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Abstract

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Keywords

oxygen, reactive, induces, activation, species, receptor, cells, p2x7, erythroid, formation, CMMB

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P2X7 receptor activation induces reactive oxygen species formation in erythroid cells

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Abstract

The presence of P2X7 on erythroid cells is well established, but its physiological role remains unclear. The current study aimed to determine if P2X7 activation induces reactive oxygen species (ROS) formation in murine erythroleukaemia (MEL) cells, a commonly used erythroid cell line. ATP induced ROS formation in a time- and concentration-dependent fashion. The most potent P2X7 agonist, 2'(3')-O-(4-benzoylbenzoyl)ATP, but not UTP or ADP, also induced ROS formation. The P2X7 antagonist, A-438079, impaired ATP-induced ROS formation. The ROS scavenger, *N*-acetyl-L-cysteine, and the ROS inhibitor, diphenyleneiodonium, also impaired P2X7-induced ROS formation, but use of enzyme-specific ROS inhibitors failed to identify the intracellular source of P2X7-induced ROS formation. P2X7-induced ROS formation was impaired partly by physiological concentrations of Ca^{2+} and Mg^{2+} , and almost completely in cells in *N*-methyl-D-glucamine chloride medium. The p38 MAPK inhibitors SB202190 and SB203580, and the caspase inhibitor Z-VAD-FMK, but not *N*-acetyl-L-cysteine, impaired P2X7-induced MEL cell apoptosis. ATP also stimulated p38 MAPK and caspase activation, both of which could be impaired by A-438079. In conclusion, these findings indicate that P2X7 activation induces ROS formation in MEL cells and that this process may be involved in events downstream of P2X7 activation, other than apoptosis, in erythroid cells.

Keywords: P2X7; extracellular ATP; red blood cell; reactive oxygen species; apoptosis

Introduction

The P2X7 receptor is a trimeric ATP-gated cation channel that mediates the influx of extracellular Ca^{2+} and Na^+ , and efflux of intracellular K^+ , as well as the uptake of organic cations including *N*-methyl-D-glucamine⁺ (NMDG⁺) and fluorescent dyes such as ethidium⁺ [1]. P2X7 is expressed on haematopoietic, epithelial, bone and neuronal cells, where it functions in inflammation and immunity [2], and cellular homeostasis [3]. Due to these and other properties of P2X7 activation, this receptor plays important roles in health and disease [4]. The various biological roles attributed to P2X7 are the result of various signalling events downstream of P2X7 activation including reactive oxygen species (ROS) formation [5]. P2X7 activation induces ROS generation in a variety of cell types resulting in transcription factor activation [6], pro-inflammatory cytokine release [7, 8], microbial killing [9], cell death [10, 11] and autophagy [12]. However, it remains unknown if P2X7 activation can induce ROS formation in erythroid cells.

The presence of P2X7 on erythroid cells is well established, but its physiological role remains unclear. P2X7 is present on mature red blood cells (erythrocytes) [13, 14] where it participates in cation fluxes [13, 15-17], cell death [16, 18-20] and the release of lipid-derived mediators [21]. P2X7 is also present on erythroid precursors including murine erythroleukaemia (MEL) cells [22, 23]. Our group directly demonstrated that MEL cells express P2X7, and that activation of this receptor induces the uptake of organic cations, rapid phosphatidylserine exposure, microparticle release and apoptosis in these cells [23]. In the current study, we demonstrate that P2X7 activation also induces ROS formation in MEL cells,

supporting a potential role for P2X7-induced ROS formation in intracellular signalling or oxidative stress in erythroid cells.

Material and methods

Reagents RPMI-1640 medium, L-glutamine, gentamicin and 2',7'-dichlorodihydrofluorescein diacetate (H₂DCFDA) were from Invitrogen (Grand Island, NJ, USA). FCS (heat inactivated before use) was from Lonza (Basel, Switzerland) or Bovogen Biologicals (East Kellior, Australia). ATP, 2'(3')-O-(4-benzoylbenzoyl)ATP (BzATP), ADP, UTP, rotenone, allopurinol, N_ω-nitro-L-arginine methyl ester hydrochloride (L-NAME), EGTA and BAPTA-AM were from Sigma Chemical Co (St Louis, MO, USA). A-438079 was from Tocris Bioscience (Ellisville, MO, USA). N-acetyl-L-cysteine (NAC) and 7-aminoactinomycin D (7AAD) were from Alexis Biochemicals (Lausen, Switzerland). Diphenyleneiodonium (DPI) was from Cayman Chemical Company (Ann Arbor, MI, USA). Ethidium bromide was from Amresco (Solon, OH, USA). Apocynin and SB202190 were from Calbiochem (San Diego, CA, USA). Annexin-V-FLUOS and DNase I were from Roche Applied Science (Pensberg, Germany). SB203580 and Z-VAD-FMK were from Jena Bioscience (Jena, Germany) and BioVision (Mountain View, CA, USA), respectively.

Cell culture MEL cells were maintained in complete culture medium (RPMI-1640 medium containing 10% FCS, 2 mM L-glutamine and 5 ng/ml gentamicin) as described [23].

ROS formation measurements ATP-induced ROS formation was assessed in MEL cells loaded with H₂DCFDA, which is converted to the fluorescent compound 2',7'-dichlorofluorescein (DCF) upon oxidation by ROS. Briefly, MEL cells in NaCl medium (145 mM NaCl, 5 mM KCl, 5 mM D-glucose, 10 mM HEPES, pH 7.4) (1×10^6 cells/ml) were incubated with 5 μ M H₂DCFDA for 5 min at 37°C. Cells were centrifuged and washed once with NaCl medium. H₂DCFDA-loaded cells in NaCl medium were incubated in the absence or presence of nucleotides (as indicated) at 37°C. Incubations were stopped by addition of an equal volume of ice-cold MgCl₂ medium (145 mM NaCl, 5 mM KCl, 20 mM MgCl₂, 10 mM HEPES, pH 7.4) and centrifugation. Cells were washed once with NaCl medium. The mean fluorescence intensity (MFI) of DCF fluorescence (ROS formation) was determined using a BD (San Diego, CA, USA) LSR II flow cytometer and FlowJo software (Tree Star, Ashland, OR, USA).

In some experiments, ATP-induced ROS formation was assessed in cells pre-incubated in the absence or presence of different compounds (as indicated) at 37°C before the addition of ATP. In other experiments, ATP-induced ROS formation was assessed in cells in NaCl medium containing 2 mM CaCl₂ and 1 mM MgCl₂, KCl medium (150 mM KCl, 5 mM D-glucose, 10 mM HEPES, pH 7.4), *N*-methyl-D-glucamine chloride (NMDG Cl) medium (145 mM NMDG Cl, 5 mM KCl, 5 mM D-glucose, 10 mM HEPES, pH 7.4), complete culture medium, or in NaCl medium containing either 1 mM CaCl₂, 100 μ M EGTA or 1 mM Mg²⁺. Since free Ca²⁺ or Mg²⁺ can lower the concentration of ATP⁴⁻, cells in the presence of 1 mM Ca²⁺ or 1 mM Mg²⁺ in some experiments were incubated with 1.54 or 1.74 mM ATP, respectively to provide equimolar ATP⁴⁻ concentrations (415 μ M) as calculated using the

Bound and Determined Program [24].

Ethidium⁺ uptake measurements P2X7-induced pore formation in MEL cells suspended in NaCl medium was assessed using a fixed-time ethidium⁺ uptake assay as described [23]. The MFI of ethidium⁺ uptake (pore formation) was determined by flow cytometry. In some experiments, ATP-induced ethidium⁺ uptake was assessed in cells pre-incubated with different compounds or incubated in different media as above (for ROS formation).

Apoptosis measurements P2X7-induced apoptosis of MEL cells was performed as described [23]. Briefly, cells were pre-incubated in the absence or presence of different compounds (as indicated) at 37°C, followed by incubation in the absence or presence of 1 mM ATP for 24 h. Following ATP incubation, the percentage of Annexin-V⁺/7AAD⁻ (early apoptotic) and Annexin-V⁺/7AAD⁺ (late apoptotic) cells was determined by flow cytometry.

p38 MAPK activation measurements MEL cells in complete culture medium in 24-well plates (5×10^5 cells/ml/well) were pre-incubated in the absence or presence of 10 μ M A-438079 for 15 min at 37°C, followed by incubation in the absence or presence of 1 mM ATP for 5 min. Harvested cells were suspended in 100 μ L ice-cold SDS sample buffer containing 5% 2-mercaptoethanol, and sonicated in the presence of DNase I (10 μ g/ml) for 5 min. Proteins were separated by SDS-PAGE (4% stacking and 10% resolving gels) and transferred to nitrocellulose membrane (Bio-Rad, Hercules, CA, USA). Immunoblotting of p38 MAPK and phospho-p38 MAPK was performed using the PhosphoPlus p38 MAPK (Thr180/Tyr182)

Antibody Kit (Cell Signaling Technology, Danvers, MA, USA) according to the manufacturer's instructions.

Caspase activation measurements Caspase activation was assessed using the CaspGLOW Fluorescein Active Caspase Staining Kit (BioVision) according to the manufacturer's instructions. Briefly, MEL cells in complete culture medium in 24-well plates (5×10^5 cells/ml/well) were pre-incubated in the absence or presence of 10 μ M A-438079 for 15 min at 37°C, followed by incubation in the absence or presence of 1 mM ATP for 6 h. Following ATP incubation, cells were incubated with fluorescein isothiocyanate (FITC)-VAD-FMK for 1 h at 37°C; 7AAD was added during the final 5 min of incubation. Cells were harvested and washed twice with Wash Buffer. The percentage of FITC⁺/7AAD⁺ (caspase activated) cells was determined by flow cytometry.

Presentation of data and statistical analysis Results are presented as means \pm SD. Data was compared using one-way analysis of variance (using Tukey's post test) or (when indicated) the unpaired two-tailed Student's *t*-test using Prism 5 (GraphPad Software, San Diego, CA, USA) with $P < 0.05$ considered significant.

Results

P2X7 activation induces ROS formation in MEL cells To determine whether P2X7 activation can induce ROS formation in MEL cells, cells were loaded with with H₂DCFDA, which is

converted to the fluorescent compound DCF upon oxidation by ROS. Cells in NaCl medium (nominally free of Ca^{2+} and Mg^{2+}) were then incubated in the absence or presence of ATP for up to 30 mins, and the amount of ROS formation determined by flow cytometry. ATP induced ROS formation in MEL cells in a time-dependent fashion (Fig. 1A). Moreover, ATP induced ROS formation in a concentration-dependent fashion, with a maximal response at 2 mM ATP and an EC_{50} of 151 μM (95% confidence interval of 120-190 μM) (Fig. 1B). This is similar to the EC_{50} for ATP-induced ethidium⁺ uptake in MEL cells [23] and for cation fluxes mediated by recombinant murine P2X7 [25].

To determine further if the ATP-induced ROS formation in MEL cells was mediated by P2X7 activation, cells were incubated in the absence or presence of the most potent P2X7 agonist BzATP, or the non-P2X7 agonists ADP or UTP [25]; although it should be noted that BzATP also activates P2X1-P2X5 [26, 27]. ATP was included as a positive control. Both ATP and BzATP induced significant ROS formation in MEL cells compared to cells incubated in the absence of nucleotide (Fig. 1C). In contrast, ADP or UTP failed to induce ROS formation (Fig. 1C). Finally, to confirm that ATP-induced ROS formation was mediated by P2X7 activation, MEL cells were pre-incubated in the absence or presence of the P2X7 antagonist A-438079, before incubation in the absence or presence of ATP. As above (Fig. 1A-C), ATP induced significant ROS formation in MEL cells compared to cells incubated in the absence of ATP (Fig. 1D). Pre-incubation of MEL cells with 10 μM A-438079 impaired ATP-induced ROS formation by $87 \pm 1\%$ (Fig. 1D). Collectively, these results indicate that extracellular ATP induces ROS formation in MEL cells by P2X7 activation.

The above studies were conducted with MEL cells suspended in NaCl medium

nominally free of Ca^{2+} and Mg^{2+} . Therefore, to assess if ATP could induce ROS formation in MEL cells in medium containing physiological concentrations of divalent cations, MEL cells were suspended in NaCl medium containing 2 mM Ca^{2+} and 1 mM Mg^{2+} , and the ATP-induced ROS formation was assessed. Due to the known inhibitory actions of Ca^{2+} and Mg^{2+} on P2X7 [28, 29], cells were exposed to 1 mM ATP (as above), as well as 2 and 3 mM ATP. ATP induced ROS formation in MEL cells in NaCl medium (containing physiological concentrations of divalent cations) in a concentration-dependent fashion (Fig. 1E).

P2X7-induced ROS formation in MEL cells is impaired by NAC and DPI To confirm that P2X7 activation induced ROS formation in MEL cells, cells in NaCl medium (nominally free of Ca^{2+} and Mg^{2+}) were pre-incubated in the absence or presence of the ROS scavenger NAC, or in the presence of DMSO diluent control or the broad-spectrum ROS inhibitor DPI before incubation in the absence or presence of ATP. As above (Fig. 1), ATP induced significant ROS formation in MEL cells compared to cells incubated in the absence of ATP (Fig. 2A,B). Pre-incubation of MEL cells with 10 mM NAC or 20 μM DPI impaired ATP-induced ROS formation by $70 \pm 7\%$ and $50 \pm 15\%$, respectively (Fig. 2A,B). To determine if NAC or DPI directly affected P2X7, ATP-induced ethidium⁺ uptake was measured in the absence or presence of each compound. Pre-incubation of MEL cells with NAC or DPI (as above) did not alter the amount of ATP-induced ethidium⁺ uptake (Fig. 2A,B).

ROS can be generated from numerous sources within cells. Therefore, in an attempt to identify the intracellular source of ROS generated downstream of P2X7 activation, MEL cells in NaCl medium (nominally free of Ca^{2+} and Mg^{2+}) were pre-incubated in the presence of

diluent control (as indicated), or apocynin, rotenone, allopurinol or L-NAME, which impair NADPH oxidase, mitochondrial complex I, xanthine oxidase or nitric oxide synthase, respectively, before incubation in the absence or presence of ATP. Pre-incubation times and antagonist concentrations were based on previous studies with murine macrophages [10, 11]. Again, ATP induced significant ROS formation in MEL cells compared to cells incubated in the absence of ATP (Fig. 2C-F). Pre-incubation of MEL cells with either 100 μ M apocynin, 5 μ M rotenone, 100 μ M allopurinol or 1 mM L-NAME had no significant effect on ATP-induced ROS formation compared to cells incubated with both the respective diluent control and ATP (Fig. 2C-F). In the absence of ATP, ROS formation in the presence of apocynin, allopurinol or L-NAME was not significantly different to the respective diluent control (Fig. 2C,E,F). In contrast, in the absence of ATP, rotenone induced significant ROS formation compared to DMSO (Fig. 2D). Shorter (45 min) incubations with different concentrations of rotenone in the absence of ATP also induced ROS formation in MEL cells in concentration-dependent fashion, with significant ROS formation observed with a minimum of 0.5 μ M rotenone and with an EC_{50} of >2 μ M (results not shown). While 30 min pre-incubation of MEL cells with 0.1 μ M rotenone potentiated ATP-induced ROS formation compared to cells incubated with both DMSO and ATP (results not shown). Pre-incubation of MEL cells with either apocynin, rotenone, allopurinol or L-NAME (as above) had no significant effect on ATP-induced ethidium⁺ uptake compared to cells incubated with both the respective diluent control and ATP (Fig. 2C-F).

P2X7-induced ROS formation is impaired in MEL cells in NaCl medium containing Ca^{2+} or

Mg^{2+} , or in NMDG Cl medium P2X7 is a trimeric ATP-gated cation channel [1], therefore a series of experiments were undertaken to explore a possible role for divalent or monovalent cations in P2X7-induced ROS formation. For each experiment, cells in different media were compared to cells in NaCl medium (nominally free of Ca^{2+} and Mg^{2+}). P2X7-induced ROS formation is dependent on Ca^{2+} influx in rat, but not murine, submandibular gland cells [30, 31]. Therefore, to determine if P2X7-induced ROS formation in MEL cells is dependent on Ca^{2+} influx, ATP-induced ROS formation was compared in cells in the absence or presence of additional Ca^{2+} . ATP induced significant ROS formation in MEL cells in either the absence or presence of 1 mM Ca^{2+} compared to similarly treated cells incubated in the absence of ATP (Fig. 3A). Of note, ATP-induced ROS formation was significantly lower in the presence of Ca^{2+} compared to ATP-induced ROS formation in the absence of Ca^{2+} (Fig. 3A). This result paralleled observations with ATP-induced ethidium⁺ uptake, with ATP-induced ethidium⁺ uptake significantly lower in the presence of Ca^{2+} compared to ATP-induced ethidium⁺ uptake in the absence of Ca^{2+} (Fig. 3A).

NaCl medium may contain nominal amounts of Ca^{2+} . Therefore, to further exclude a role for Ca^{2+} influx in P2X7-induced ROS formation, ATP-induced ROS formation was compared in MEL cells in the absence or presence the Ca^{2+} chelator EGTA. ATP induced significant ROS formation in MEL cells in either the absence or presence of 100 μ M EGTA compared to similarly treated cells incubated in the absence of ATP (Fig. 3B). Moreover, ATP-induced ethidium⁺ uptake was similar in MEL cells in either the absence or presence of EGTA (Fig. 3B) indicating that neither EGTA nor nominal amounts of extracellular Ca^{2+} had a direct effect on P2X7-induced pore formation.

Next, the potential role of intracellular Ca^{2+} increase in P2X7-induced ROS formation in MEL cells was examined by comparing ATP-induced ROS formation in MEL cells pre-incubated in the presence of DMSO or the intracellular Ca^{2+} chelator BAPTA-AM. ATP induced significant ROS formation in MEL cells in either the presence of DMSO or 10 μM BAPTA-AM compared to similarly treated cells incubated in the absence of ATP (Fig. 3C). Moreover, ATP-induced ethidium⁺ uptake was similar in MEL cells in either the presence of DMSO or BAPTA-AM (Fig. 3C) indicating that neither BAPTA-AM nor intracellular Ca^{2+} increase had a direct effect on P2X7-induced pore formation.

Since the current study predominately assessed P2X7-induced ROS formation in MEL cells in NaCl medium, which may contain nominal amounts of Mg^{2+} , the possibility remained that this process was dependent on Mg^{2+} efflux. Therefore, ATP-induced ROS formation was compared in cells in the absence or presence of 1 mM Mg^{2+} , a concentration similar to that of free intracellular Mg^{2+} . ATP induced significant ROS formation in MEL cells in either the absence or presence of Mg^{2+} compared to similarly treated cells incubated in the absence of ATP (Fig. 3D). As above for Ca^{2+} (Fig. 3A), ATP-induced ROS formation was significantly lower in the presence of Mg^{2+} compared to ATP-induced ROS formation in the absence of Mg^{2+} (Fig. 3D). This result paralleled observations with ATP-induced ethidium⁺ uptake, with ATP-induced ethidium⁺ uptake significantly lower in the presence of Mg^{2+} compared to ATP-induced ethidium⁺ uptake in the absence of Mg^{2+} (Fig. 3D).

Both ROS formation and K^{+} efflux are involved in P2X7-induced interleukin-1 β release from monocytes, although it remains to be determined if these upstream processes are linked [32]. Therefore, to determine if P2X7-induced ROS formation in MEL cells is dependent on

K⁺ efflux, ATP-induced ROS formation was compared in cells in NaCl medium and KCl medium, which prevents the loss of intracellular K⁺. The amount of ATP-induced ROS formation in MEL cells in KCl medium was similar to ROS formation in cells in NaCl medium (Fig. 3E). To determine if the inability of high extracellular K⁺ to impair ATP-induced ROS formation was not due to altered P2X7 function in KCl medium, ATP-induced ethidium⁺ uptake was assessed in cells in both media. ATP-induced ethidium⁺ uptake was similar in MEL cells incubated in either medium (Fig. 3E).

The above results with KCl medium (Fig. 3E) indirectly exclude a role for Na⁺ influx in P2X7-induced ROS formation in MEL cells. It is well known however that high concentrations of extracellular K⁺ will depolarise cells and interfere with various transport systems. Therefore, ATP-induced ROS formation was compared in cells in NaCl medium and NMDG Cl medium, which is nominally free of Na⁺. Both media were nominally free of Ca²⁺ and Mg²⁺. In contrast to observations with KCl medium above (Fig. 3E), ATP-induced ROS formation was impaired by 88 ± 1% in MEL cells in NMDG Cl medium compared to cells in NaCl medium (Fig. 3F). To determine if this inhibition was due to impaired P2X7 function in NMDG Cl medium, ATP-induced ethidium⁺ uptake was assessed in cells in both media. In contrast to ROS formation, ATP-induced ethidium⁺ uptake was approximately two-fold greater in MEL cells incubated in NMDG Cl medium compared to cells in NaCl medium (Fig. 3F).

p38 MAPK and caspase activation, but not ROS formation, mediates P2X7-induced apoptosis of MEL cells Our group previously demonstrated that P2X7 activation induces apoptosis of

MEL cells [23], however the intracellular signalling pathways involved were not determined. ROS formation, as well as the p38 MAPK and caspase pathways are involved in P2X7-induced apoptosis of RAW264.7 macrophages [10]. Therefore, the roles of these pathways in P2X7-induced MEL cell apoptosis were examined. First, cells were pre-incubated in the absence or presence of NAC before incubation in the absence or presence of ATP. As previously observed [23], incubation of MEL cells with 1 mM ATP induced a significant increase in the percentage of Annexin-V⁺/7AAD⁻ (early apoptotic) and Annexin-V⁺ /7AAD⁺ (late apoptotic) cells (Fig. 4A). Pre-incubation of MEL cells with 10 mM NAC failed to impair ATP-induced early or late apoptosis (Fig. 4A). Moreover, incubation of MEL cells in complete culture medium with 1 mM ATP (as used in the apoptosis assay) did not induce significant ROS formation despite 1 mM ATP causing significant ROS formation in MEL cells in NaCl medium (nominally free of Ca²⁺ and Mg²⁺) (Fig. 4B).

Next, cells were pre-incubated in the presence of DMSO or the p38 MAPK inhibitors SB202190 or SB203580 before incubation in the absence or presence of ATP. Again ATP induced a significant increase in early and late apoptotic cells (Fig. 4C). Pre-incubation of MEL cells with 20 μ M SB202190 significantly impaired ATP-induced early and late apoptosis by $56 \pm 5\%$ and $56 \pm 9\%$, respectively (Fig. 4C). Similarly, pre-incubation of MEL cells with 20 μ M SB203580 impaired the amount of ATP-induced early and late apoptosis by $48 \pm 17\%$ and $52 \pm 6\%$, respectively (Fig. 4C).

Next, cells were pre-incubated in the presence of DMSO, or the broad-spectrum caspase inhibitor Z-VAD-FMK before incubation in the absence or presence of ATP. Again ATP

induced a significant increase in early and late apoptotic cells (Fig. 4D). Pre-incubation of MEL cells with 20 μ M Z-VAD-FMK significantly impaired the amount of ATP-induced late apoptotic cells by $36 \pm 2\%$ (Fig. 4D). In contrast, pre-incubation of cells with Z-VAD-FMK failed to impair the amount of early apoptotic cells induced by ATP (Fig. 4D).

p38 MAPK and caspase inhibitors have been reported to impair P2X7-induced pore formation [33, 34]. To exclude the possibility that the above compounds impaired P2X7-induced apoptosis of MEL cells by directly inhibiting P2X7, cells were pre-incubated in the presence of DMSO, SB202190, SB203580 or Z-VAD-FMK (as above), and the P2X7-induced ethidium⁺ uptake assessed. ATP induced significant ethidium⁺ uptake into MEL cells compared to cells incubated in the absence of ATP (Fig. 4E). Pre-incubation of MEL cells with any of the compounds failed to alter the amount of ATP-induced ethidium⁺ uptake (Fig. 4E). Collectively, these results indicate that p38 MAPK and caspase activation, but not ROS formation, are involved in P2X7-induced apoptosis of MEL cells.

P2X7 activation induces p38 MAPK and caspase activation in MEL cells To support the role of p38 MAPK activation in the P2X7-induced apoptosis of MEL cells, ATP-induced p38 MAPK phosphorylation was examined by immunoblotting. Immunoblotting with an anti-phospho-p38 MAPK antibody revealed that incubation of cells with ATP for 5 min induced greater amounts of p38 MAPK activation compared to cells incubated in the absence of ATP (Fig. 5A). Moreover, pre-incubation of cells with the P2X7 antagonist A-438079 (10 μ M) impaired the amount ATP-induced p38 activation compared to ATP alone (Fig. 5A).

To support the role of caspase activation in the P2X7-induced apoptosis of MEL cells,

ATP-induced caspase activity was examined using a FITC-conjugated caspase substrate and flow cytometry. MEL cells were incubated in the absence or presence of ATP for 6 h, and then for a further 1 h with FITC-VAD-FMK and the percentage of FITC⁺/7AAD⁺ cells (caspase activation) measured by flow cytometry. Incubation of cells with ATP resulted in a 2.6-fold increase in the relative percentage of cells with activated caspases compared to cells incubated in the absence of ATP (Fig. 5B). Pre-incubation of cells with 10 μ M A-438079 impaired the amount ATP-induced caspase activation by $95 \pm 9\%$ compared to cells incubated with ATP alone (Fig. 5B). Collectively, these results indicate that P2X7 activation stimulates p38 MAPK and caspases, and indirectly supports a role for these enzymes in the P2X7-induced apoptosis of MEL cells.

Discussion

The current study demonstrates that P2X7 activation results in the formation of ROS in MEL cells, an erythroid cell line. P2X7-induced ROS formation has been previously observed in myeloid cells [6-12, 35], glomerular mesangial cells [36] and submandibular gland cells [30, 31], but to the best of our knowledge the current study is the first to demonstrate that P2X7 activation causes ROS formation in erythroid cells. It will be of future importance to determine if P2X7 induces ROS formation in primary erythroid cells. Moreover, it will be important to identify the intracellular source of ROS formation downstream of P2X7 activation in erythroid cells. In the current study, use of apocynin, rotenone, allopurinol and L-NAME, which impair NADPH oxidase, mitochondrial complex I, xanthine oxidase and

nitric oxide synthase, respectively, failed to identify the intracellular source of P2X7-induced ROS formation in MEL cells; however the mitochondrial respiratory chain cannot be excluded as a source. Although rotenone is commonly used as an inhibitor of mitochondrial ROS formation, it has been widely reported to also induce mitochondrial ROS formation [37]. Consistent with the latter, rotenone repeatedly increased basal ROS formation in MEL cells, making it difficult to ascertain if mitochondria were the source of P2X7-induced ROS formation. Moreover, DPI can impair mitochondrial ROS formation, as well as NADPH oxidase ROS formation [38], supporting a possible role for mitochondria in P2X7-induced ROS formation. The failure of apocynin to impair P2X7-induced ROS formation excludes a role for NADPH oxidase in this process; however further studies using gene knockdown approaches are required to support this conclusion. Nevertheless based on the current results, the mitochondrial respiratory chain or enzyme systems other than those examined in the current study remain the most likely sources of P2X7-induced ROS formation in erythroid cells.

The physiological role of P2X7-induced ROS formation in erythroid cells remains to be established. Accumulating evidence suggests that ROS are involved in cell homeostasis, proliferation, differentiation and death [39, 40]. Specifically in relation to erythroid cells, low amounts of ROS are involved in the differentiation of erythroid progenitors, through interactions with FoxO3 and other transcription factors [41]. Whereas, excess amounts of ROS may shorten the life-span of mature erythroid cells [41]. In the current study however, the presence of the ROS scavenger NAC failed to impair P2X7-induced apoptosis of MEL cells, suggesting that P2X7-induced ROS formation is more likely to be involved in

signalling events related to erythroid cell homeostasis, proliferation or differentiation rather than erythroid cell death. An alternative, but not mutually exclusive, concept is that P2X7-induced ROS formation may result in phosphatidylserine exposure (in the absence of immediate cell death) and the subsequent removal of erythroid precursors from the circulation in certain disease states such as thalassaemia. In this regard, thalassaemic DNA-containing reticulocytes and erythrocytes have a higher content of ROS and exposed phosphatidylserine compared to DNA-free counterparts, leading the authors to postulate that oxidative stress-induced phosphatidylserine exposure is involved in the clearance of these cells from the circulation [42].

The current study demonstrates that Na^+ influx but not K^+ efflux may have a role in P2X7-induced ROS formation in MEL cells. NMDG Cl medium, which is nominally free of Na^+ , almost completely impaired P2X7-induced ROS formation, however KCl medium, which is also nominally free of Na^+ , did not. These conflicting results may be explained by at least two, but not mutually exclusive, possibilities. First, since P2X7 activation can induce NMDG⁺ uptake [43], the possibility remains that P2X7-induced ROS formation is impaired by increased intracellular NMDG⁺ rather than through the absence of Na^+ influx. Second, since high concentrations of extracellular K^+ can depolarise cells and interfere with various transport systems, the possibility remains that under these (KCl medium) but not other conditions (NaCl medium) that Na^+ influx is not essential for P2X7-induced ROS formation. Reports for a role for Na^+ influx in P2X7-induced signalling events are few, however others have shown that Na^+ influx is involved in P2X7-induced rapid exposure of phosphatidylserine in murine thymocytes [44] and P2X7-induced mitochondrial membrane

depolarisation in rat submandibular glands [45]. Thus, the current study may offer another example of the importance of Na^+ influx in a P2X7-induced signalling event. In contrast, this study indicates that K^+ efflux is not essential for P2X7-induced ROS formation in MEL cells. Both ROS formation and K^+ efflux are involved in P2X7-induced interleukin-1 β from monocytes, although it remains unknown if these upstream processes are linked [32]. The absence of a role for K^+ efflux in P2X7-induced ROS formation in MEL cells suggests that K^+ efflux and ROS formation may be involved in distinct pathways in monocyte interleukin-1 β release downstream of P2X7 activation.

The current study also demonstrates that Ca^{2+} influx or Mg^+ efflux, or an increase in intracellular Ca^{2+} is not essential for P2X7-induced ROS formation in MEL cells. Others have shown that P2X7-induced ROS formation is dependent on Ca^{2+} influx in rat, but not murine, submandibular gland cells [30, 31]. Thus, the current study with a murine erythroid cell line parallel those of murine submandibular gland cells. Of note however, 1 mM extracellular Ca^{2+} or Mg^{2+} , but not nominal concentrations of extracellular Ca^{2+} (as observed with the use of EGTA), impaired both P2X7-induced ROS formation and pore formation in MEL cells despite the presence of equivalent ATP^{4-} concentrations. This suggests that mM (or perhaps μM) concentrations of extracellular Ca^{2+} and Mg^{2+} may act as allosteric inhibitors of murine P2X7 as originally observed for rat P2X7 [46], and established by more recent studies of rat P2X7 [28, 29]. The current study also demonstrates that an increase in intracellular Ca^{2+} is not essential for P2X7-induced ROS formation (and pore formation) in MEL cells. To the best of our knowledge the potential role of intracellular Ca^{2+} on P2X7-induced ROS formation in any cell type has not been reported, although others have

excluded a role for an increase in intracellular Ca^{2+} in ATP-induced organic cation uptake into RAW264.7 macrophages and HEK-293 cells transfected with rat P2X7 [47].

The current study confirms that P2X7 activation induces apoptosis of MEL cells, and demonstrates for the first time that this process involves activation of p38 MAPK and caspases, but not ROS formation. The involvement of p38 MAPK and caspase activation in P2X7-induced MEL cell apoptosis parallels observations with P2X7-induced apoptosis of RAW264.7 cells [10]. In contrast, this previous study also demonstrated a role for ROS formation in RAW264.7 cell apoptosis, which was not required for MEL cell apoptosis in the current study. This difference was not due to impaired scavenger action of NAC, as this compound impaired P2X7-induced ROS formation in MEL cells. Collectively, the above indicates that p38 MAPK and caspase activation, and ROS formation are independent processes downstream of P2X7 activation in MEL cells. Although the current study demonstrated that caspase activation mediates the P2X7-induced apoptosis of MEL cells, the identity of the caspases involved remains unknown. Others have shown that high pressure-induced apoptosis of MEL cells involves both the extrinsic and intrinsic pathways mediated by caspase-8 and caspase-9, respectively, as well as the downstream activation of caspase-3 [48, 49]. Thus, it is likely that either or both caspase-8 and caspase-9, as well as caspase-3 are involved in P2X7-induced MEL cell apoptosis.

The physiological role of P2X7-induced apoptosis of erythroid cells, as well as other cell types remains poorly defined. Studies of P2X7-deficient mice suggest that the P2X7 is not involved in regulating cell numbers under normal, physiological conditions [50, 51], except in bone homeostasis [52] and the peripheral T cell compartment [53]. Nevertheless

P2X7-induced apoptosis remains a well-described event [2, 3]. Therefore, it is likely that P2X7-induced apoptosis may play a role in removing damaged erythroid cells to prevent the development of anaemia or leukaemia [54], or to promote their rapid clearance from the circulation to prevent autoimmunity [55]. Alternatively, errant ATP release and subsequent P2X7-induced apoptosis of erythroid progenitors may be associated with the premature loss of these cells to cause anaemia [54]. This may be particularly relevant to chemotherapy, which can cause the release of ATP from cells [56] and induce cytotoxicity in immature erythroid progenitors to cause anaemia in cancer patients [57].

Finally, the current study demonstrates that neither the ROS scavenger (NAC) nor any of the ROS inhibitors (DPI, apocynin, rotenone, allopurinol and L-NAME) had a direct effect on P2X7 itself. The inability of these compounds to impair P2X7-induced pore formation in MEL cells is consistent with similar findings in murine J774 cells and primary murine macrophages [11]. Thus, these observations validate the current and future use of these compounds to study ROS formation downstream of P2X7 activation, and indicate that ROS formation does not modulate P2X7-induced pore formation. In contrast, P2X7 in some cell types or studies can be directly impaired by a number of enzyme inhibitors [33, 34, 58-60]. Of relevance to the current study, the p38 MAPK inhibitors SB202190 or SB203580 can impair P2X7-induced pore formation in human THP-1 myeloid cells, murine 2BH4 thymic epithelial cells and primary murine macrophages [33, 34]. Although others have demonstrated that these compounds impair ATP-induced pore formation mediated by human, but not rat or murine, recombinant P2X7 [61]. This latter finding is consistent with the absence of an inhibitory effect of SB202190 or SB203580 on P2X7-induced pore formation

in MEL cells. The caspase-1 and caspase-3 inhibitors YVAD and DEVD, respectively have also been shown to impair P2X7-induced pore formation in THP-1 cells [33]. However, the broad-spectrum caspase inhibitor Z-VAD-FMK failed to impair P2X7-induced pore formation in MEL cells.

In conclusion, these findings indicate that P2X7 activation induces ROS formation in MEL cells and that this process may be involved in events downstream of P2X7 activation, other than apoptosis, in erythroid cells. This study highlights the potential role of P2X7 activation in intracellular signalling pathways or oxidative stress in erythroid cells.

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Figure legends

Fig. 1 P2X7 activation induces ROS formation in MEL cells. H₂DCFDA-loaded MEL cells in (A-D) NaCl medium or (E) NaCl medium (containing 2 mM Ca²⁺ and 1 mM Mg²⁺) were incubated at 37°C in the absence (Basal) or presence of (A) 1 mM ATP for up to 30 min, (B) varying ATP concentrations for 15 min, (C) 1 mM ATP, UTP or ADP, or 200 μM BzATP for 15 min, (D) 10 μM A-438079 for 15 min, and then in the absence or presence of 1 mM ATP for 15 min, or (E) 1, 2 or 3 mM ATP for 15 min. (A-E) Incubations were stopped by addition of MgCl₂ medium and centrifugation, and the mean fluorescence intensity (MFI) of DCF (ROS formation) was determined by flow cytometry. ROS formation is expressed as (A and C-E) MFI or (B) percentage of maximum response compared to 2 mM ATP. Results are mean ± SD (A and C-E, *n* = 3; B, *n* = 6); where the SD is not visible, SD is ≤ 2.6; **P* < 0.05, ***P* < 0.01 compared to corresponding Basal; ††*P* < 0.01 compared to ATP alone (C and D, ANOVA; E, Student's *t*-test).

Fig. 2 P2X7-induced ROS formation in MEL cells is impaired by NAC and DPI. H₂DCFDA-loaded MEL cells (*left panels*) or MEL cells (*right panels*) in NaCl medium were pre-incubated at 37°C in the (A, F) absence (Control), or presence of (A) 10 mM NAC for 30 min or (F) 1 mM L-NAME for 60 min, or in the presence of (B) DMSO or 20 μM DPI for 30 mins, (C) DMSO or 100 μM apocynin for 60 mins, (D) DMSO or 5 μM rotenone for 60 mins, or (E) NaOH or 100 μM allopurinol for 30 mins. (A-F) Cells were then incubated in the absence (Basal) or presence of 1 mM ATP at 37°C for 15 min; 25 μM ethidium⁺ was also

present (*right panels*). Incubations were stopped by addition of MgCl_2 medium and centrifugation, and the mean fluorescence intensity (MFI) of DCF (ROS formation; *left panels*) or ethidium⁺ uptake (pore formation; *right panels*) analysed by flow cytometry. Results are mean \pm SD ($n = 3$); * $P < 0.05$, ** $P < 0.01$ compared to corresponding Basal; $^{\dagger\dagger}P < 0.01$ compared to ATP without NAC or antagonist; $^{\dagger\dagger\dagger}P < 0.01$ compared to Basal control.

Fig. 3 P2X7-induced ROS formation is impaired in MEL cells in NaCl medium containing Ca^{2+} or Mg^{2+} , or in NMDG Cl medium. H_2DCFDA -loaded MEL cells (*left panels*) or MEL cells (*right panels*) in (A, B and D) NaCl medium in the absence (Control) or presence of (A) 1 mM Ca^{2+} , (B) 100 μM EGTA or (D) 1 mM Mg^{2+} , (C) NaCl medium in the presence of DMSO or 10 μM BAPTA-AM (pre-incubated at 37°C for 30 min), (E) NaCl or KCl medium, or (F) NaCl or NMDG Cl medium were incubated in the absence (Basal) or presence of 415 μM ATP⁴⁻ (1, 1.54 or 1.74 mM ATP as explained in Materials and methods) at 37°C for 15 min; 25 μM ethidium⁺ was also present (*right panels*). (A-F) Incubations were stopped by addition of MgCl_2 medium and centrifugation, and the mean fluorescence intensity (MFI) of DCF (ROS formation; *left panels*) or ethidium⁺ uptake (pore formation; *right panels*) analysed by flow cytometry. Results are mean \pm SD ($n = 3$); * $P < 0.05$, ** $P < 0.01$ compared to corresponding Basal; $^{\dagger}P < 0.05$, $^{\dagger\dagger}P < 0.01$ compared to ATP alone; $^{\dagger\dagger\dagger}P < 0.01$ compared to Basal control.

Fig. 4 p38 MAPK and caspase activation, but not ROS formation, mediates P2X7-induced apoptosis in MEL cells. (A, C and D) MEL cells in complete culture medium were

pre-incubated at 37°C for 30 min (A) in the absence (Control) or presence of 10 mM NAC, in the presence of (C) DMSO, 20 µM SB202190 or 20 µM SB203580, or (D) DMSO or 20 µM Z-VAD-FMK. Cells were then incubated in the absence (Basal) or presence of 1 mM ATP at 37°C for 24 h. Cells were harvested, and labelled with Annexin-V-FLUOS and 7AAD, and the percentage of Annexin-V⁺/7AAD⁻ (early apoptotic; *left panels*) or Annexin-V⁺/7AAD⁺ (late apoptotic; *right panels*) cells analysed by flow cytometry. (B) H₂DCFDA-loaded MEL cells in NaCl medium or complete culture medium (CC) were incubated in the absence (Basal) or presence of 1 mM ATP at 37°C for 15 min. Incubations were stopped by addition of MgCl₂ medium and centrifugation, and the mean fluorescence intensity (MFI) of DCF (ROS formation) was determined by flow cytometry. (E) MEL cells in NaCl medium were pre-incubated at 37°C for 30 min in the presence of DMSO or 20 µM SB202190 (*left panel*), DMSO or 20 µM SB203580 (*centre panel*), or DMSO or 20 µM Z-VAD-FMK (*right panel*). Ethidium⁺ (25 µM) was added and cells were incubated in the absence (Basal) or presence of 1 mM ATP at 37°C for 15 min. Incubations were stopped by addition of MgCl₂ medium and centrifugation, and the mean fluorescence intensity (MFI) of ethidium⁺ uptake (pore formation) analysed by flow cytometry. (A-E) Results are mean ± SD (*n* = 3); **P* < 0.05, ***P* < 0.01 compared to corresponding Basal; †*P* < 0.05, ††*P* < 0.01 compared to ATP alone.

Fig. 5 P2X7 activation induces p38 MAPK and caspase activation in MEL cells. (A) MEL cells in complete culture medium were pre-incubated in the absence or presence of 10 µM A-438079 at 37°C for 15 min, and then in the absence or presence of 1 mM ATP at 37°C for 5 min. Whole lysates of MEL cells were then separated by SDS-PAGE, transferred to

nitrocellulose membrane and probed with (*top panel*) anti-phospho-p38 (p-p38) and (*bottom panel*) anti-p38 (p38) antibodies. (B) MEL cells in complete culture medium were pre-incubated in the absence (Control) or presence of 10 μ M A-438079 at 37°C for 15 min, and then in the absence (Basal) or presence of 1 mM ATP at 37°C for 6 h. Cells were then incubated with FITC-VAD-FMK at 37°C for 1 h and 7AAD for the final 5 min, and the percentage of FITC⁺/7AAD⁺ (caspase activated) cells analysed by flow cytometry. Results are mean \pm SD ($n = 3$); ** $P < 0.01$ compared to corresponding Basal; $^{\dagger\dagger}P < 0.01$ compared to ATP alone.