

Distribution and Ca²⁺ signalling of fibroblast-like (PDGFR α ⁺) cells in the murine gastric fundus

Salah A. Baker, Grant W. Hennig, Anna K. Salter, Masaki Kurahashi, Sean M. Ward and Kenton M. Sanders

Department of Physiology and Cell Biology, University of Nevada School of Medicine, Reno, NV 89557, USA

Key points

- A new class of interstitial cells, PDGFR α ⁺ cells, is distributed densely in the proximal stomachs of mice.
- PDGFR α ⁺ cells express the molecular apparatus necessary for transduction of inputs from enteric inhibitory motor neurons.
- PDGFR α ⁺ cells generate spontaneous Ca²⁺ transients and display dynamic Ca²⁺ oscillations in response to purines.
- Purinergic responses are mediated by P2Y1 receptors and by Ca²⁺ release from intracellular Ca²⁺ stores.
- Ca²⁺ release in PDGFR α ⁺ cells is the likely means by which purinergic neurotransmitters activate Ca²⁺-activated K⁺ channels (SK) and hyperpolarization in GI muscles to elicit inhibitory motor responses.
- Spontaneous Ca²⁺ transients may be a means of regulating basal excitability of fundus muscles and release of purines from motor neurons may contribute to the control of pressure during filling in the proximal stomach.

Abstract Platelet-derived growth factor receptor α positive (PDGFR α ⁺) cells are suggested to mediate purinergic inputs in GI muscles, but the responsiveness of these cells to purines *in situ* has not been evaluated. We developed techniques to label and visualize PDGFR α ⁺ cells in murine gastric fundus, load cells with Ca²⁺ indicators, and follow their activity via digital imaging. Immunolabelling demonstrated a high density of PDGFR α ⁺ cells in the fundus. Cells were isolated and purified by fluorescence-activated cell sorting (FACS) using endogenous expression of enhanced green fluorescent protein (eGFP) driven off the *Pdgfra* promoter. Quantitative PCR showed high levels of expression of purinergic P2Y1 receptors and SK3 K⁺ channels in PDGFR α ⁺ cells. Ca²⁺ imaging was used to characterize spontaneous Ca²⁺ transients and responses to purines in PDGFR α ⁺ cells *in situ*. ATP, ADP, UTP and β -NAD elicited robust Ca²⁺ transients in PDGFR α ⁺ cells. Ca²⁺ transients were also elicited by the P2Y1-specific agonist (*N*)-methanocarba-2MeSADP (MRS-2365), and inhibited by MRS-2500, a P2Y1-specific antagonist. Responses to ADP, MRS-2365 and β -NAD were absent in PDGFR α ⁺ cells from *P2ry1*^(-/-) mice, but responses to ATP were retained. Purine-evoked Ca²⁺ transients were mediated through Ca²⁺ release mechanisms. Inhibitors of phospholipase C (U-73122), IP3 (2-APB), ryanodine receptors (Ryanodine) and SERCA pump (cyclopiazonic acid and thapsigargin) abolished Ca²⁺ transients elicited by purines. This study provides a link between purine binding to P2Y1 receptors and activation of SK3 channels in PDGFR α ⁺ cells. Activation of Ca²⁺ release is likely to be the signalling mechanism in PDGFR α ⁺ cells responsible for the transduction of purinergic enteric inhibitory input in gastric fundus muscles.

(Received 9 September 2013; accepted after revision 15 October 2013; first published online 21 October 2013)

Corresponding author K. M. Sanders: Department of Physiology and Cell Biology, University of Nevada School of Medicine, MS 352, Reno, NV 89557, USA. Email: ksanders@medicine.nevada.edu

Abbreviations: 2-APB, 2-aminoethyl diphenylborinate; c , number of cells; CM, circular muscle; CPA, cyclopiazonic acid; cpm, cycles per minute; eGFP, enhanced green fluorescent protein; FACS, fluorescence-activated cell sorting; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; GI, gastrointestinal; ICC, interstitial cells of Cajal; IJP, inhibitory junction potential; IM, intramuscular; IP₃, inositol 1,4,5-trisphosphate; KRB, Krebs–Ringer bicarbonate; LM, longitudinal muscle; MY, myenteric plexus; β -NAD, β -nicotinamide adenine dinucleotide; L-NNA, N^{ω} -nitro-L-arginine; nNOS, neuronal nitric oxide synthase; PDGFR α , platelet-derived growth factor receptor α ; PGP 9.5, protein gene product 9.5; PLC, phospholipase C; P2Y1, purinergic receptor subtype; ROI, region of interest; SERCA, sarcoendoplasmic reticulum (SR) Ca²⁺ ATPase; SIP, SMC/ICC/PDGFR α ⁺; SK3, small conductance Ca²⁺-activated K⁺ channel; SMC, smooth muscle cell; TTX, tetrodotoxin; UTP, uridine 5'-triphosphate; vAChT, vesicular acetylcholine transporter.

Introduction

Superimposed upon myogenic control in the gastrointestinal (GI) tract are a variety of hierarchical regulatory systems that generate the coordinated muscular movements of normal GI motility. Smooth muscle cells (SMCs), for example, are coupled via gap junctions to at least two distinct classes of interstitial cells, interstitial cells of Cajal (ICC) and PDGFR α ⁺ cells (Komuro *et al.* 1999; Fujita *et al.* 2003). Thus, these three cell types form an electrical syncytium we have referred to as the SMC/ICC/PDGFR α ⁺ (SIP) syncytium (Sanders *et al.* 2012). Inward and outward conductances in any of the SIP cells modulate to overall muscle excitability and responses to other regulatory inputs. ICCs serve as pacemaker cells (Ward *et al.* 1994; Torihashi *et al.* 1995) and mediate and integrate inputs from motor neurons (Burns *et al.* 1996; Ward *et al.* 1998, 2000; Ward & Sanders, 2006). It was recently shown that PDGFR α ⁺ cells are likely to mediate purinergic inputs from enteric inhibitory motor neurons (Kurahashi *et al.* 2011).

PDGFR α ⁺ cells share similar anatomical distributions with ICC, and the study of these cells was advanced when it was shown that antibodies to PDGFR α label cells formerly referred to generically as 'fibroblast-like cells' (Iino *et al.* 2009; Iino & Nojyo, 2009; Sanders, 2010; Kurahashi *et al.* 2011). ICC and PDGFR α ⁺ cells share a similar mesenchymal origin, but they form distinct populations of mature cells based on ultrastructural properties, morphology, expression of specific proteins and function (Komuro *et al.* 1999; Horiguchi & Komuro, 2000; Iino & Nojyo, 2009; Kurahashi *et al.* 2011). Distributions of PDGFR α ⁺ cells in the tunica muscularis have been described in several GI regions of laboratory animals and humans, including the colon, small intestine and sphincters (Iino *et al.* 2009; Iino & Nojyo, 2009; Cobine *et al.* 2011; Kurahashi *et al.* 2011, 2012; Blair *et al.* 2012; Grover *et al.* 2012), and double labelling immunohistochemistry has shown these cells to be closely

associated with enteric motor neurons (Kurahashi *et al.* 2011, 2012; Blair *et al.* 2012). Their close associations with nerve terminals suggest these cells, like ICC, might receive and transduce neurotransmitter input from enteric motor neurons (Komuro, 1999; Horiguchi & Komuro, 2000; Iino & Nojyo, 2009; Kurahashi *et al.* 2011).

PDGFR α ⁺ cells have abundant expression of small conductance Ca²⁺-activated K⁺ channels (SK3 channels) and P2Y1 receptors (Klemm & Lang, 2002; Vanderwinden *et al.* 2002; Fujita *et al.* 2003; Iino *et al.* 2009; Iino & Nojyo, 2009; Kurahashi *et al.* 2011, 2012), which are the major receptors and effectors mediating purinergic enteric inhibitory regulation of GI muscles (Gallego *et al.* 2006, 2011, 2012; Zhang *et al.* 2010; Hwang *et al.* 2012). Recently PDGFR α ⁺ cells were isolated and shown to generate large amplitude apamin- and Ca²⁺-sensitive outward currents in response to purines (ATP, ADP and β -NAD; Kurahashi *et al.* 2011). The current density of SK-like currents in colonic smooth muscle cells was too small to account for large inhibitory junction potentials recorded from intact muscles in response to enteric inhibitory neurotransmission. Thus, it appears that PDGFR α ⁺ cells may contribute significantly to purinergic enteric neural control of GI motility. Similar cells, expressing similar ion channels and manifesting similar functions, may also be important regulators of compliance in detrusor muscles of the bladder (Koh *et al.* 2012; Monaghan *et al.* 2012; Lee *et al.* 2013).

No studies to date have investigated the responsiveness of PDGFR α ⁺ cells in intact muscles directly nor shown how purinergic signals are transduced into activation of SK-dependent outward currents. This is because it is very difficult to distinguish these cells in the mixed-cell populations that constitute GI muscles. In the present study we have used mice in which PDGFR α ⁺ cells are constitutively labelled with enhanced green fluorescent protein (eGFP), loaded cells with fluorescent Ca²⁺ indicators, and investigated spontaneous Ca²⁺ signalling and responses to purines in this new class of interstitial

cell. Our observations are compatible with the hypothesis that PDGFR α ⁺ cells are responsive to the purines and pathways released and activated during enteric inhibitory neural responses.

Methods

Animals

PDGFR α ^{tm11(EGFP)Sor/J} mice, *P2ry1*^(-/-) mice (B6.129P2-*P2ry1*^{tm1Bhk/J}) and their wild-type siblings (C57BL/6) were obtained from the Jackson Laboratory (Bar Harbor, MN, USA). Animals between the ages of 5 and 7 weeks (age-matched) were anaesthetized by inhalation of isoflurane (Baxter, Deerfield, IL, USA) and killed by cervical dislocation.

The animals used and the experiments performed in this study were maintained in accordance with the National Institutes of Health *Guide for the Care and Use of Laboratory Animals*. All procedures were approved by the Institutional Animal Use and Care Committee at the University of Nevada.

Tissue preparation

Stomachs, including portions of the oesophagus and duodenum, were removed from mice via an abdominal incision immediately after cervical dislocation. The stomachs were bathed in Krebs–Ringer bicarbonate solution (KRB), opened along the lesser curvature and the gastric contents washed away with KRB solution. The fundus region was identified and isolated (3 mm from the fundus–corpus border) and the mucosa and sub-mucosa layers were removed by sharp dissection.

Drugs and solutions

Tissues were maintained and perfused with KRB containing (mmol l⁻¹): NaCl, 120.35; KCl, 5.9; NaHCO₃, 15.5; NaH₂PO₄, 1.2; MgCl₂, 1.2; CaCl₂, 2.5; and glucose, 11.5. KRB was bubbled with a mixture of 97% O₂–3% CO₂ and warmed to a physiological temperature of 37 ± 0.2°C. For experiments with 0 mM [Ca²⁺]_o, no CaCl₂ was added to the KRB, thus the solutions were nominally free of Ca²⁺, but there may have been trace amounts of Ca²⁺ in these solutions. We refer to this as a 0 mM external Ca²⁺ solution.

Atropine, *N*^ω-nitro-L-arginine (L-NNA), adenosine triphosphate (ATP), adenine diphosphate (ADP), β -nicotinamide adenine dinucleotide (β -NAD), caffeine, 2-aminoethyl diphenylborinate (2-APB), cyclopiazonic acid (CPA) and thapsigargin were purchased from Sigma-Aldrich (St Louis, MO, USA). Ryanodine was

purchased from Ascent Scientific (Princeton, NJ, USA). (*N*)-Methanocarba-2MeSADP (MRS-2365), 2-iodo-6-(methylamino)-9*H*-purin-9-yl]-2-(phosphonoxy)bicyclohexane-1-methanol dihydrogen phosphate ester tetraammonium salt (MRS-2500) and tetrodotoxin (TTX) were purchased from Tocris Bioscience (Ellisville, MO, USA). All drugs were dissolved in the manufacturer-recommended solvent to make stock solutions and the final concentration stated in the Results section.

Immunohistochemistry

Whole mounts of fundus tissues were processed for immunohistochemical analysis. Tissues were pinned onto the base of a Sylgard dish after sharp dissecting and were fixed in acetone or paraformaldehyde (4°C; 10 min) as previously described (Kurahashi *et al.* 2011). Following fixation, preparations were washed for 30 min in phosphate-buffered saline (PBS; 0.1 M pH 7.4). Non-specific antibody binding was reduced by incubation in 1% bovine serum albumin for 1 h at room temperature. The tissues were incubated with primary antibodies for 48 h at 4°C and with secondary antibodies for 1 h at room temperature. The antibodies and dilutions used were as described previously (Kurahashi *et al.* 2011). Whole mounts were examined using a Zeiss LSM 510 Meta laser-scanning confocal microscope. Confocal micrographs are digital composites of Z-series scans of 0.5–1.0 optical sections through a depth of 5–40 μ m. Final images were constructed using Zeiss LSM software.

Cell isolation and fluorescence-activated cell sorting (FACS)

Fundus muscles of PDGFR α -eGFP mice were dissected as described above. Muscles were equilibrated in Ca²⁺-free Hanks' solution consisting of (in mM): 125 NaCl, 5.36 KCl, 15.5 NaHCO₃, 0.336 Na₂HPO₄, 0.44 KH₂PO₄, 10 glucose, 2.9 sucrose, and 11 Hepes adjusted to pH 7.2 with NaOH, for 30 min. The buffer was replaced with an enzyme solution containing 2 mg ml⁻¹ collagenase type 2 (Worthington Biochemical, Lakewood, NJ, USA), 4 mg ml⁻¹ bovine serum albumin (Sigma), and 4 mg ml⁻¹ trypsin inhibitor (Sigma). The tissues were placed in a 37°C water bath for 15 min and then triturated through a pipette tip to disperse the cells. PDGFR α -positive (eGFP) cells were sorted by FACS with a Becton-Dickinson FACSaria II instrument using an excitation laser (488 nm) and emission filter (530/30 nm). Sorting was performed using a 130 μ m nozzle at a sheath pressure of 12 p.s.i. and sort rate of 1000 to 3000 events s⁻¹. Live cells, gated on exclusion of the Hoechst 33258 viability indicator

(data not shown), were subsequently gated on eGFP fluorescence intensity.

RNA extraction and quantitative PCR

Total RNA was isolated from collected PDGFR α ⁺ cells and unsorted cells (total cell population from tissue dispersions before cell sorting) from six mice using an illustra RNAspin Mini RNA Isolation Kit (GE Healthcare, Pittsburgh, PA, USA), and first-strand cDNA was synthesized using SuperScript III (Invitrogen, Carlsbad, CA, USA), according to the manufacturer's instructions. The PCR primers used and their GenBank accession numbers are listed in Supplemental Table S1 (available online). Using GoTaq DNA Polymerase (Promega, Madison, WI, USA), PCR products were analysed on 2% agarose gels and visualized by ethidium bromide. Quantitative PCR (qPCR) was performed with the same primers as PCR using SYBR green chemistry on the 7500 HT Real-time PCR System (Applied Biosystems, Grand Island, NY, USA). Regression analysis of the mean values of eight multiplex qPCRs for the log₁₀ diluted cDNA was used to generate standard curves. Unknown amounts of mRNA were plotted relative to the standard curve for each set of primers and graphically plotted using Microsoft Excel (Microsoft, Redmond, WA, USA). This gave a transcriptional quantification of each gene relative to the endogenous glyceraldehyde-3-phosphate dehydrogenase (GAPDH) standard after log transformation of the corresponding raw data. A total of six mice were divided into two groups consisting of three mice each; qPCR analysis was run in duplicate in the two groups. There was high repeatability between the two groups, and the average was used for comparison between PDGFR α ⁺ cells and unsorted cells.

Calcium imaging

Fluorescent dye loading. Gastric fundus muscle strips were pinned to the base of a Sylgard-coated dish. After an equilibration period of (1 h), the preparation was loaded with Oregon Green 488 BAPTA-2 AM (10 μ g ml⁻¹; Molecular Probes, Eugene, OR, USA), in a solution of 0.02% DMSO and 0.01% non-toxic detergent Cremophor EL for 30 min at 25°C. After incubation, the preparation was perfused with warm KRB solution at 37°C for 40 min to allow for de-esterification of the dye.

Visualization of Ca²⁺ transients. Preparations were visualized and imaged using an Eclipse E600FN microscope (Nikon Inc., Melville, NY, USA) equipped with $\times 20$ to $\times 40$ lenses (Nikon Plan Fluor). The indicator was excited at 488 nm (T.I.L.L. Polychrome IV, Grafelfing, Germany) and the fluorescence emission (> 515 nm) was

detected using a cooled, interline transfer CCD-camera system (Imago, T.I.L.L. Photonics, Grafelfing, Germany). Image sequences were collected after 4×4 binning of the 344×260 line image in TILLvisION software (T.I.L.L. Photonics GmbH, Grafelfing, Germany). Movies and image sequences of Ca²⁺ activity in fundus cells were analysed and constructed using custom software (Volumetry G7wv, GWH). Movies were further processed for playback viewing using Quick Time player 7 (Apple Inc., C4 USA). A pseudo-coloured blue–green lookup table was used to recolour the movie. The average fluorescence throughout the entire movie was subtracted from a duplicate copy of the movie, then temporal averaging ($t \pm 0.08$ s) was applied. The average frequency of Ca²⁺ transients was calculated for 20 s before and after drug application. The following abbreviations were used throughout the analysis and figures: cpm, cycles per minute; c, cells; asterisk (*) motion/focus artifact).

Statistical analysis

The figures were constructed from digitized data using Adobe Photoshop 4.0.1 (Adobe Co., Mountain View, CA, USA), Excel and PowerPoint 2011 (Microsoft, Redmond, WA, USA). The bar graphs represent the means from each experiment and 'n' values refer to the number of animals. Data are expressed as means \pm standard error of the mean. Statistical significance was calculated using either Student's *t* test or a one-way ANOVA followed by a post Newman–Keuls test. *P* values of < 0.05 were considered to represent significant changes.

Results

Distribution of PDGFR α ⁺ cells in the gastric fundus

The distribution of PDGFR α ⁺ cells in the fundus included intramuscular (IM) cells that were intermingled in circular muscle (CM) bundles, similar intramuscular cells that were present in longitudinal muscle (LM) bundles, and more stellate-shaped cells that were found in the plane between CM and LM, in the region of the myenteric plexus (MY). These cells were found in equivalent density and with similar distributions in wild-type and genetically engineered eGFP-PDGFR α ⁺ mice. Immunohistochemistry using PDGFR α antibodies in wild-type fundus showed that intramuscular PDGFR α ⁺ cells (PDGFR α ⁺-IM) were arranged in parallel to smooth muscle bundles in the CM and LM layers (Fig. 1A and D). In the myenteric region, myenteric PDGFR α ⁺ cells (PDGFR α ⁺-MY) were characterized as extensively branching cells with multipolar cell bodies and long processes (Fig. 1G and J). PDGFR α immunopositive cells were distinct from ICC-IM (Kit⁺) (Fig. 1B and E) in

the CM and LM muscle layers as shown by double labelling (Fig. 1C and F). No Kit-like immunoreactivity was resolved in the myenteric region (Fig. 1H); however, PDGFR α ⁺-MY cells were prominent (Fig. 1G and I). These data clearly show that PDGFR α ⁺ cells are a class of interstitial cells distinct from ICC in the gastric fundus.

PDGFR α ⁺ cells express SK3 channels

We and others have shown that SK3 channel immunoreactivity is prominent in PDGFR α ⁺ cells from GI muscles and bladder of several species (i.e. colon, small intestine, detrusor) (Fujita *et al.* 2003; Iino & Nojyo, 2009; Kurahashi *et al.* 2011, 2012; Grover *et al.* 2012; Koh *et al.* 2012; Lee *et al.* 2013). SK3-like immunoreactivity was also present in PDGFR α ⁺ cells in fundus muscles. PDGFR α ⁺ cells co-labelled with SK3 antibodies labelled PDGFR α ⁺-IM

and PDGFR α ⁺-MY throughout the fundus (Fig. 1K and L, Supplemental Fig. S1, available online).

Distribution of PDGFR α ⁺ cells in relation to motor nerves

The distribution of PDGFR α ⁺ cells in relation to enteric neurons was evaluated in whole mount preparations double immunolabelled for PDGFR α and the neuronal marker, PGP 9.5. PDGFR α ⁺ cells were closely associated with PGP 9.5 immunoreactive neurons in CM and LM (Fig. 2A–C and Supplemental Fig. S2). PDGFR α ⁺ cells were also in close proximity to neuronal nitric oxide synthase (nNOS)-positive inhibitory motor neurons, as observed in the double immunolabelling in all muscle layers (Fig. 2D–F and Supplemental Fig. S3). PDGFR α ⁺ cells were also closely associated with vesicular acetylcholine transporter (vAChT)-positive neurons (Fig. 2G–I and Supplemental Fig. S4).

Nuclear eGFP as a reporter for PDGFR α ⁺ cells

Mice that express a histone 2B–eGFP fusion protein in cells expressing PDGFR α receptors were used to

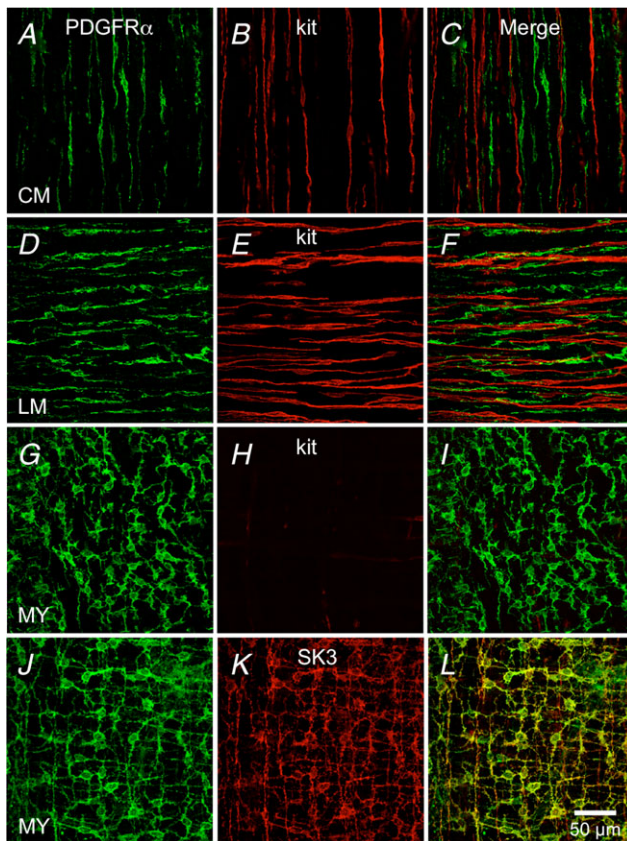


Figure 1. Distribution of PDGFR α ⁺ cells in the fundus

Whole mount double immunolabelling for PDGFR α (green) and Kit (red) in circular muscle layer (CM; A–C), in the longitudinal muscle layer (LM; D–F) and in the region of the myenteric plexus (MY; G–I). No co-localization was observed between PDGFR α ⁺ cells and Kit⁺ cells in any muscle layer (C, F and I). J–L, immunolabelling for PDGFR α (green) and SK3 (red) in MY region of gastric fundus. Expression of SK3 channels is observed in PDGFR α cells in the fundus. Scale bar in L is 50 μ m and pertains to all panels.

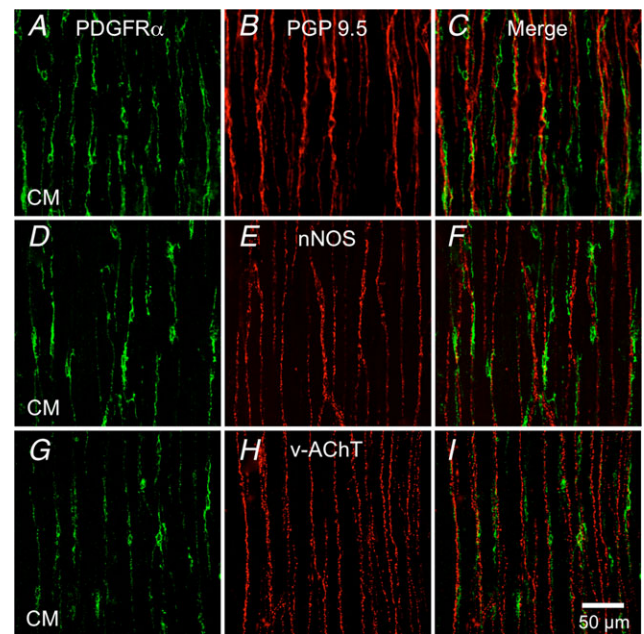


Figure 2. Relation of PDGFR α ⁺ cells to motor nerves in the fundus

Whole mount double immunolabelling for PDGFR α (green) and the pan neuronal marker PGP 9.5 (red) (A–C), double immunolabelling for PDGFR α (green) and nNOS (red) (D–F), double immunolabelling for PDGFR α (green) and vAChT (red) (G–I) in circular muscle layer (CM) of the fundus. PDGFR α ⁺ cells track closely along the processes of enteric motor neurons in both muscle layers. Scale bar in I is 50 μ m and pertains to all panels.

identify PDGFR α ⁺ cells in fundus muscles. A network of fluorescent cells (with eGFP confined to the nuclei) was observed in fundus muscles of these mice (Fig. 3A, D and G), and double labelling with PDGFR α antibodies confirmed that eGFP nuclei cells are PDGFR α ⁺ cells in all muscle layers (Fig. 3C, F and I).

Molecular expression of P2Y receptors and SK channels in PDGFR α ⁺ cells

It has been suggested that PDGFR α ⁺ cells are involved in purinergic neurotransmission via P2Y1 and subsequent SK channel activation (Cobine *et al.* 2011; Kurahashi *et al.* 2011; Lee *et al.* 2013). Therefore we examined the expression of P2Y receptors and SK channels in PDGFR α ⁺ cells. Fundus muscles were dispersed enzymatically, and PDGFR α ⁺ cells were collected by FACS using the eGFP reporter in eGFP-PDGFR α ⁺ muscles. Quantitative PCR (qPCR) was performed on cDNAs prepared from sorted PDGFR α ⁺ cells and compared to unsorted cells representing the total cell population from fundus muscles. We evaluated the expression of P2Y receptors (i.e. *P2ry1*, *P2ry2*, *P2ry4*, *P2ry6*, *P2ry12*, *P2ry13* and *P2ry14*) and found robust expression of *P2ry1* in comparison to other receptors

examined (fold change = 7.1 ± 0.76 , PDGFR α ⁺ sorted cells = 0.18 ± 0.02 , unsorted cells = 0.025 ± 0.002 , *P* value = 0.002; Fig. 4A). We also observed a higher expression of *P2ry2* (fold change = 1.6 ± 0.16 , PDGFR α ⁺ sorted cells = 0.02 ± 0.002 , unsorted cells = 0.012 ± 0.001 , *P* value = 0.003; Fig. 4A) and *P2ry4* receptors in PDGFR α ⁺ cells (fold change = 1.67 ± 0.09 , PDGFR α ⁺ sorted cells = 0.009 ± 0.0005 , unsorted cells = 0.005 ± 0.0003 , *P* value = 0.003; Fig. 4A). Other P2Y receptors (*P2ry6*, *P2ry12*, *P2ry13* and *P2ry14*) demonstrated no significant increase in expression in PDGFR α ⁺ cells (fold change: *P2ry6* = -1.1 ± 0.1 , *P2ry12* = -1.2 ± 0.08 , *P2ry13* = -1.25 ± 0.09 and *P2ry14* = -1.0 ± 0.06 ; Fig. 4A). We also evaluated the expression of SK channels (*Kcnn1*, *Kcnn2* and *Kcnn3*) and found high expression of *Kcnn3* (SK3) subtype in PDGFR α ⁺ cells in comparison to unsorted

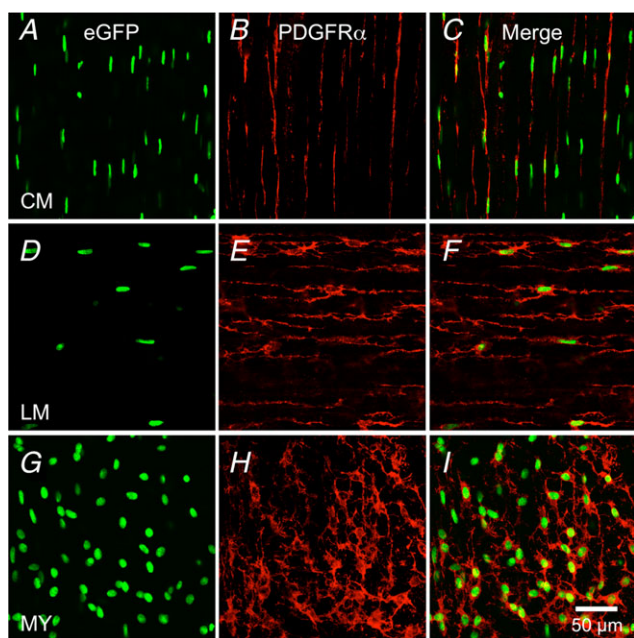


Figure 3. Nuclear eGFP driven off promoter for *Pdgfra* is an accurate reporter for PDGFR α ⁺ cells

Whole mount immunolabelling for eGFP (green) and PDGFR α (red) in circular muscle layer (CM; A–C), in the longitudinal muscle layer (LM; D–F) and in the region of the myenteric plexus (MY; G–I) of murine gastric fundus. eGFP is confined to nuclei because it is fused to histone 2B, while PDGFR α is expressed in plasma membranes. Scale bar in I is 50 μ m and pertains to all panels.

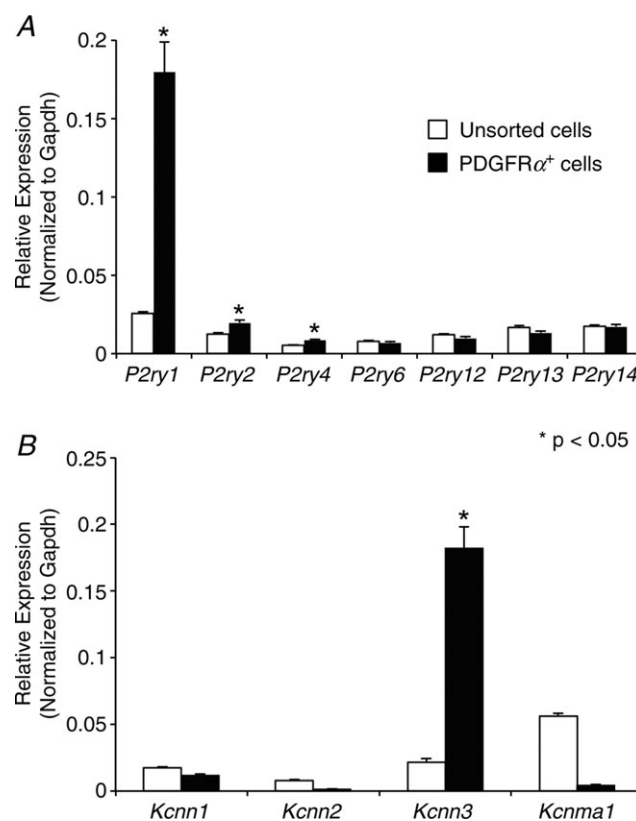


Figure 4. Expression of purinergic P2Y receptors and SK channels in PDGFR α ⁺ cells

A, relative expression comparison of P2Y receptor transcripts (*P2ry1*, *P2ry2*, *P2ry3*, *P2ry4*, *P2ry6*, *P2ry12*, *P2ry13* and *P2ry14*) in sorted PDGFR α ⁺ cells vs. unsorted cells (i.e. dispersions of fundus muscles prior to sorting) revealed by qPCR. Note *P2ry1* relative expression was 7.1-fold higher in PDGFR α ⁺ cells (*n* = 6, *P* value = 0.002). B, qPCR comparison of SK1, SK2, SK3 and α -Slo expression (*Kcnn1*, *Kcnn2*, *Kcnn3* and *Kcma1*) in sorted-PDGFR α ⁺ cells vs. unsorted cells. Note: *Kcnn3* expression was 8.4-fold higher in sorted PDGFR α ⁺ cells (*n* = 6, *P* value = 0.001). The relative expression of each gene was normalized to the house-keeping gene GAPDH.

cells (fold change = 8.4 ± 0.75 , PDGFR α ⁺ sorted cells = 0.18 ± 0.016 , unsorted cells = 0.02 ± 0.002 , P value = 0.001; Fig. 4B). *Kcnn1* and *Kcnn2* demonstrated low expression in PDGFR α ⁺ cells, *Kcnn1* (fold change = -1.42 ± 0.06) and *Kcnn2* (fold change = -5.18 ± 0.53 ; Fig. 4B). We also examined the expression of *Kcnn1* (BK channel α subunit, α -Slo) in PDGFR α ⁺ cells and found a low expression of α -Slo in this cell type (fold change = -11.8 ± 0.56 ; Fig. 4B). The data demonstrate robust expression of P2Y1 receptors and SK3 channels in PDGFR α ⁺ cells of the fundus.

Ca²⁺ signalling in PDGFR α ⁺ cells

Ca²⁺ imaging was performed on muscles of eGFP-PDGFR α ⁺ mice to examine spontaneous Ca²⁺ transients and purinergic responses of PDGFR α ⁺ cells. This technique allowed unequivocal identification of PDGFR α ⁺ cells (eGFP expression in nuclei) and did not obscure resolution of cytoplasmic Ca²⁺ transients. PDGFR α ⁺-MY (Fig. 5A) were found at an average density of 408 ± 28 cells mm⁻² ($n = 14$) and with average minimum separation between cell bodies of 34.6 ± 1.8 μ m ($n = 14$; $c = 213$; Fig. 5B).

Under basal conditions Ca²⁺ transients were resolved in many PDGFR α ⁺ cells within a given field (e.g. $24.2 \pm 5\%$ of cells generated spontaneous Ca²⁺ transients at a frequency of 7.1 ± 0.8 cpm, (range 2–16 cpm); $n = 10$; Fig. 5C). Spontaneous Ca²⁺ transients were not resolved in the remainder of PDGFR α ⁺ cells. Pretreatment of fundus tissues with 1 μ M tetrodotoxin (TTX) decreased, but did not abolish spontaneous Ca²⁺ transients in PDGFR α ⁺ cells (i.e. frequency: 5.1 ± 0.5 cpm, $n = 6$, $c = 12$, compared to the control frequency of 8.6 ± 0.94 cpm in the muscles, $n = 6$, $c = 12$, P value = 0.008; Fig. 5D and E, respectively).

Next we examined the responses of PDGFR α ⁺ cells to a variety of purine agonists and antagonists. All of these experiments were performed in the continued presence of L-NNA (100 μ M) and atropine (1 μ M) to distinguish purinergic responses from nitroergic and cholinergic influences. The average activity of spontaneous Ca²⁺ transients in PDGFR α ⁺ cells used in this part of the study was 6.68 ± 1.4 cpm ($n = 25$, $c = 64$).

A single bolus application of ATP (100 μ M) increased the frequency of Ca²⁺ transients in PDGFR α ⁺ cells (Fig. 6A and B). These responses were typically characterized by an initial sustained rise in fluorescence that tapered off gradually and lasted, on average for

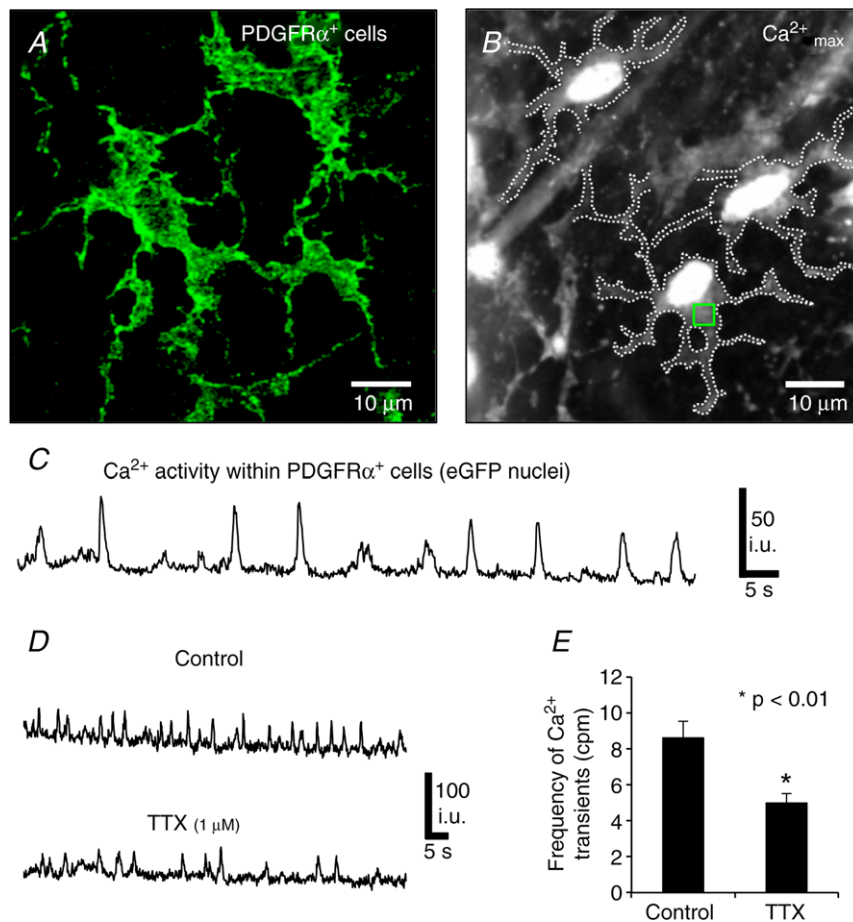


Figure 5. Spontaneous Ca²⁺ transients in PDGFR α ⁺ cells

A, PDGFR α ⁺ cells in the myenteric region at high magnification ($\times 100$). B, An image of maximum Ca²⁺ fluorescence within eGFP-PDGFR α ⁺ cells. Dotted lines demarcate the cell edges and the square (green) refers to the region of interest in which Ca²⁺ transients were measured (ROI). C, Spontaneous Ca²⁺ transients in PDGFR α ⁺ cells in the absence of any added drugs. D, Spontaneous Ca²⁺ transients in PDGFR α ⁺ cells before and after TTX (1 μ M). E, Summary of TTX effects on the frequency of spontaneous Ca²⁺ transients in PDGFR α ⁺ cells ($n = 6$, $c = 12$; P value = 0.008). The bar graphs represent the average means from each experiment and n number represents each experiment.

8.7 ± 0.86 s ($n = 12$, $c = 24$; Fig. 6A). The sustained rise was followed by a period of oscillatory Ca^{2+} waves that lasted through the recording window (Fig. 6A and B). ATP increased the frequency of Ca^{2+} transients to 20 ± 2.16 cpm ($n = 12$, $c = 24$, P value = 0.0001; Fig. 6A and K).

ADP ($100 \mu\text{M}$) evoked Ca^{2+} responses in the $\text{PDGFR}\alpha^+$ cells consisting of a sustained Ca^{2+} transient that gradually tapered off after 11.6 ± 0.9 s ($n = 12$, $c = 24$; Fig. 6C and D). The sustained phase was followed by increased Ca^{2+} waves at a frequency of 32.4 ± 2.53 cpm ($n = 12$, $c = 24$, P value = 0.0001; Fig. 6C and K and Supplemental Movie S1, available online).

β -NAD, a candidate for the purine neurotransmitter in GI muscles (Mutafova-Yambolieva *et al.* 2007), also evoked Ca^{2+} responses in $\text{PDGFR}\alpha^+$ cells similar to those evoked by ATP and ADP (Fig. 6E and F). A brief bolus application of β -NAD ($100 \mu\text{M}$) increased the frequency of Ca^{2+} transients. The response included an initial sustained

rise in fluorescence lasting 3.3 ± 0.31 s ($n = 12$, $c = 22$; Fig. 6E) and then sustained increase in Ca^{2+} oscillations averaging 12.2 ± 1.55 cpm ($n = 12$, $c = 22$, P value = 0.02; Fig. 6E and K).

The effects of uridine 5'-triphosphate (UTP) and uridine 5'-diphosphate (UDP) were also examined. UTP ($100 \mu\text{M}$) evoked Ca^{2+} responses in the $\text{PDGFR}\alpha^+$ cells consisting of a sustained Ca^{2+} transient that gradually tapered off after 4.9 ± 0.51 s ($n = 6$, $c = 14$; Fig. 6I). The sustained phase was followed by increased Ca^{2+} waves at a frequency of 20.5 ± 0.9 cpm ($n = 6$, $c = 14$, P value = 0.0001; Fig. 6I and K). UDP evoked a small sustained Ca^{2+} transient that gradually tapered off after 2.4 ± 0.42 s ($n = 5$, $c = 10$; Fig. 6J). UDP had no effect on the frequency of Ca^{2+} waves, except in a few quiescent cells that showed a small increase in Ca^{2+} transient frequency of 1.89 ± 0.57 cpm ($n = 5$, $c = 10$, P value = 0.0001; Fig. 6J and K). These experiments demonstrate the ability of $\text{PDGFR}\alpha^+$ cells to generate intracellular Ca^{2+} waves in

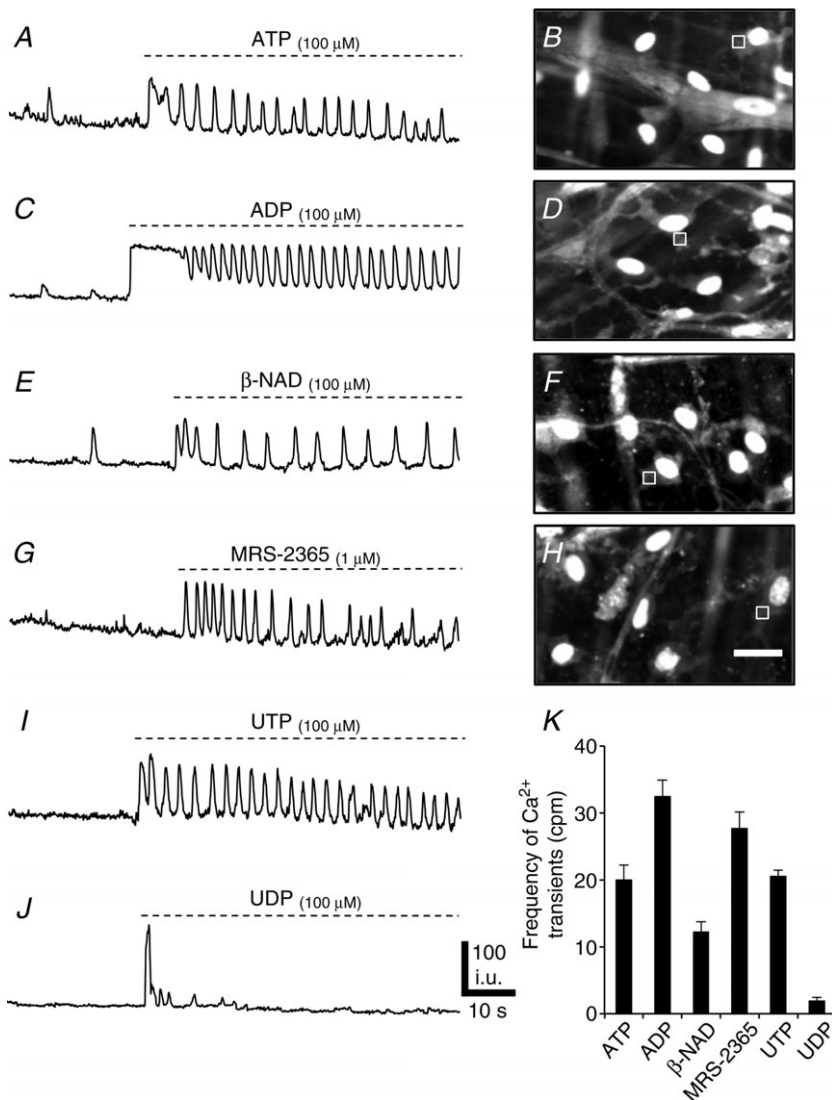


Figure 6. Responses of $\text{PDGFR}\alpha^+$ cells to purines (ATP, ADP, β -NAD, UTP and UDP) and P2Y1 agonist MRS-2365

A, Ca^{2+} transients in an ROI (square in B) in a $\text{PDGFR}\alpha^+$ cell elicited by ATP ($100 \mu\text{M}$) ($n = 12$, $c = 24$; P value = 0.0001). B, image of Ca^{2+} transients in eGFP- $\text{PDGFR}\alpha^+$ (eGFP nuclei) cells in response to ATP. The Ca^{2+} transients were recorded in ROI denoted by the square. C, an example of Ca^{2+} transients in an ROI (square in D) in a $\text{PDGFR}\alpha^+$ cell elicited by ADP ($100 \mu\text{M}$) ($n = 12$, $c = 24$; P value = 0.0001). D, image of Ca^{2+} transients in eGFP- $\text{PDGFR}\alpha^+$ cells in response to ADP. E, Ca^{2+} transients elicited in a $\text{PDGFR}\alpha^+$ cell (ROI: square in F) by β -NAD ($100 \mu\text{M}$) ($n = 12$, $c = 22$; P value = 0.02). F, image of Ca^{2+} transients in eGFP- $\text{PDGFR}\alpha^+$ (eGFP nuclei) in response to β -NAD. G, Ca^{2+} transients elicited in a $\text{PDGFR}\alpha^+$ cell (ROI: square in H) in response to MRS-2365 ($1 \mu\text{M}$). H, image of Ca^{2+} transients in eGFP- $\text{PDGFR}\alpha^+$ in response to MRS-2365 ($n = 12$, $c = 22$; P value = 0.0001). I, Ca^{2+} transients elicited in a $\text{PDGFR}\alpha^+$ cell in response to UTP ($100 \mu\text{M}$) ($n = 6$, $c = 14$; P value = 0.0001). J, Ca^{2+} transients elicited in a $\text{PDGFR}\alpha^+$ cell in response to UDP ($100 \mu\text{M}$) ($n = 5$, $c = 10$; P value = 0.0001). K, summary of changes in the frequency of Ca^{2+} transients in response to purines (each bar in the graph represents the average of the frequency of Ca^{2+} transients in response to a given purine and the n value represents the number tissues exposed to each purine). Scale bar in H is $20 \mu\text{m}$ and pertains to B, D and F panels.

response to purine compounds suggested to be purine neurotransmitters. In comparison, it should be noted that we did not resolve Ca²⁺ transients in the adjacent SMCs after the addition of purines.

Ca²⁺ transients were asynchronous in PDGFR α ⁺ cells

PDGFR α ⁺ cells exhibited spontaneous asynchronous Ca²⁺ waves that spread across cell bodies and occasionally invaded processes. Ca²⁺ waves spread with an average velocity within cells of $45.2 \pm 1.4 \mu\text{m s}^{-1}$ ($c = 37$, $n = 6$; Fig. 7A). There was no apparent synchronization in the occurrence of spontaneous Ca²⁺ transients in adjacent PDGFR α ⁺ cells, nor was there any evidence for propagation of spontaneous transients from cell to cell (Fig. 7B and C), suggesting that these events were localized and stochastic phenomena in cells. As described above, purines (e.g. ADP) increased Ca²⁺ transients. Within individual PDGFR α ⁺ cells the propagation velocity of Ca²⁺ waves increased to $51 \pm 2 \mu\text{m s}^{-1}$ in the presence of ADP ($c = 34$, $n = 6$, P value = 0.026; Fig. 7D). Ca²⁺ waves in adjacent cells were not synchronized by the addition of purines (Fig. 7D), and no evidence for cell-to-cell propagation of Ca²⁺ waves was observed.

Role of P2Y1 receptors in purinergic responses of PDGFR α ⁺ cells

Molecular studies showed a robust expression of P2Y1 receptors in PDGFR α ⁺ cells of the fundus. Previous studies have shown that purinergic stimulation activates outward currents in single isolated PDGFR α ⁺ cells (Kurahashi *et al.* 2011). Therefore, we evaluated the role of P2Y1 receptors in mediating the increase in Ca²⁺ transients in response to purinergic stimulation of PDGFR α ⁺ cells.

Similar to the response of cells to primary purines, a robust increase in Ca²⁺ transients was elicited in PDGFR α ⁺ cells in response to the P2Y1 receptor-specific agonist MRS-2365 ($1 \mu\text{M}$; Fig. 6G and H). MRS-2365 increased Ca²⁺ transient frequency to 27.7 ± 2.4 cpm ($n = 12$, $c = 22$; P value = 0.0001; Fig. 6G and H; Supplemental Movie S2). These responses were also characterized by a small, sustained rise in fluorescence that lasted for 0.8 ± 0.22 s ($n = 12$, $c = 12$; Fig. 6G).

Pre-treatment of muscles with MRS-2500 ($1 \mu\text{M}$), a highly selective antagonist of P2Y1 receptors, abolished Ca²⁺ waves elicited by β -NAD and MRS-2365 (Fig. 8D and E, respectively). Responses to ADP were also greatly diminished by MRS-2500, and the sustained Ca²⁺ oscillations typical of ADP stimulation were blocked in many cells (Fig. 8C). In 3 of 7 muscles Ca²⁺ oscillations persisted in a few PDGFR α ⁺ cells per field of view ($36 \pm 4.34\%$; frequency = 17.4 ± 3.23 cpm). In contrast, after MRS-2500 pretreatment, ATP evoked

Ca²⁺ transients in $60.2 \pm 5.7\%$ of PDGFR α ⁺ cells (i.e. frequency = 14.84 ± 2.6 cpm, $n = 6$, $c = 20$; Fig. 8A and B, Cell 2), but failed to elicit Ca²⁺ responses in some cells (e.g. Fig. 8B, Cell 1).

Responses of PDGFR α ⁺ cells from P2ry1^(-/-) mice

To verify the specificity of pharmacological treatments we also performed experiments on PDGFR α ⁺ cells from mice deficient in P2Y1 receptors. We bred our reporter strain (Pdgfra^{tm11(EGFP)Sor/J}) mice into a strain of P2ry1^(-/-) mice,

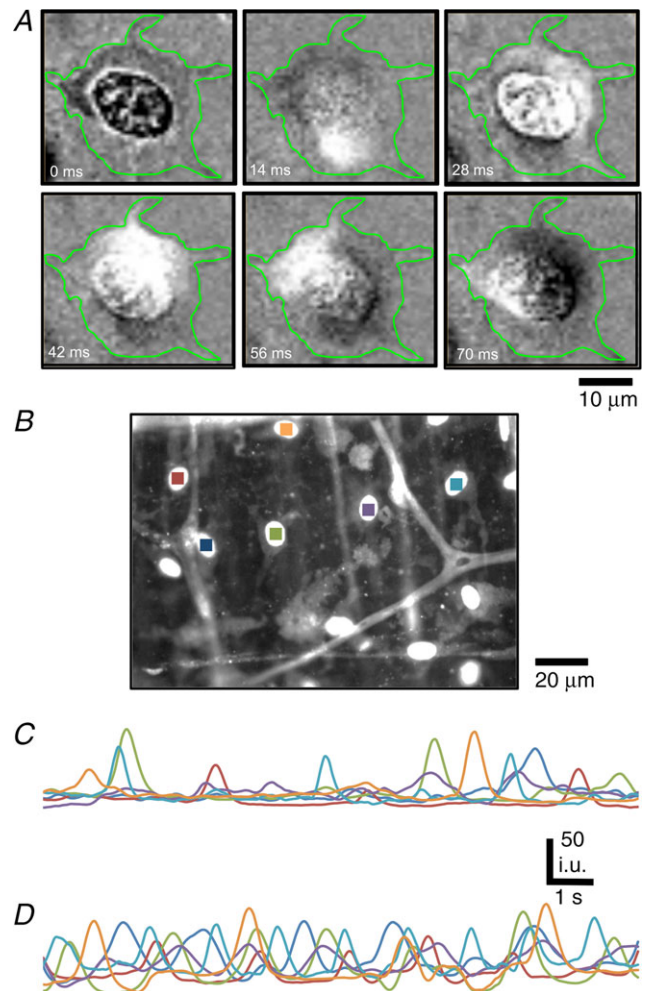


Figure 7. Asynchronous Ca²⁺ waves in PDGFR α ⁺ cells

A, representative time-sequence images of a spontaneous Ca²⁺ wave in a PDGFR α ⁺ cell. The duration of the Ca²⁺ wave was 0.56 s. Images were background corrected and the green outlines in each image denote the cell boundary ($n = 6$, $c = 37$; scale bar is $10 \mu\text{m}$). B–D, Ca²⁺ transients were monitored in PDGFR α ⁺ cells before and after ADP. B, Ca²⁺ transients were recorded in arbitrary units from ROIs defined in several PDGFR α ⁺ cells (coloured squares). Scale bar is $20 \mu\text{m}$. C, cytosolic Ca²⁺ dynamics in 6 PDGFR α ⁺ cells shown in coloured ROIs in C under control conditions and after ADP (D). Ca²⁺ transients were stochastic and non-propagating events in PDGFR α ⁺ cells before and after stimulation with ADP ($n = 6$).

creating mice deficient in P2Y1 receptors with constitutive expression of eGFP in PDGFR α ⁺ cells. We confirmed PDGFR α and SK3 channel expression in these mice using immunohistochemistry and found normal density and

distributions of PDGFR α ⁺ cells in the absence of P2Y1 receptors (Fig. 9A and B).

Next, we examined the effects of purines on PDGFR α ⁺ cells in the absence of P2Y1 receptors. ATP (100 μ M) application to *P2ry1*^(-/-) muscles yielded

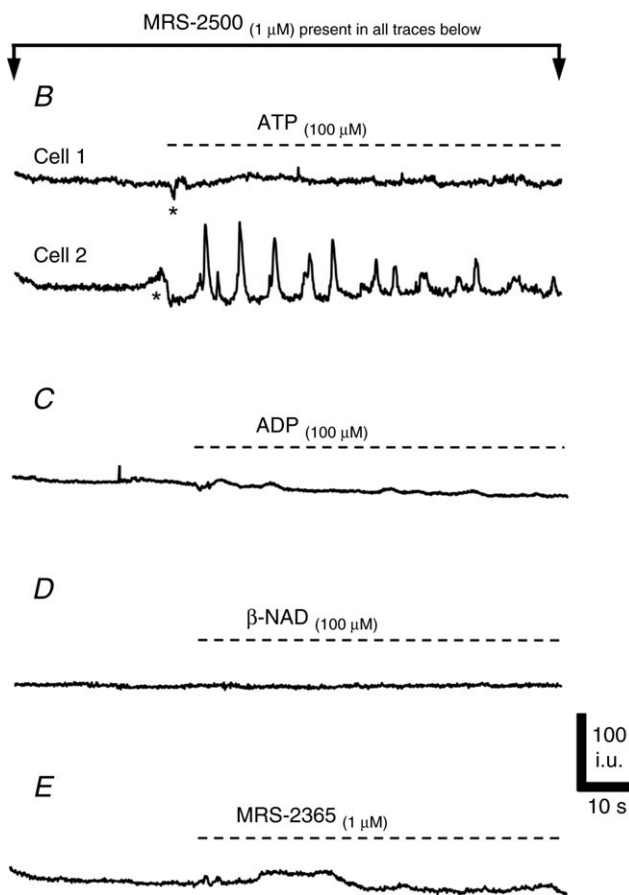
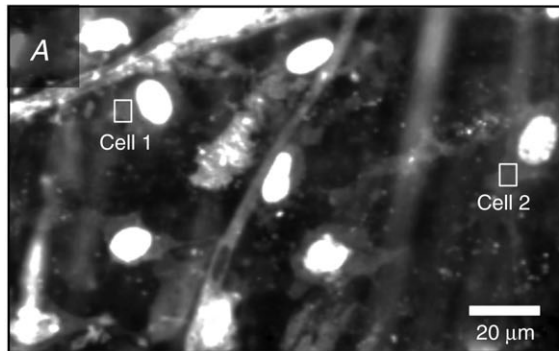


Figure 8. Role of P2Y1 receptors in mediating Ca²⁺ transients in PDGFR α ⁺ cells

A–E, Ca²⁺ responses of PDGFR α ⁺ cells in the presence of a P2Y1-receptor specific antagonist MRS-2500 (1 μ M). A, example image of Ca²⁺ transients in eGFP-PDGFR α ⁺ cells (eGFP nuclei) in response to ATP. B, example of Ca²⁺ transients in PDGFR α ⁺ cells in response to ATP (100 μ M). Cell 1 showed no response to ATP in the presence of MRS-2500 and Cell 2 showed an increase in Ca²⁺ transients ($n = 6$, $c = 20$; P value = 0.014). (ROIs of Cells 1 and 2 are marked by the squares in A) Ca²⁺ responses to ADP (C), β -NAD (D) and MRS-2365 (E) in PDGFR α ⁺ cells were blocked by MRS-2500.

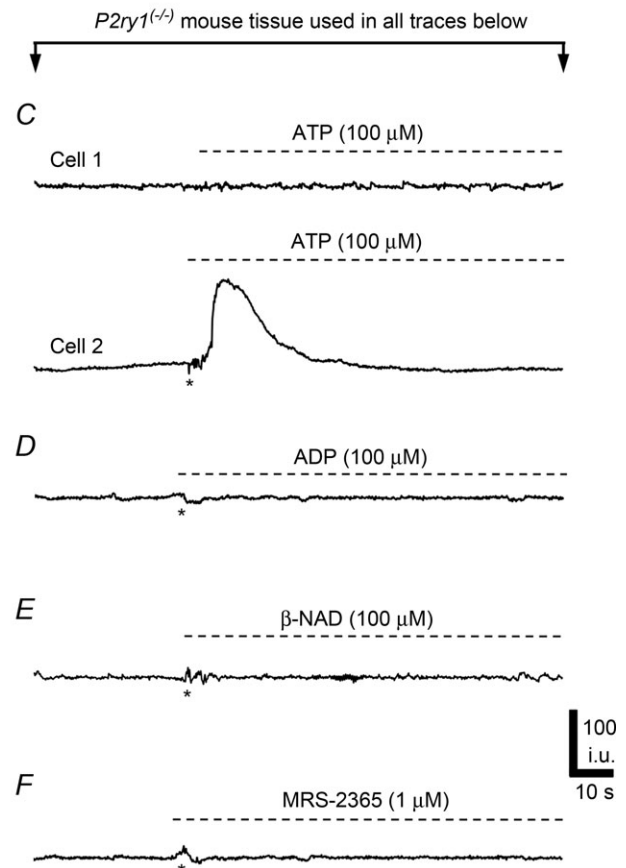
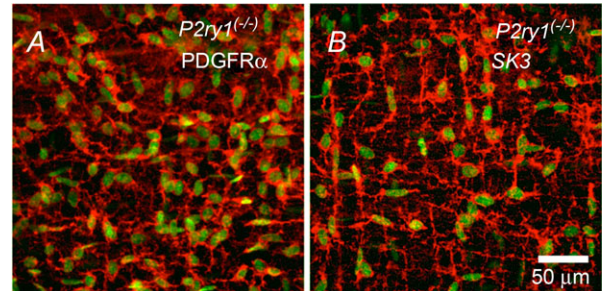


Figure 9. Effects of ATP, ADP, β -NAD and MRS-2365 in PDGFR α ⁺ cells from *P2ry1*^(-/-) mice

A, whole mount double immunolabelling for eGFP-PDGFR α (green eGFP nuclei) and PDGFR α (red, A) and SK3 (red, B) antibodies in PDGFR α ⁺ cells from *P2Y1*^(-/-) mice. C–F, Ca²⁺ transients in response to purines were attenuated in PDGFR α ⁺ cells of *P2Y1*^(-/-) mice. C, example of Ca²⁺ transients in PDGFR α ⁺ cells in response to ATP (100 μ M). No response was elicited in some cells (e.g. Cell 1), but others showed a rise in Ca²⁺ that gradually tapered off during the exposure (e.g. Cell 2; $n = 5$, $c = 10$). The other purines tested (ADP (D), β -NAD (E) and MRS-2365 (F)) failed to elicit Ca²⁺ transients in PDGFR α ⁺ cells of *P2Y1*^(-/-) mice. (For all C, D, E and F, $n = 5$).

mixed responses; some PDGFR α ⁺ cells showed little or no response to ATP (Fig. 9C, Cell 1) but others (32.6 ± 3.38%) displayed Ca²⁺ responses consisting of a sustained rise in Ca²⁺ that tapered off gradually and lasted 27.4 ± 2.16 s (*n* = 5, *c* = 10; Fig. 9C, Cell 2). Ca²⁺ oscillations were not observed in PDGFR α ⁺ cells from *P2ry1*^(-/-) mice. These data show that ATP can mediate its post-junctional effects in PDGFR α ⁺ cells through purinergic receptors other than P2Y1 receptors. In contrast, the P2Y1-specific agonist MRS-2365 (1 μM) had no effect on Ca²⁺ transients in PDGFR α ⁺ cells of *P2ry1*^(-/-) mice (Fig. 9F). Similarly, neither ADP nor β-NAD (both 100 μM) elicited Ca²⁺ responses in PDGFR α ⁺ cells deficient in P2Y1 receptors (Fig. 9D and E, respectively).

Contributions from extracellular and intracellular Ca²⁺ to transients in PDGFR α ⁺ cells

The source(s) of Ca²⁺ contributing to Ca²⁺ responses in PDGFR α ⁺ cells were examined in another series of experiments. Contributions from external Ca²⁺ ([Ca²⁺]_o) were tested by substituting the normal 2.5 mM [Ca²⁺]_o in KRB with KRB containing 0 mM [Ca²⁺]_o (nominally free Ca²⁺ solution, 20 min). ATP and ADP elicited Ca²⁺ transients in PDGFR α ⁺ cells in the presence of 0 mM [Ca²⁺]_o. The frequency of the Ca²⁺ transients was reduced by 21.8% in response to ATP (frequency = 17.98 ± 2.14 cpm, *n* = 7, *c* = 14, *P* value = 0.02; Fig. 10B) and 26.5% in response to ADP (frequency = 20.45 ± 2.1 cpm, *n* = 7, *c* = 14, *P* value = 0.0017; Fig. 10A and B). We also noted a 27.3% reduction in Ca²⁺ transients frequency in response to MRS-2365 in the presence of 0 mM [Ca²⁺]_o (frequency = 17.66 ± 1.2 cpm, *n* = 7, *c* = 14, *P* value = 0.01; Fig. 10B). These data suggest that purinergic responses of PDGFR α ⁺ cells are not immediately dependent upon Ca²⁺ influx mechanisms; however, responses run down when [Ca²⁺]_o is not sufficient to maintain intracellular stores.

We also examined the role of Ca²⁺ uptake and release mechanisms on purinergic responses of PDGFR α ⁺ cells. CPA (10 μM) and thapsigargin (1 μM) pre-treatment blocked Ca²⁺ responses to ATP, ADP and β-NAD in PDGFR α ⁺ cells (Fig. 10C and D). 2-APB (50 μM) caused significant reductions in Ca²⁺ transients elicited in PDGFR α ⁺ cells by ATP (e.g. frequency = 7.5 ± 1.84 cpm, *n* = 5, *c* = 12, *P* value = 0.001; Fig. 11D), ADP (frequency = 5.51 ± 1.21 cpm, *n* = 5, *c* = 12, *P* value = 0.001; Fig. 11B and D), and β-NAD (frequency = 2.4 ± 0.5 cpm, *n* = 5, *c* = 10, *P* value = 0.001; Fig. 11D). 2-APB (100 μM) further attenuated purinergic responses: ATP (frequency = 0.55 ± 0.37 cpm, *n* = 5, *c* = 12, *P* value = 0.001; Fig. 11D), ADP (frequency = 0.38 ± 0.27 cpm, *n* = 5, *c* = 12, *P* value = 0.001; Fig. 11C

and D), and β-NAD (frequency = 0.4 ± 0.24 cpm, *n* = 5, *c* = 10, *P* value = 0.001, Fig. 11D). The phospholipase C (PLC) inhibitor, U-73122 (10 μM) inhibited purinergic responses (*n* = 5, Fig. 11E); however, its inactive analogue U-73343 (10 μM) had no significant effect and the frequency of Ca²⁺ oscillations was not significantly different from control responses (*P* > 0.05, *n* = 4; Fig. 11F).

Caffeine (10 mM), an activator of ryanodine-sensitive channels, reduced Ca²⁺ transients in PDGFR α ⁺ cells in response to purines: ATP (frequency =

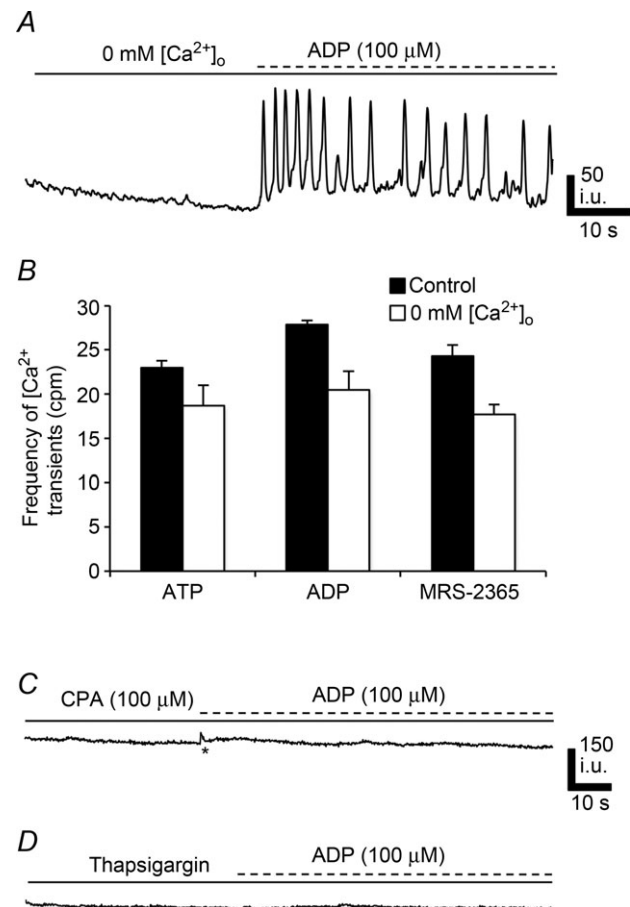


Figure 10. Contribution of extracellular [Ca²⁺]_o and SERCA pump to purinergic Ca²⁺ transients in PDGFR α ⁺ cells

A, Ca²⁺ responses in PDGFR α ⁺ cells to ADP (100 μM) were not blocked after bathing cells in 0 mM [Ca²⁺]_o for 20 min. B, summary of Ca²⁺ transient frequency in PDGFR α ⁺ cells in the presence of 2.5 mM [Ca²⁺]_o (black) vs. 0 mM [Ca²⁺]_o (white) in response to ATP (*n* = 7, *c* = 14; *P* value = 0.02), ADP (*n* = 7, *c* = 14; *P* value = 0.0017) and MRS-2365 (*n* = 7, *c* = 14; *P* value = 0.01). (Each bar in the graph represents the average of the frequency of Ca²⁺ transients in response to a given purine and the *n* value represents the number of tissues exposed to each purine). C, the SERCA pump inhibitor CPA (10 μM) blocked Ca²⁺ transients in PDGFR α ⁺ cells in response to ADP (100 μM). D, pretreatment of PDGFR α ⁺ cells with thapsigargin (1 μM) inhibited Ca²⁺ transients in response to ADP (100 μM). All *n* = 5.

14.9 ± 2.03 cpm, $n = 5$, $c = 12$, P value = 0.01; Fig. 12E), ADP (frequency = 17.5 ± 1.15 cpm, $n = 5$, $c = 12$, P value = 0.001; Fig. 12B and E), and β -NAD (frequency = 5.5 ± 0.57 cpm, $n = 5$,

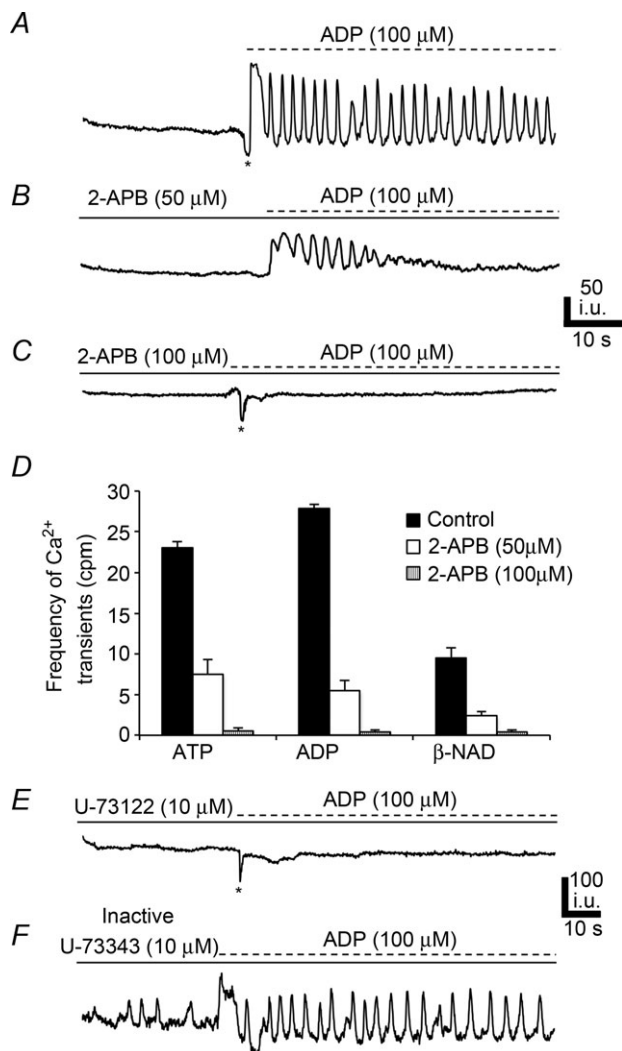


Figure 11. Role of IP₃ receptor-operated stores and phospholipase C (PLC) in purinergic responses of PDGFR α ⁺ cells

A, Ca²⁺ transients elicited in PDGFR α ⁺ cells in response to ADP (100 μ M). L-NNA (100 μ M) and atropine (1 μ M) present in all panels. B, 2-APB (50 μ M) reduced Ca²⁺ transient frequency in PDGFR α ⁺ cells ($n = 5$, $c = 12$; P value = 0.001). C, 2-APB (100 μ M) blocked most of the response to ADP ($n = 5$, $c = 12$; P value = 0.001). D, summary of Ca²⁺ transient frequency in PDGFR α ⁺ cells in response to purines (ATP, ADP and β -NAD) in control (black) and in the presence of 2-APB (50 μ M; white) and 2-APB (100 μ M; white with vertical lines) All $n = 5$. (Each bar in the graph represents the average of the frequency of Ca²⁺ transients in response to a given purine and the n value represents the number of tissues exposed to each purine). E, PLC inhibitor U-73122 (10 μ M; $n = 5$) blocked responses to ADP in PDGFR α ⁺ cells. However, the inactive analogue U-73343 (10 μ M; $n = 4$) had no significant effect (F).

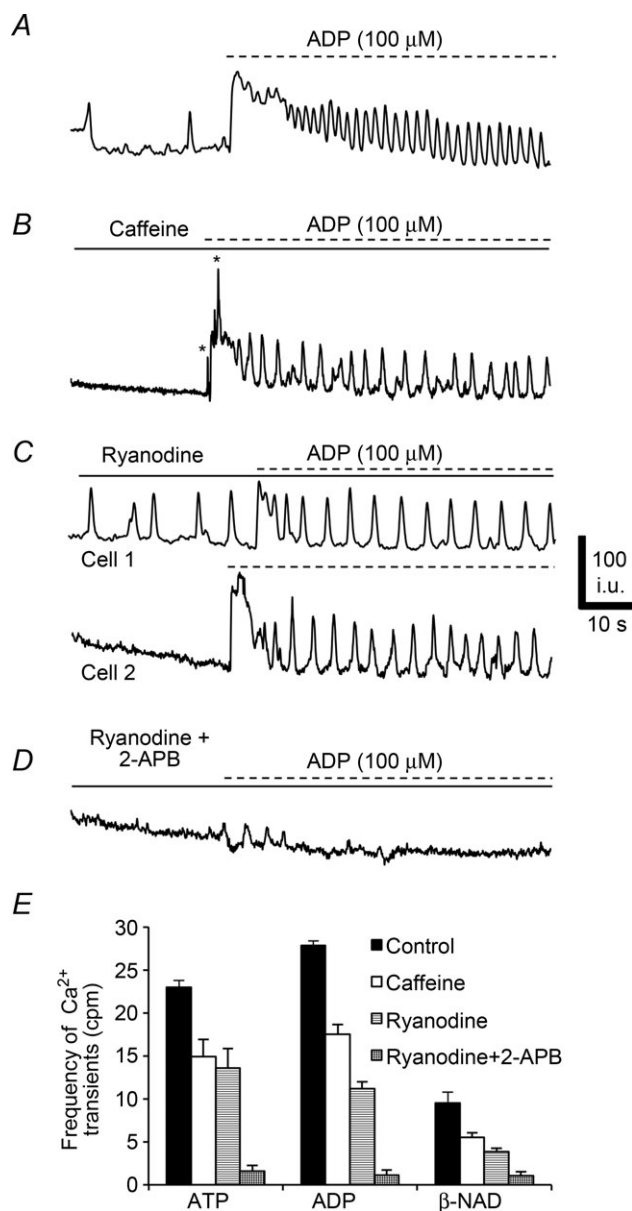


Figure 12. Role of ryanodine receptors in purinergic responses of PDGFR α ⁺ cells

A, Ca²⁺ transients in PDGFR α ⁺ cells after ADP (100 μ M). All traces were recorded in the presence of L-NNA (100 μ M) and atropine (1 μ M). B, caffeine (10 mM) reduced the frequency of Ca²⁺ transients in PDGFR α ⁺ cells elicited by ADP ($n = 5$, $c = 12$; P value = 0.001). C, ryanodine (50 μ M) also reduced Ca²⁺ transients in PDGFR α ⁺ cells (see Cell 1 and Cell 2; $n = 5$, $c = 14$; P value = 0.001). Some PDGFR α ⁺ cells (e.g. Cell 1) displayed increased spontaneous Ca²⁺ transients in the presence of ryanodine. D, a combination of ryanodine (50 μ M) and 2-APB (50 μ M) blocked most of the Ca²⁺ transients in PDGFR α ⁺ cells and responses to purines ($n = 5$, $c = 12$; P value = 0.001). E, summary of Ca²⁺ transient frequency in PDGFR α ⁺ cells in response to purines (ATP and ADP and β -NAD, $n = 15$) control (black) and in the presence of caffeine (10 mM; white), ryanodine (50 μ M; white with horizontal lines) and ryanodine and 2-APB together (50 μ M; white cross hatched). Each bar in the graph represents the average of the frequency of Ca²⁺ transients in response to a given purine and the n value represents the number of tissues exposed to each purine.

$c = 12$, P value = 0.01; Fig. 12E). Ryanodine (50 μ M) also attenuated Ca²⁺ transients in response to purines: ATP (frequency = 13.55 ± 2.29 cpm, $n = 5$, $c = 14$, P value = 0.01; Fig. 12E), ADP (frequency = 11.15 ± 0.85 cpm, $n = 5$, $c = 14$, P value = 0.001; Fig. 12C and E), and β -NAD (frequency = 3.85 ± 0.42 cpm, $n = 5$, $c = 14$, P value = 0.001, Fig. 12E). It should also be noted that there was an increase in the number of PDGFR α ⁺ cells exhibiting spontaneous Ca²⁺ transients in the presence of ryanodine in comparison to control conditions (i.e. $52.87 \pm 5.1\%$ displayed spontaneous transients in the presence of ryanodine and the frequency = 6.86 ± 0.89 cpm (Fig. 12C, Cell 1). Pre-exposure to ryanodine and 2-APB (50 μ M) almost eliminated Ca²⁺ responses to ATP (frequency = 1.57 ± 0.71 cpm, $n = 5$, $c = 12$, P value = 0.001; Fig. 12E), ADP (frequency = 1.14 ± 0.59 cpm, $n = 5$, $c = 12$, P value = 0.001; Fig. 12C and E), and β -NAD (frequency = 1.05 ± 0.5 cpm, $n = 5$, $c = 12$, P value = 0.001, Fig. 12E).

Discussion

The present study documents the distribution of PDGFR α ⁺ cells through the tunica muscularis of the murine gastric fundus. These cells were identified as 'fibroblast-like' in the classic morphological literature, but their close association with motor neurons, expression of receptors and effectors involved in purinergic motor responses, and gap junction coupling with smooth muscle cells suggests a more dynamic role for these cells in motor regulation than just maintenance of the extracellular matrix. There is an abundance of PDGFR α ⁺ cells in the fundus, and significant numbers of these cells generated spontaneous Ca²⁺ transients. Spontaneous Ca²⁺ transients were isolated, stochastic events characterized by localized Ca²⁺ waves that failed to propagate to cells nearby. Purines activated dynamic Ca²⁺ transients in PDGFR α ⁺ cells that were mediated by P2Y1 receptors and oscillatory release of Ca²⁺ from Ca²⁺ stores. Ca²⁺ transients evoked by purines were also localized to individual cells. Because PDGFR α ⁺ cells also express small conductance Ca²⁺-activated K⁺ channels (SK3), it is likely that spontaneous and purine-activated Ca²⁺ transients are coupled to activation of outward currents in PDGFR α ⁺ cells (as has been described in the colon; Kurahashi *et al.* 2011). Electrical coupling to SMCs suggests that purinergic responses in PDGFR α ⁺ cells would couple to relaxation responses in the fundus.

We confirmed that PDGFR α ⁺ cells are a special population of interstitial cells, distinct from ICCs in the gastric fundus, in agreement with previous reports in several GI muscles (Iino *et al.* 2009; Cobine *et al.* 2011; Kurahashi *et al.* 2011; Blair *et al.* 2012). One recent study claimed that Kit-positive cells (ICC) are also

immunopositive for PDGFR α in bladder lamina propria and detrusor muscle (Monaghan *et al.* 2012). However, the absence of ICC-MY in the fundus myenteric region and the presence of an extensive branching network of PDGFR α ⁺ cells observed in this study clearly confirms the existence of two independent populations of interstitial cells. Furthermore, we demonstrated the close association of PDGFR α ⁺ cells to motor neurons in the gastric fundus as previously described for other GI muscles (Iino *et al.* 2009; Iino & Nojyo, 2009; Kurahashi *et al.* 2011, 2012). This morphological evidence of the distribution of PDGFR α ⁺ cells across the gastric fundus as an integral part of the SIP syncytium indicates a role for them in regulating the excitability of gastric muscles.

Considerable evidence now suggests that PDGFR α ⁺ cells are primary targets for purinergic neurotransmission in visceral smooth muscles (Kurahashi *et al.* 2011; Lee *et al.* 2013). Purinergic inhibitory responses in post-junctional cells are usually attributed to the activation of small conductance Ca²⁺-activated K⁺ (SK) channels (Banks *et al.* 1979; Mutafova-Yambolieva *et al.* 2007). In this study we found robust expression of SK3 channels in PDGFR α ⁺ cells and not in SMCs or ICC. Interstitial cells (ICC-IM) are distributed in the gastric fundus and have been found to have a role in the mediation of excitatory cholinergic and inhibitory nitrenergic neurotransmission signals (Burns *et al.* 1996; Ward *et al.* 2000, 2004; Beckett *et al.* 2002; Suzuki *et al.* 2003; Ward & Sanders, 2006). It has been observed that purinergic neurotransmission, which is manifest as fast inhibitory junction potentials, is intact in muscles of *W/W^V* mice that lack ICC-IM (Burns *et al.* 1996), suggesting that ICC may have little role in purinergic responses. The direct involvement of SMCs in purinergic transmission is also unlikely since the net response of these cells to ATP at physiological holding potentials is activation of non-selective cationic channels and membrane depolarization (Monaghan *et al.* 2006). Additionally, SMCs respond to purines with a net inward current in bladder SMCs (Lee *et al.* 2013). These studies suggest the inability of ICC or SMCs to mediate purinergic responses, and they cannot account for the overall hyperpolarizing membrane potential by purines observed in GI muscles. PDGFR α ⁺ cells have the necessary machinery to generate outward currents in response to purines (Kurahashi *et al.* 2011; Lee *et al.* 2013). This study adds another important piece of information by showing that PDGFR α ⁺ cells also have the appropriate apparatus to generate intracellular Ca²⁺ transients in response to purines that may be the link to activate SK3 channels. Outward currents and hyperpolarization responses generated in PDGFR α ⁺ cells can conduct to SMCs and ICC via gap junction couplings between these cells (Komuro *et al.* 1999; Fujita *et al.* 2003). Thus, purinergic responses in PDGFR α ⁺ cells

can be conveyed and regulate the excitability of the SIP syncytium.

Spontaneous Ca^{2+} transients in $\text{PDGFR}\alpha^+$ cells were attenuated by the neuronal blocker TTX, suggesting a possible role of $\text{PDGFR}\alpha^+$ cells in the basal regulation of tone in the proximal stomach through an ongoing release of neurotransmitters (purines). In a related study examining how purinergic neurotransmission affects spontaneous activity in colon muscle tissues, it was found that TTX and the SK channel blocker, apamin, inhibit spontaneous inhibitory junction potentials (IJP) and increase basal contractile activity (Gil *et al.* 2010). It was also found that spontaneous IJPs are due to activation of the P2Y1 receptor, because the antagonist MRS-2500 inhibited IJPs.

It is widely accepted that P2Y receptors mediate purinergic neurotransmission in the gut (Giaroni *et al.* 2002; Gallego *et al.* 2006; Grasa *et al.* 2009a; Zhang *et al.* 2010), and P2Y1 receptors are the primary receptors responsible for fast IJPs (Gallego *et al.* 2006, 2012; Grasa *et al.* 2009a; Hwang *et al.* 2012). Several compounds have been claimed to be specific antagonists for P2Y1 receptors, but MRS-2500 has been reported to be the most potent and was therefore chosen for this study (Cattaneo *et al.* 2004; Grasa *et al.* 2009a,b). Ca^{2+} transients elicited by ADP, β -NAD and MRS-2365 in $\text{PDGFR}\alpha^+$ cells were blocked by MRS-2500. However, Ca^{2+} transients in response to ATP were retained in many $\text{PDGFR}\alpha^+$ cells, indicating that additional purinergic receptors are expressed by $\text{PDGFR}\alpha^+$ cells and available to ATP. Use of $P2ry1^{-/-}$ mice provided results similar to the results of studies using pharmacological agonists and antagonists of P2Y1 receptors. Ca^{2+} transients evoked by ADP, MRS-2365 and β -NAD were not observed in $\text{PDGFR}\alpha^+$ cells deficient in P2Y1 receptors. Responses to ATP could still be elicited in $P2ry1^{-/-}$ $\text{PDGFR}\alpha^+$ cells. P2Y2 receptors may have mediated the effects of ATP in $\text{PDGFR}\alpha^+$ cells in muscles of $P2ry1^{-/-}$ mice, because these receptors were expressed by $\text{PDGFR}\alpha^+$ cells and UTP also elicited Ca^{2+} transients in $\text{PDGFR}\alpha^+$ cells. P2Y2 receptors are equally sensitive to ATP and UTP but not sensitive to metabolites of ATP such as ADP (Velázquez *et al.* 2000). Like P2Y1 receptors, P2Y2 receptors are coupled to Ca^{2+} release via the PLC pathway (Murthy & Makhlouf, 1998). Expression of P2Y2 receptors by $\text{PDGFR}\alpha^+$ cells may explain the purinergic effects independent of P2Y1 receptors observed in this study.

Ca^{2+} transients in $\text{PDGFR}\alpha^+$ cells appear to be due to release of Ca^{2+} from intracellular stores, because reduction in $[\text{Ca}^{2+}]_o$ for 20 min did not block responses to purines. Blocking Ca^{2+} uptake into stores with SERCA pump inhibitory drugs blocked Ca^{2+} transients evoked by purines. Ca^{2+} release mechanisms may require synergism between IP_3 and ryanodine receptors because blockers of both channels were necessary to eliminate Ca^{2+} trans-

ients. 2-APB was used as a blocker of IP_3 -dependent Ca^{2+} release, but this drug has some non-specific effects such as inhibition of SERCA pumps and block of store-operated Ca^{2+} channels in the plasma membrane (Bootman *et al.* 2002; Peppiatt *et al.* 2003). Further investigation will be needed with more highly selective IP_3 receptor antagonists and/or experiments using IP_3 receptor knockout animals.

In summary this study established the ability of the $\text{PDGFR}\alpha^+$ class of interstitial cells to respond to purines *in situ*. Previous studies showed the activation of Ca^{2+} -activated outward currents in isolated $\text{PDGFR}\alpha^+$ cells, and this study provides a mechanism for activation of Ca^{2+} -activated currents through oscillatory release of Ca^{2+} from internal stores. A significant percentage of $\text{PDGFR}\alpha^+$ cells generated Ca^{2+} transients spontaneously, and some of this activity was related to tonic inhibitory input from motor neurons. Thus, $\text{PDGFR}\alpha^+$ cells may be involved in setting the basal excitability and tone of the proximal stomach. We also found that Ca^{2+} release was activated by a variety of purine compounds, and responses to ADP, β -NAD and MRS-2365 were mediated via P2Y1 receptors. Responses to these purines are analogous to the responses elicited in intact muscles by the purine neurotransmitter released from enteric inhibitory motor neurons. Ca^{2+} transients were also evoked in $\text{PDGFR}\alpha^+$ cells by ATP, but these responses appeared to be mediated through receptors in addition to P2Y1, possibly the P2Y2 receptors expressed by these cells. Activation of outward currents and hyperpolarization responses in $\text{PDGFR}\alpha^+$ cells can be conveyed via gap junctions to the other cell types in the SIP syncytium and thereby regulate the excitability of this cellular network in the gastric fundus.

References

- Banks BE, Brown C, Burgess GM, Burnstock G, Claret M, Cocks TM & Jenkinson DH (1979). Apamin blocks certain neurotransmitter-induced increases in potassium permeability. *Nature* **282**, 415–417.
- Beckett EA, Horiguchi K, Khoyi M, Sanders KM & Ward SM (2002). Loss of enteric motor neurotransmission in the gastric fundus of *Sl/Sl^d* mice. *J Physiol* **543**, 871–887.
- Blair PJ, Bayguinov Y, Sanders KM & Ward SM (2012). Relationship between enteric neurons and interstitial cells in the primate gastrointestinal tract. *Neurogastroenterol Motil* **24**, e437–e449.
- Bootman MD, Collins TJ, Mackenzie L, Roderick HL, Berridge MJ & Peppiatt CM (2002). 2-Aminoethoxydiphenyl borate (2-APB) is a reliable blocker of store-operated Ca^{2+} entry but an inconsistent inhibitor of InsP_3 -induced Ca^{2+} release. *FASEB J* **16**, 1145–1150.
- Burns AJ, Lomax AE, Torihashi S, Sanders KM & Ward SM (1996). Interstitial cells of Cajal mediate inhibitory neurotransmission in the stomach. *Proc Natl Acad Sci U S A* **93**, 12008–12013.

- Cattaneo M, Lecchi A, Ohno M, Joshi BV, Besada P, Tchilibon S, Lombardi R, Bischofberger N, Harden TK & Jacobson KA (2004). Antiaggregatory activity in human platelets of potent antagonists of the P2Y₁ receptor. *Biochem Pharmacol* **68**, 1995–2002.
- Cobine CA, Hennig GW, Kurahashi M, Sanders KM, Ward SM & Keef KD (2011). Relationship between interstitial cells of Cajal, fibroblast-like cells and inhibitory motor nerves in the internal anal sphincter. *Cell Tissue Res* **344**, 17–30.
- Fujita A, Takeuchi T, Jun H & Hata F (2003). Localization of Ca²⁺-activated K⁺ channel, SK3, in fibroblast-like cells forming gap junctions with smooth muscle cells in the mouse small intestine. *J Pharmacol Sci* **92**, 35–42.
- Gallego D, Gil V, Aleu J, Martinez-Cutillas M, Clave P & Jimenez M (2011). Pharmacological characterization of purinergic inhibitory neuromuscular transmission in the human colon. *Neurogastroenterol Motil* **23**, e792–e338.
- Gallego D, Gil V, Martinez-Cutillas M, Mane N, Martin MT & Jimenez M (2012). Purinergic neuromuscular transmission is absent in the colon of P2Y₁ knocked out mice. *J Physiol* **590**, 1943–1956.
- Gallego D, Hernandez P, Clave P & Jimenez M (2006). P2Y₁ receptors mediate inhibitory purinergic neuromuscular transmission in the human colon. *Am J Physiol Gastrointest Liver Physiol* **291**, G584–G594.
- Giaroni C, Knight GE, Ruan HZ, Glass R, Bardini M, Lecchini S, Frigo G & Burnstock G (2002). P2 receptors in the murine gastrointestinal tract. *Neuropharmacology* **43**, 1313–1323.
- Gil V, Gallego D, Grasa L, Martin MT & Jimenez M (2010). Purinergic and nitrgergic neuromuscular transmission mediates spontaneous neuronal activity in the rat colon. *Am J Physiol Gastrointest Liver Physiol* **299**, G158–G169.
- Grasa L, Gil V, Gallego D, Martin MT & Jimenez M (2009a). P2Y₁ receptors mediate inhibitory neuromuscular transmission in the rat colon. *Br J Pharmacol* **158**, 1641–1652.
- Grasa L, Gil V, Gallego D, Martin MT & Jimenez M (2009b). P2Y₁ receptors mediate inhibitory neuromuscular transmission in the rat colon. *Br J Pharmacol* **158**, 1641–1652.
- Grover M, Bernard CE, Pasricha PJ, Parkman HP, Abell TL, Nguyen LA, Snape W, Shen KR, Sarr M, Swain J, Kendrick M, Gibbons S, Ordog T & Farrugia G (2012). Platelet-derived growth factor receptor α (PDGFR α)-expressing “fibroblast-like cells” in diabetic and idiopathic gastroparesis of humans. *Neurogastroenterol Motil* **24**, 844–852.
- Horiguchi K & Komuro T (2000). Ultrastructural observations of fibroblast-like cells forming gap junctions in the W/W^v mouse small intestine. *J Auton Nerv Syst* **80**, 142–147.
- Hwang SJ, Blair PJ, Durnin L, Mutafova-Yambolieva V, Sanders KM & Ward SM (2012). P2Y₁ purinoreceptors are fundamental to inhibitory motor control of murine colonic excitability and transit. *J Physiol* **590**, 1957–1972.
- Iino S, Horiguchi K, Horiguchi S & Nojyo Y (2009). c-Kit-negative fibroblast-like cells express platelet-derived growth factor receptor α in the murine gastrointestinal musculature. *Histochem Cell Biol* **131**, 691–702.
- Iino S & Nojyo Y (2009). Immunohistochemical demonstration of c-Kit-negative fibroblast-like cells in murine gastrointestinal musculature. *Arch Histol Cytol* **72**, 107–115.
- Klemm MF & Lang RJ (2002). Distribution of Ca²⁺-activated K⁺ channel (SK2 and SK3) immunoreactivity in intestinal smooth muscles of the guinea-pig. *Clin Exp Pharmacol Physiol* **29**, 18–25.
- Koh BH, Roy R, Hollywood MA, Thornbury KD, McHale NG, Sergeant GP, Hatton WJ, Ward SM, Sanders KM & Koh SD (2012). Platelet-derived growth factor receptor- α cells in mouse urinary bladder: a new class of interstitial cells. *J Cell Mol Med* **16**, 691–700.
- Komuro T (1999). Comparative morphology of interstitial cells of Cajal: ultrastructural characterization. *Microsc Res Tech* **47**, 267–285.
- Komuro T, Seki K & Horiguchi K (1999). Ultrastructural characterization of the interstitial cells of Cajal. *Arch Histol Cytol* **62**, 295–316.
- Kurahashi M, Nakano Y, Hennig GW, Ward SM & Sanders KM (2012). Platelet-derived growth factor receptor α -positive cells in the tunica muscularis of human colon. *J Cell Mol Med* **16**, 1397–1404.
- Kurahashi M, Zheng H, Dwyer L, Ward SM, Don Koh S & Sanders KM (2011). A functional role for the ‘fibroblast-like cells’ in gastrointestinal smooth muscles. *J Physiol* **589**, 697–710.
- Lee H, Koh BH, Peri LE, Sanders KM & Koh SD (2013). Functional expression of SK channels in murine detrusor PDGFR α ⁺ cells. *J Physiol* **591**, 503–513.
- Monaghan KP, Johnston L & McCloskey KD (2012). Identification of PDGFR α positive populations of interstitial cells in human and guinea pig bladders. *J Urol* **188**, 639–647.
- Monaghan KP, Koh SD, Ro S, Yeom J, Horowitz B & Sanders KM (2006). Nucleotide regulation of the voltage-dependent nonselective cation conductance in murine colonic myocytes. *Am J Physiol Cell Physiol* **291**, C985–C994.
- Murthy KS & Makhlof GM (1998). Coexpression of ligand-gated P_{2X} and G protein-coupled P_{2Y} receptors in smooth muscle. Preferential activation of P_{2Y} receptors coupled to phospholipase C (PLC)- β 1 via G $\alpha_{q/11}$ and to PLC- β 3 via G $\beta\gamma_{13}$. *J Biol Chem* **273**, 4695–4704.
- Mutafova-Yambolieva VN, Hwang SJ, Hao X, Chen H, Zhu MX, Wood JD, Ward SM & Sanders KM (2007). β -Nicotinamide adenine dinucleotide is an inhibitory neurotransmitter in visceral smooth muscle. *Proc Natl Acad Sci U S A* **104**, 16359–16364.
- Peppiatt CM, Collins TJ, Mackenzie L, Conway SJ, Holmes AB, Bootman MD, Berridge MJ, Seo JT & Roderick HL (2003). 2-Aminoethoxydiphenyl borate (2-APB) antagonises inositol 1,4,5-trisphosphate-induced calcium release, inhibits calcium pumps and has a use-dependent and slowly reversible action on store-operated calcium entry channels. *Cell Calcium* **34**, 97–108.
- Sanders KM (2010). Interstitial cells in smooth muscles. Review series. *J Cell Mol Med* **14**, 1197–1198.
- Sanders KM, Koh SD, Ro S & Ward SM (2012). Regulation of gastrointestinal motility – insights from smooth muscle biology. *Nat Rev Gastroenterol Hepatol* **9**, 633–645.
- Suzuki H, Ward SM, Bayguinov YR, Edwards FR & Hirst GD (2003). Involvement of intramuscular interstitial cells in nitrgergic inhibition in the mouse gastric antrum. *J Physiol* **546**, 751–763.

- Torihashi S, Ward SM, Nishikawa S, Nishi K, Kobayashi S & Sanders KM (1995). *c-kit*-dependent development of interstitial cells and electrical activity in the murine gastrointestinal tract. *Cell Tissue Res* **280**, 97–111.
- Vanderwinden JM, Rumessen JJ, de Kerchove d'Exaerde A Jr, Gillard K, Panthier JJ, de Laet MH & Schiffmann SN (2002). Kit-negative fibroblast-like cells expressing SK3, a Ca²⁺-activated K⁺ channel, in the gut musculature in health and disease. *Cell Tissue Res* **310**, 349–358.
- Velázquez B, Garrad RC, Weisman GA & Gonzalez FA (2000). Differential agonist-induced desensitization of P2Y₂ nucleotide receptors by ATP and UTP. *Mol Cell Biochem* **206**, 75–89.
- Ward SM, Beckett EA, Wang X, Baker F, Khoyi M & Sanders KM (2000). Interstitial cells of Cajal mediate cholinergic neurotransmission from enteric motor neurons. *J Neurosci* **20**, 1393–1403.
- Ward SM, Burns AJ, Torihashi S & Sanders KM (1994). Mutation of the proto-oncogene *c-kit* blocks development of interstitial cells and electrical rhythmicity in murine intestine. *J Physiol* **480**, 91–97.
- Ward SM, Morris G, Reese L, Wang XY & Sanders KM (1998). Interstitial cells of Cajal mediate enteric inhibitory neurotransmission in the lower esophageal and pyloric sphincters. *Gastroenterology* **115**, 314–329.
- Ward SM & Sanders KM (2006). Involvement of intramuscular interstitial cells of Cajal in neuroeffector transmission in the gastrointestinal tract. *J Physiol* **576**, 675–682.
- Ward SM, Sanders KM & Hirst GD (2004). Role of interstitial cells of Cajal in neural control of gastrointestinal smooth muscles. *Neurogastroenterol Motil* **16**, Suppl. 1, 112–117.
- Zhang Y, Lomax AE & Paterson WG (2010). P2Y₁ receptors mediate apamin-sensitive and -insensitive inhibitory junction potentials in murine colonic circular smooth muscle. *J Pharmacol Exp Ther* **333**, 602–611.

Additional information

Competing interests

None declared.

Author contributions

Conception and design of the experiments: S.A.B., G.W.H., S.M.W. and K.M.S. Collection, analysis and interpretation of data: S.A.B., G.W.H., A.K.S., M.K., S.M.W. and K.M.S. Drafting the article or revising it critically for important intellectual content: S.A.B., S.M.W. and K.M.S. All authors read and approved the manuscript for submission.

Funding

Funding for this study was provided by RO1 DK091336 to K.M.S. and DK57236 to S.M.W. The animal breeding, immunohistochemistry and molecular expression studies were performed by Core Labs supported by NIDDK grant P01-DK41315.

Acknowledgements

The authors are grateful to the following people for their help: Yasuko Nakano for immunohistochemistry, Lauren Peri for molecular expression studies, Byoung Koh for FACS analysis studies, and Nancy Horowitz for mice maintenance.