11 promotes cell lysis
264 and NLRP3 activation via GSDMD pores. While the report from Yang *et al.* and
265 our study both utilised *Panx1*^{-/-} lines generated from C57BL/6 background, it
266 remains possible, but unlikely, that passenger mutation and genetic variation
267 between the *Panx1*^{-/-} lines might influence the outcome of the study. This is
268 because *Panx1*^{-/-} BMDMs used in the study of Yang *et. al* expressed
269 comparable levels of caspase-11 compared to WT cells; and are also likely to
270 express normal levels of GSDMD since *Panx1*-deficiency did not affect
271 pyroptosis upon NLRP3 and AIM2 activation in that study (ref). While we
272 provide clear evidence that pannexin-1 is dispensable for non-canonical
273 inflammasome signalling in BMDMs, at this moment, we cannot rule out the
274 possibility that that pannexin-1 may indeed confer susceptibility to murine
275 models of endotoxin shock *in vivo*. Since apoptotic caspases contribute to
276 endotoxin-induced lethality (ref), it is tempting to speculate that pannexin-1 may
277 potentially contribute to lethality by promoting NLRP3 activation, pyroptosis and
278 IL-1 β secretion in apoptotic cells (ref). Future studies should confirm the role of
279 pannexin-1 in endotoxin-induced lethality and explore the therapeutic function
280 of pannexin-1 inhibitors in such disease.

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285

286 **Materials and methods**

287

288 Mice

289 C57BL/6 and *Gsdmd*^{+/-} mice were housed in specific-pathogen free facilities in
290 the University of Lausanne. *Panx1*^{-/-} were previously described [31, 32] and
291 kindly provided by Prof. Marc Chanson and Prof. Nathalie Rouach.

292

293 Cell culture

294 Murine bone marrow-derived macrophages (BMDM) were prepared by
295 differentiating bone marrow progenitor cells for 7 days in DMEM (Gibco)
296 supplemented with 20% MCSF (3T3 supernatant), 10% heat-inactivated FCS
297 (Bioconcept), 10 mM HEPES (Bioconcept), penicillin/streptomycin
298 (Bioconcept) and non-essential amino acids (Gibco). Inflammasome and
299 apoptosis assays were performed on mature BMDMs on day 7–9 of
300 differentiation. All WT versus *Panx1*^{-/-} data presented in the main manuscript
301 were generated using *Panx1*^{-/-} and WT control BMDM provided by Prof Marc
302 Chanson [32], while all WT versus *Panx1*^{-/-} data in the supplementary data
303 were generated using *Panx1*^{-/-} and WT controls provided by Prof Nathalie
304 Rouach [31].

305

306 Apoptosis assay

307 BMDMs were plated in 96-well plates at a density of 5×10^4 cells per well in
308 complete media a day prior to stimulation. BMDMs were stimulated with a
309 combination of ABT-737 and S63845 (all 0.5 μ M; Selleckchem) or a
310 combination of recombinant murine TNF (100 ng/ml; Enzo) and the SMAC-
311 mimetic AZD5582 (0.5 μ M; Selleckchem) in Opti-MEM for 5 h to induce intrinsic
312 or extrinsic apoptosis respectively. To measure IL-1 β secretion, BMDMs were
313 primed for 4 h in Opti-MEM and stimulated with ABT-737 and S63845 (all 1 μ M;
314 Selleckchem) for 16 h. Where indicated, cells were treated with probenecid (1
315 mM; Sigma), trovafloxacin (10 μ M; Sigma) or spironolactone (20 μ M; Sigma)
316 30 min prior to cell stimulation.

317

318 Inflammasome assay

319 BMDMs were plated in 96-well plates at a density of 5×10^4 cells per well in
320 complete media a day prior to stimulation. To activate the canonical NLRP3
321 inflammasome, cells were primed with ultrapure *E. coli* 055:B5 LPS (100 ng/ml;
322 InvivoGen) for 4 h in Opti-MEM and stimulated with nigericin (5 μ M; Invivogen)
323 or ATP (2.5 mM; Sigma) for 90 min, or MSU (150 μ g/ml; Invivogen) for 4 h. To
324 activate the non-canonical inflammasome, cells were primed with Pam3CSK4
325 (1 μ g/ml; Invivogen) for 4 h in Opti-MEM and transfected with ultrapure *E. coli*
326 0111:B4 LPS (2 μ g/ml) using 0.25% Fugene HD (Promega) and centrifuged for
327 500 *g* for 10 min at 37°C. BMDMs were harvested 16 h post LPS transfection.
328 Where indicated, cells were treated with probenecid (1 mM; Sigma),
329 trovafloxacin (10 μ M; Sigma) or spironolactone (20 μ M; Sigma) at the last 20–
330 30 min of priming.

331

332 Immunoblotting

333 Cell extracts were lysed in boiling lysis buffer (66 mM Tris–Cl pH 7.4, 2%
334 SDS, 10 mM DTT, 1x NuPage LDS sample buffer; Thermo Fisher) and
335 resuspended with methanol/chloroform-precipitated supernatant. Mixed
336 supernatant and extracts were separated on 14% polyacrylamide gels and
337 transferred onto nitrocellulose membrane using Trans-blot Turbo (Bio-rad).
338 Antibodies for immunoblot were against caspase-1 p20 (casper-1; Adipogen;
339 1:1000), GSDMD (EPR19828; Abcam; 1:1000), pro-IL-1 β (AF-401-NA, R&D;
340 1:1000), pannexin-1 (D9M1C; Cell Signaling; 1:1000) and alpha-tubulin (DM1A;
341 Abcam; 1:5000).

342

343 LDH assay

344 Lactate dehydrogenase (LDH) release into the cell culture supernatant was
345 quantified using the TaKaRa LDH cytotoxicity detection kit (Clontech) according
346 to the manufacture's protocol.

347

348 IL-1 β ELISA

349 IL-1 β release into the cell culture supernatant was quantified using the mouse
350 IL-1 beta DuoSet ELISA Kit (R&D systems) according to the manufacture's
351 protocol.

352

353 Statistical analyses

354 Statistical analyses were performed using GraphPad Prism 7 software. All data
355 sets were analysed using the parametric t-test. Data were considered
356 significant when *P ≤ 0.05, **P ≤ 0.01.

357

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366

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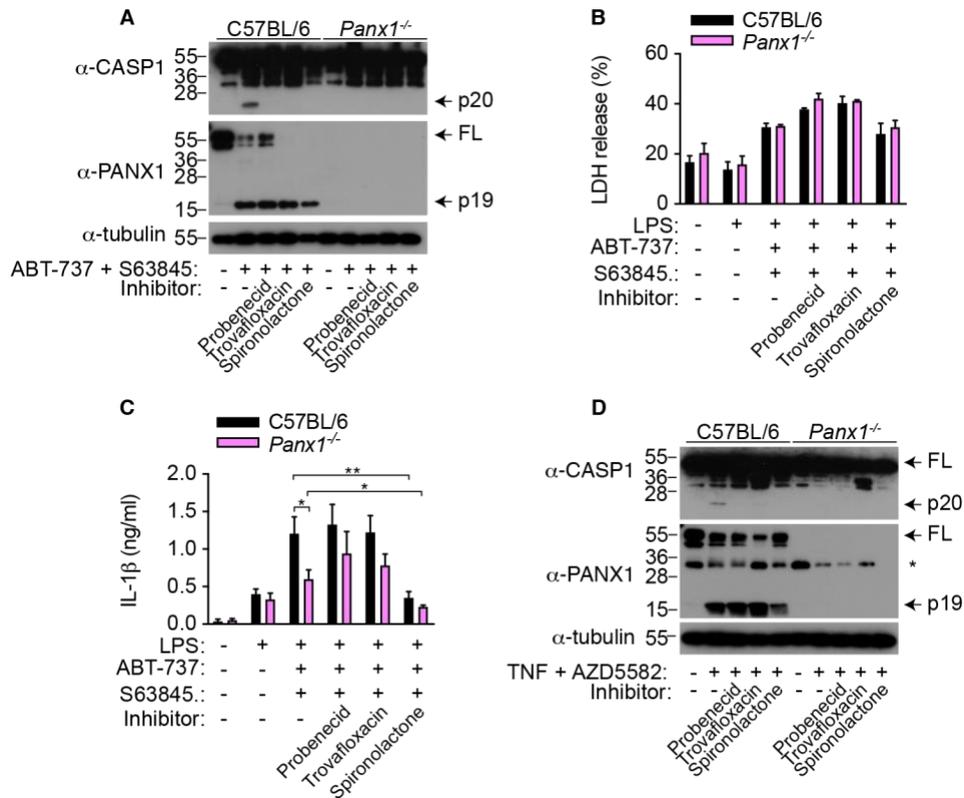
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494 **Figure legends**

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497 **Figure 1. Channel activity of pannexin-1 is required for NLRP3**

498 **inflammasome activation during apoptosis.** (A) BMDMs were

499 stimulated with ABT-737 (0.5 μM) and S63845 (0.5 μM) for 5 h to induce

500 intrinsic apoptosis. (B-C) BMDMs were primed with LPS (100 ng/ml) for 3 h and

501 stimulated with ABT-737 (1 μM) and S63845 (1 μM) for 16 h. (D) BMDMs were

502 stimulated with TNF (100 ng/ml) and AZD8882 (0.5 μM) for 5 h to induce

503 intrinsic apoptosis. Where indicated, pannexin-1 channel inhibitors probenecid

504 (1 mM), trovafloxacin (10 μM) and spironolactone (20 μM) were added 30 min

505 prior to cell stimulation. (B, C) Data are mean ± S.E.M of pooled data from 3

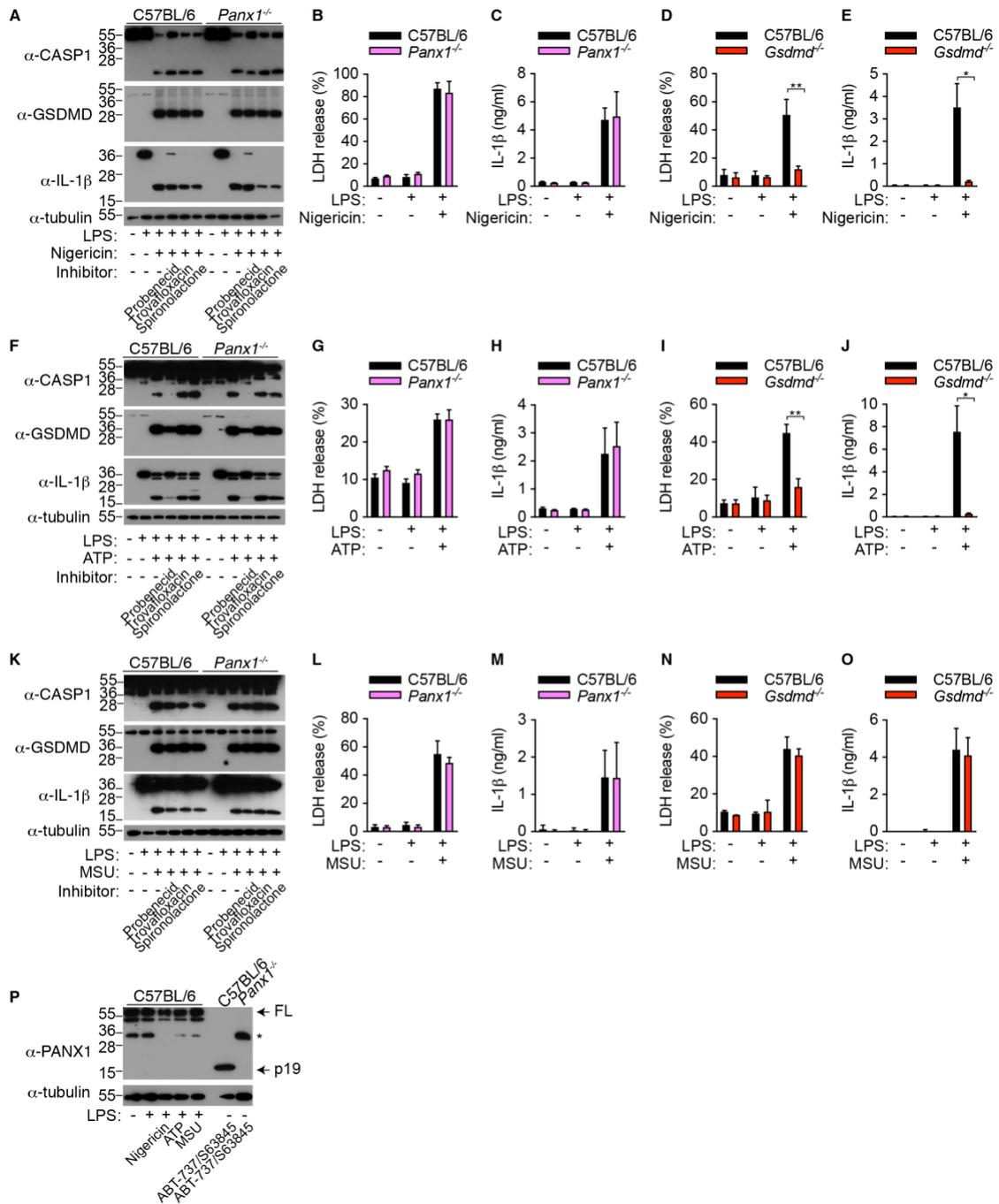
506 independent experiments. Data were considered significant when *P < 0.05,

507 **P < 0.01. Immunoblots show mixed supernatant and cell extracts and are

508 representative of 3 independent experiments. * indicates a cross-reactive band.

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513 **Figure 2. Pannexin-1 is dispensable for canonical NLRP3 inflammasome**

514 **activation.** (A-P) BMDMs were primed with 100 ng/ml ultrapure LPS for 4 h

515 and stimulated with nigericin (0.5 μM; 90 min), ATP (2.5 mM; 90 min) or MSU

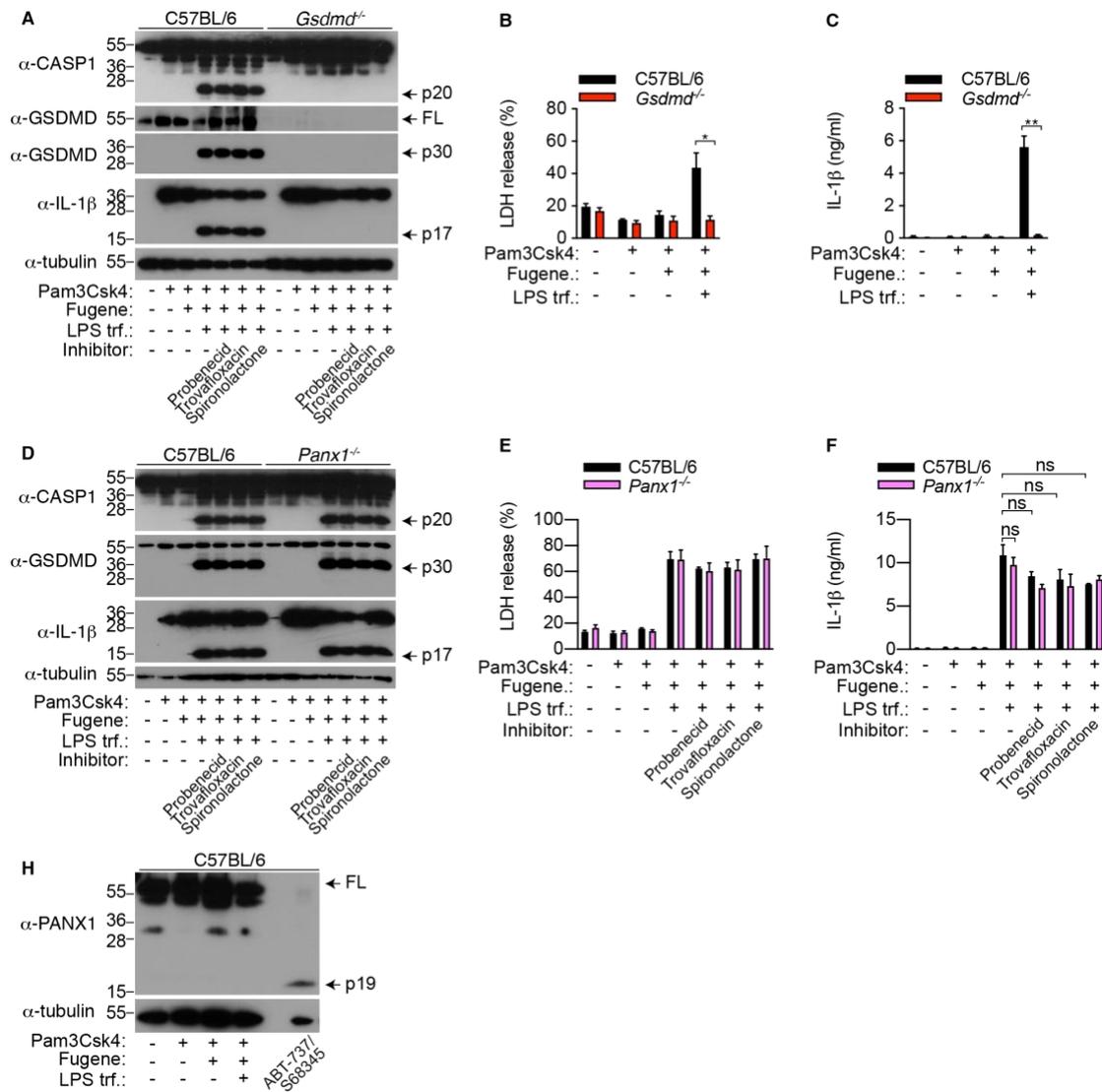
516 (150 μg/ml; 4 h), or stimulated with ABT-737 (0.5 μM) and S63845 (0.5 μM) for

517 5 h. Where indicated, pannexin-1 channel inhibitors probenecid (1 mM),

518 trovafloxacin (10 μM) and spironolactone (20 μM) were added 30 min prior to

519 cell stimulation. (A, F, K, P) Mixed supernatant and cell extracts were examined

520 by immunoblotting and are representative of at 2-3 independent experiments.
 521 Data are mean \pm S.D. of triplicate cell stimulation, representative of 3 (B, C, G,
 522 H, L, M) or 2 (N, O) independent experiments; (D, E, I, J) data are mean \pm
 523 S.E.M of pooled data from 3 independent experiments. Data were considered
 524 significant when *P < 0.05, **P < 0.01.
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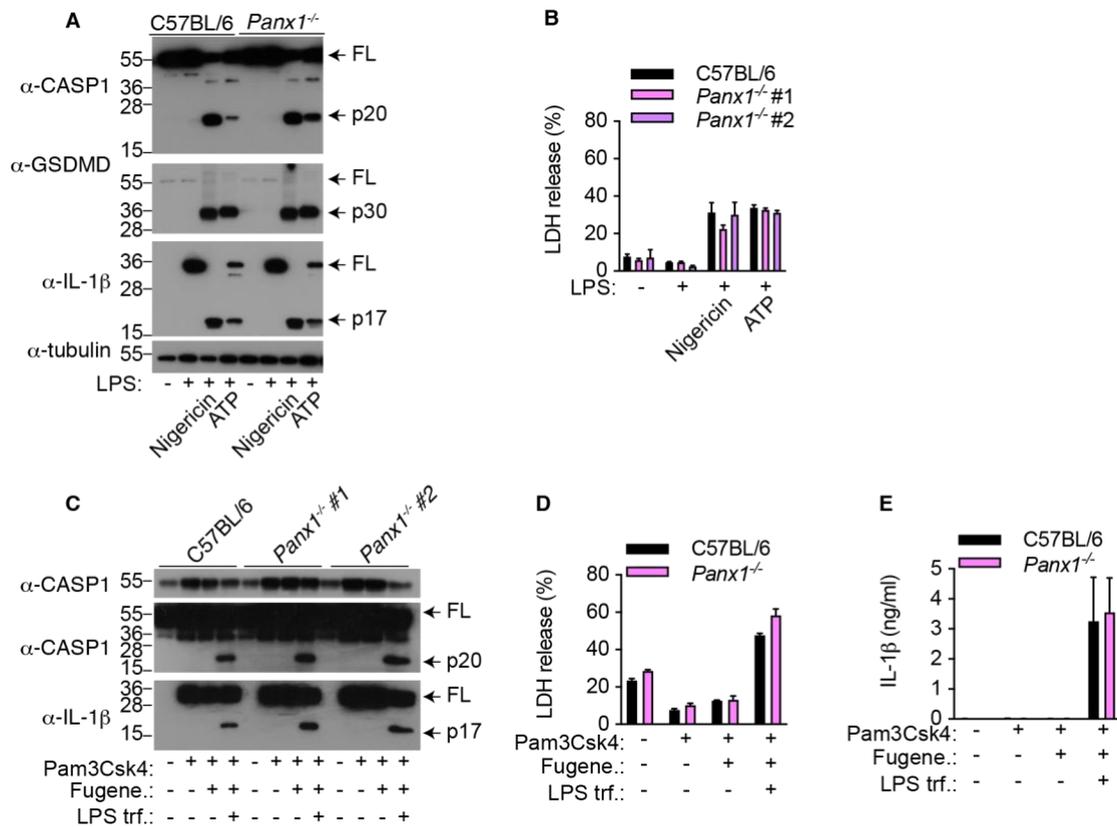


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527 **Figure 3. Non-canonical inflammasome drives pyroptosis and NLRP3**
 528 **activation via GSDMD but not pannexin-1.** (A-G) BMDMs were primed with
 529 1 μ g/ml Pam3CSK4 for 4 h and transfected with 2 μ g/ml ultrapure 0111:B4 LPS
 530 using Fugene for 16 h, or treated stimulated with ABT-737 (0.5 μ M) and S63845
 531 (0.5 μ M) for 5 h. (A, D, G) Mixed supernatant and cell extracts were examined
 532 by immunoblotting and are representative of at 2-3 independent experiments.
 533 (B-C) Data are mean \pm S.E.M of pooled data from 3 independent experiments

534 or (E-F) mean \pm S.D of triplicate cell stimulation representative of 3
 535 independent experiments. Data were considered significant when *P < 0.05,
 536 **P < 0.01.

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541 **Supplementary Figure 1. Pannexin-1 is dispensable for canonical or non-**
 542 **canonical inflammasome signalling.** (A-B) BMDMs were primed with 100
 543 ng/ml ultrapure LPS for 4 h and stimulated with nigericin (0.5 μ M; 90 min) or
 544 ATP (2.5 mM; 90 min). (C-E) BMDMs were primed with 1 μ g/ml Pam3CSK4 for
 545 4 h and transfected with 2 μ g/ml ultrapure 0111:B4 LPS using Fugene for 16 h.
 546 (B, D, E) Data are mean \pm S.D of triplicate cell stimulation representative of 2-
 547 3 independent experiments.