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1 **Altered Urothelial ATP Signaling in Major Subset of Human Overactive Bladder**
2 **Patients with Pyuria**

3
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24 **Abstract (200 words):**

25 Overactive Bladder (OAB) is an idiopathic condition, characterized by urgency,
26 urinary frequency and urgency incontinence, in the absence of routinely traceable
27 urinary infection. We have described microscopic pyuria (≥ 10 wbc μl^{-1}) in patients
28 suffering from the worst symptoms. It is established that inflammation is associated
29 with increased ATP release from epithelial cells, and extracellular ATP originating
30 from the urothelium following increased hydrostatic pressure, is a mediator of
31 bladder sensation. Here, using bladder-biopsy samples, we have investigated
32 urothelial ATP signaling in OAB patients with microscopic pyuria.

33 Basal, but not stretch-evoked, release of ATP was significantly greater from
34 urothelium of OAB patients with pyuria than from non-OAB patients or OAB patients
35 without pyuria (< 10 wbc μl^{-1}). Basal ATP release from urothelium of OAB patients
36 with pyuria was inhibited by the P2 receptor antagonist suramin and abolished by
37 the hemichannel blocker carbenoxolone, which differed from stretch-activated ATP
38 release. Altered P2 receptor expression was evident in urothelium from pyuric OAB
39 patients. Furthermore, intracellular bacteria were visualized in shed urothelial cells
40 from ~80% of OAB patients with pyuria.

41 These data suggest that increased ATP release from the urothelium, involving
42 bacterial colonization, may play a role in the heightened symptoms associated with
43 pyuric OAB patients.

44

45

46 **Introduction:**

47 Overactive bladder syndrome (OAB) is an idiopathic condition where the bladder
48 detrusor urinae muscle spontaneously contracts before the bladder is full. In the
49 USA, it is ranked in the top 10 of common chronic conditions, competing with both
50 diabetes and depression, with a reported prevalence of up to 31-42% in the adult
51 population (2).

52 OAB is currently characterized by symptoms of urgency, with or without
53 urgency incontinence, with increased frequency, and nocturia, and in some cases
54 pain, in the absence of urinary tract infection (UTI) or other defined underlying
55 pathology (54). The exclusion of infection is determined by failure to isolate $\geq 10^5$
56 colony forming units (CFU)/ml of a single species of bacteria from culture of a
57 midstream urine (MSU) specimen (23) and negative leukocyte esterase and/or
58 nitrate urinalysis by dipstick (25). Controversy exists as to whether current methods
59 used to determine UTI are fully accurate (28). It has been established that the
60 bacterial threshold of $\geq 10^5$ CFU/ml, in the presence of symptoms, is not identified in
61 approximately 50% of UTI (31). In addition, we have recently reported the low
62 sensitivity and specificity of routine urinary dipstick tests (20, 51). Interestingly, in
63 recent studies where the threshold has been reduced to $\geq 10^2$ CFU/ml, bacterial
64 cystitis has been identified in approximately one third of patients with refractory
65 OAB, suggesting bacteria may play a significant role in the aetiology of OAB, in at
66 least a subset of patients with OAB (22, 37, 49). It is generally accepted, that the
67 best indicator of UTI is the detection of ≥ 10 white blood cells (wbc) in 1 μ l of fresh,
68 un-spun, urine examined using a haemocytometer (16, 44); however nowadays this
69 is not normal clinical practice. Using this methodology, we have identified a low-
70 grade inflammatory response (pyuria with ≥ 10 wbc μ l⁻¹) in 10-35% of MSU
71 specimens from patients with OAB (*i.e.* symptoms of urgency, with or without
72 urgency incontinence, with frequency and nocturia, in the absence of UTI) (45, and
73 new data not shown). Interestingly, our observation is that these patients showed
74 the worst symptoms of frequency. Most apposite to this finding is that persistent
75 inflammation, caused by infection and thereby accompanied by pyuria, is associated
76 with increased nucleotide (primarily adenosine 5'-triphosphate [ATP]) release from
77 epithelial cells and nucleotide-activated P2 receptor signaling (4, 5, 36, 55).

78 Extracellular nucleotide signaling via P2 receptor activation is important in
79 the regulation of bladder function (3, 7, 8). Bladder stretch, during filling, induces a
80 cytosolic Ca²⁺ increase via multiple proposed pathways (9) to promote release of ATP
81 through conductive or vesicular pathways (46), and probably other nucleotides from
82 superficial urothelium. Once released, nucleotides bind to P2 receptors on
83 suburothelial sensory afferents (P2X2, 3 and/or 2/3 subtypes) to trigger nerve
84 activation and the sensation of bladder fullness and the urge to urinate (52). Given
85 the rapid breakdown of extracellular nucleotides by nucleotidases, this route should
86 involve intermediate signaling steps involving activation of P2 receptors on other
87 urothelial cells to release additional nucleotides to act in an autocrine/paracrine
88 manner (17, 19, 48, 58). Nucleotide signaling in human urothelium is enhanced in
89 interstitial cystitis (IC) in humans and felines (17, 58), a condition symptomatically
90 similar to OAB yet with the presence of pain and an inflammatory aetiology. In IC,
91 stretch-activated ATP release is significantly increased, P2R expression profiles are
92 altered in the urothelium, and ATP breakdown is purportedly decreased (15, 58).
93 Interestingly, studies on humans with acute cystitis/urinary infection demonstrate
94 that uropathogenic *E. coli* (UPEC) can thrive in the urothelium as intracellular
95 bacterial colonies resisting host immunity (29), that should be accompanied by
96 increased nucleotide release from epithelial cells (see paragraph above).

97 We hypothesize that in a subset of OAB patients (*i.e.* those with pyuria ≥ 10
98 wbc μl^{-1}) there is increased release of ATP, and other nucleotides, from the
99 urothelium caused by low-grade inflammation, which ultimately results in increased
100 sensory nerve excitation and the enhanced symptoms of OAB. Furthermore, in
101 these patients the low-grade inflammation is associated with intracellular bacteria
102 colonization of the urothelium. Here we investigated our hypothesis using human
103 bladder urothelium obtained using flexible cystoscopy.

104

105 **Methods:**

106 *Reagents.* Chemical reagents were purchased from Sigma-Aldrich (Poole, UK); with
107 the exception of suramin obtained from Bayer AG (Leverkusen, Germany) and DAPI-
108 containing microscope slide mountant obtained from Vector Labs (Peterborough,

109 UK). HPLC column and solid phase extraction cartridges were obtained from
110 Phenomenex (Macclesfield, UK).

111

112 *Human tissue and urine samples.* All procedures were performed with consent and
113 approval from the Moorfields and Whittington Hospitals Research Ethics Committee
114 (London, UK) and the NHS Research Authority South East Coast (Kent). Informed
115 written consent was obtained from volunteers involved in the study. Bladder
116 biopsies and urine samples were obtained from either the Whittington Hospital
117 Campus, University College London or Medway Maritime Hospital. Bladder biopsies
118 were obtained using a flexible cystoscope under local (or rarely general) anaesthesia.
119 A catheter specimen of urine (CSU) was obtained from female patients and a MSU
120 from male patients and non-OAB controls. Patients with idiopathic OAB were
121 recruited from Incontinence Clinics and non-OAB controls were recruited from the
122 Haematuria Clinic. The inclusion criteria for OAB, reconfirmed at the time of sample
123 collection, were adults aged ≥ 18 years of either sex with frequency ≥ 8 per day,
124 urgency with or without urgency incontinence, and the absence of a UTI or severe
125 concomitant urinary tract pathology (*e.g.* chronic obstruction, catheterisation,
126 neurological disease, prior radiotherapy, anatomical defects, implanted devices,
127 pregnancy, bleeding disorders, and/or anticoagulant therapy). All urine samples
128 obtained were sent for conventional urinalysis and bacteria culture at the time of
129 collection, if a UTI was subsequently identified the sample was retrospectively
130 removed from the study. The inclusion criteria for non-OAB were a single historical
131 episode of microscopic haematuria with no underlying pathology. All OAB and non-
132 OAB patients were additionally investigated for pyuria by trained clinicians as
133 previously described (20); significant pyuria was taken as of ≥ 10 wbc μl^{-1} of fresh un-
134 spun urine. Thus the following experimental groups were used in this investigation:
135 *i) non-OAB controls, ii) OAB patients without pyuria*, that is patients with no wbc or
136 with < 10 wbc μl^{-1} , and *iii) OAB patients with pyuria*, that is patients with ≥ 10 wbc μl^{-1} .
137 Urine was either stored on ice and processed within 1 h (for microscopy) or
138 immediately snap-frozen and kept at -80°C until the time of processing (for HPLC).
139 Bladder biopsies were either immediately transported to the laboratory in ice-cold
140 sterile saline and processed within 1 h of collection or placed in 10% formalin for 48

141 h prior to histological scrutiny.

142

143 *Bioopsy histology.* Following 48 h in 10% formalin, biopsy tissue was dehydrated
144 with alcohol and xylene in a vacuum infiltration-processing machine for 12 h. The
145 tissue was then impregnated with paraffin wax. A microtome (Microm HM355S;
146 Thermo Fisher Scientific, Loughborough, UK) was used to cut sections of 6 μm
147 thickness. Slices were placed on a glass slide then dried at 60°C for 10 min. The
148 sections were then stained with H&E to evaluate the morphological characteristics of
149 the tissue. A Leica DM4000B upright light microscope (Wetlar, Germany) was used to
150 image the sections.

151

152 *Biopsy ATP release.* A Luciferin Luciferase ATP Bioluminescence Assay Kit was used to
153 quantify ATP release from intact, live, bladder urothelium according to the
154 manufacturer's protocol. In brief, the urothelial cell layer was manually isolated
155 from the underlying tissue of the bladder biopsy using fine forceps, scalpel and a
156 dissection microscope (final wet tissue weight being 3.4 ± 0.7 mg [n=33]). Two
157 working solutions containing the luciferin luciferase reagents were made up, one in
158 PBS (phosphate buffered saline [1X] containing [mM]: Na_2HPO_4 [10], KH_2PO_4 [1.8]
159 NaCl [137], KCl [2.7], pH 7.4; Solution-A [isotonic solution]) and one with distilled
160 water (Solution-B; hypotonic solution). The intact urothelial cell layer was then
161 incubated for 1 h in Solution-A (100 μl) at 37°C in a 96-well plate (Nunc, Roskilde,
162 Denmark). Basal readings of luminescence were taken after 1 h incubation with the
163 tissue still immersed in Solution-A. Stimulated readings of luminescence (*i.e.*
164 stretch-evoked ATP release) were taken following the addition of Solution-B (100 μl)
165 for 1 min. Where stated, stimulated readings of luminescence were taken over a
166 period of 15 min with 3 min interval recordings to observe degradation of ATP with
167 time. Two ATP standard curves, one in Solution-A (100 μl volume), and the other in
168 Solution-A and Solution-B (50:50, 200 μl volume), in combination with blank wells,
169 were prepared alongside each given experiment with concentrations ranging from
170 10^{-10} to 10^{-7} M. ATP-evoked luminescence was quantified using a luminometer
171 (Synergy 2, BioTek, Winooski, USA). ATP concentration from samples was calculated

172 from the ATP standards using linear regression analysis. All data were normalised as
173 nM ATP per 5 mg of wet tissue, and stimulated readings presented following the
174 subtraction of basal readings. In experiments investigating the effect of drugs, the
175 same protocol was followed, allowing a minimum of 5 min for the drugs to take
176 effect before luminescence was read. Parallel standard curves were also run in the
177 presence of drugs to investigate any possible interactions with the luciferin
178 luciferase reaction.

179

180 *Biopsy vesicle staining.* Following immediate transport to the laboratory in ice-cold
181 sterile saline, biopsy tissue was placed in PBS (1X) and incubated for 1 h at 37 °C to
182 create *resting* conditions, or, to imitate *stretch* conditions (*i.e.* bladder filling), biopsy
183 tissue was then incubated in hypotonic PBS (0.5X) for 1 min. Both the *resting* and
184 *stretch* biopsy tissues were immediately transferred to PFA (4%) for 48 h, before
185 being wax-embedded and sectioned (as above: *Biopsy histology*). The tissue sections
186 were permeated with a 0.1% Triton X100 solution and incubated with quinacrine
187 (100 µM) for 30 min before being washed with PBS (1X). Sections were mounted
188 with DAPI-containing mountant, and visualised under an inverted confocal
189 microscope using the x63 oil immersion objective (Leica SP5; Wetzlar, Germany).
190 Using ImageJ software, mean fluorescence intensity was measured in all images
191 taken from quinacrine-labelled tissue.

192

193 *Biopsy RNA extraction and RT-PCR.* Following immediate transport to the laboratory,
194 in cold sterile saline, the urothelial cell layer was manually isolated from the
195 underlying tissue of the bladder biopsy using fine forceps, scalpel and a dissection
196 microscope. Intact urothelium was homogenised in Tri-reagent and then
197 freeze/thawed in liquid nitrogen followed by chloroform extraction and ethanol
198 precipitation. Precipitated RNA was loaded onto Qiagen RNeasy columns (Qiagen,
199 Crawley, UK) for DNase treatment and further purification. RNA concentration was
200 measured using a spectrometer, Nano N-1000 system (Nanodrop Technologies,
201 Wilmington, USA).

202 Total RNA (10 ng) from each sample was reverse transcribed in a 20 µl reaction
203 volume using the One-step Quantitect Reverse Transcription Kit (Qiagen, Crawley,

204 UK) according to manufacturer's instructions. Primers for P2 receptor subtypes were
205 designed using Primer 3 Web-software (Whitehead Institute for Biomedical
206 Research, Cambridge, USA) and manufactured by Eurofins-MWG (Ebersberg,
207 Germany) (see Appendix for primers and mRNA accession numbers used). The real-
208 time PCR, based on SyBR green detection, (Qiagen, Crawley, UK) was performed
209 using a Chromo-4 thermal cycler (Bio-Rad, Hemel Hempstead, UK) with 2 μ l total
210 RNA. Using a standard curve, created by duplicate serial dilutions of standard DNA
211 (target sequence of interest) over 12 logarithmic orders and the thermal cycler
212 software the relative concentrations of the target amplicons were determined. In
213 addition, the standard curve was used to verify the linearity of amplification of each
214 transcript; $r^2 > 0.99$ in all cases. The relative concentrations of target in each run were
215 expressed as a ratio to the housekeeping gene, GAPDH. All PCR products were
216 checked for specificity and purity from a melting curve profile created after each run
217 by the thermal cycler software. Homology of the PCR products was further checked
218 for size by agarose gel electrophoresis.

219

220 *Urine sediment immunofluorescence.* Fresh urine (50 μ l; within 1 h of collection and
221 stored on ice) was spun onto slides using a cytospin (Sandon Cytospin 4, York, UK) at
222 800 rpm for 5 min. The deposit was stained with acridine orange (0.5% in Gey's
223 solution) for 30 min; previously shown to fluoresce green in the presence of viable
224 organism DNA after excitation by a laser at 436-490 nm (12, 32). In order to
225 differentiate extracellular from intracellular bacteria, a crystal violet counter-stain
226 (0.1% in 150 mM NaCl) was added to quench the fluorescence of extracellular
227 microorganisms. Images were acquired using an upright fluorescence microscope
228 (Leica DM4000B, Wetlar, Germany), samples were excited at 488 nm and emitted
229 light collected with a 505-550 nm band pass filter. On those slides found to have
230 cells containing bacteria, the deposit was fixed with PFA (4%) for 2 min, then further
231 treated with anti-Uroplakin III (Santa Cruz Biotechnology Inc; Santa Cruz, USA) for 12
232 h, a specific marker of urothelial cells (10). Alexa 555 (Invitrogen, UK) was used as a
233 secondary antibody and incubated for 2 h, after which the slides were mounted with
234 DAPI-containing mountant. Images were acquired with a confocal microscope (Leica
235 SP5; Wetlar, Germany) and Z-series processed using Volocity software (Improvision,

236 Coventry, UK). Alexa-555 and DAPI were excited at 543 nm and 405 nm, and
237 emitted light was collected using a 560 nm long-pass filter and 420 nm long pass
238 filter, respectively.

239

240 *Urine nucleotide and nucleoside quantification.* Urine, that had previously been
241 immediately snap-frozen and kept at -80°C, was thawed and sterile filtered through
242 a 0.22 µm membrane prior to HPLC analysis. HPLC analysis was performed as
243 previously described (11). In brief, the samples were then subjected to a solid phase
244 extraction cleaning procedure through Strata-X columns (30 mg/ml; Phenomenex,
245 Macclesfield, UK). Elution was achieved using 25 mM ethanolamine at pH 5.0 and
246 30% methanol in ethanolamine (pH 5.0). A standard curve was prepared alongside
247 each HPLC experiment consisting of ADP only. Samples (100 µl) were injected into
248 the column (Polar-RP 4 µm 80A 250x4.6 mm, Phenomenex, Macclesfield, UK) and a
249 gradient profile of 2%-30% acetonitrile in phosphate buffer was run for 20 min, with
250 an additional 15 min stabilisation period. Nucleotides and nucleosides were UV-
251 detected at 254 nm and areas measured with the Agilent software (Agilent
252 Technologies, Wokingham, UK). Areas were then adjusted to the SPE cartridge
253 performance and compared against standard curves to obtain the final
254 concentration. Urinary creatinine levels were measured using a commercial test
255 (R&D Systems, Abingdon, UK) as per manufacturer's instructions to ensure data
256 were comparable.

257

258 *Statistical analysis.* All numerical data were assessed for normality using the
259 Kolmogorov-Smirnov test. Significance level was evaluated by two-tailed paired and
260 unpaired *t*-tests, parametric and non-parametric one-way ANOVA with appropriate
261 post-hoc tests. *P* values less than 0.05 were considered statistically significant. All
262 data presented as mean±SEM, and *n* equals number of patients or samples.

263

264

265 **Results:**

266 *Bladder biopsies obtained from patients using flexible cystoscopy have full-thickness*
267 *intact urothelium.* H&E staining of wax-embedded and sliced biopsies obtained using
268 flexible cystoscopy routinely demonstrated intact urothelium of full-thickness in
269 samples of all three experimental groups of patients (*i.e.* non-OAB [8 of 8
270 specimens], OAB without pyuria [11 of 13 specimens], and OAB with pyuria [6 of 6
271 specimens]) (Figure 1A). Morphologically distinct umbrella cells were evident on the
272 luminal side of the urothelium further demonstrating the integrity of the tissue
273 (Figure 1B).

274

275 *Basal ATP release is significantly greater from urothelium of OAB patients with*
276 *pyuria.* To measure ATP release from microdissected urothelium, we used a luciferin
277 luciferase assay. ATP levels became detectable after 15 min and stabilized at 30-40
278 min (data not shown) and consequently recordings were taken at 60 min. The subtle
279 increase in concentration of ATP, which stabilized with time, was taken to represent
280 basal (*i.e.* unstimulated) release of ATP. Basal release of ATP was significantly
281 greater from urothelium of OAB patients with pyuria (78.1 ± 20.6 nM/5 mg of wet
282 tissue [hereafter referred to as simply 'nM'], $n=15$, $P<0.05$) than from non-OAB
283 patients (1.9 ± 1.5 nM, $n=9$) or OAB patients without pyuria (2.2 ± 1.7 nM, $n=33$)
284 (Figure 2A). Application of a hypotonic solution, to mimic stretch and thus bladder
285 filling, resulted in substantial, and similar increase in ATP release from the
286 urothelium of all three experimental groups (non-OAB patients, 129 ± 48 nM, $n=9$;
287 OAB patients without pyuria, 38 ± 18 nM, $n=33$; OAB patients with pyuria, 268 ± 188
288 nM, $n=15$) (Figure 2B). Peak stretch-evoked ATP levels (*i.e.* the highest
289 concentration of ATP measured following stimulation) was achieved within 1 min
290 irrespective of experimental group ($n=18$; data not shown). The concentration of
291 ATP decreased during hypotonic insult suggesting degradation by endogenous tissue
292 ATPases. As expected, increasing hypotonicity caused additional ATP release from
293 urothelium (~ 2 -fold with 25% hypotonic buffer, ~ 10 -fold with 50% hypotonic buffer,
294 and ~ 20 -fold with 75% hypotonic buffer). The concentration of ATP released from
295 urothelium following osmotic insult decreased by $51 \pm 8\%$ ($n=9$) and $52 \pm 11\%$ ($n=9$)

296 after 3 min for samples from non-OAB patients and OAB patients with pyuria,
297 respectively, and levels returned to basal concentrations within 10-12 min in both
298 cases. However, for samples from OAB patients without pyuria, the decrease in
299 concentration of stimulated-ATP release was significantly less after 3 min ($26\pm 7\%$;
300 $n=9$; $P<0.05$) and returned to basal concentrations in ~ 30 min (Figure 2C). The
301 slower rate of stimulated ATP decay seen with urothelium obtained from OAB
302 patients without pyuria was similar to that with urothelium obtained from non-OAB
303 patients in the presence of the ATPase inhibitor ARL 67156 ($100\ \mu\text{M}$; $n=5$) (Figure
304 2C); ARL 67156 did not alter the peak concentration of stimulation-evoked ATP
305 release (data not shown). For ATP concentration measurements following
306 stimulation, results obtained using HPLC correlated well with results from the
307 luciferin luciferase assay (data not shown); however it was not possible to use HPLC
308 to measure basal ATP release as, in some cases, concentrations were below the level
309 of accurate quantification using the HPLC technique.

310

311 *Basal ATP release mechanisms differ from stretch-evoked stimulated release.* We
312 pharmacologically investigated the molecular mechanism(s) by which ATP is released
313 (primarily stretch-evoked) from human urothelium obtained from non-OAB patients,
314 OAB patients without pyuria, and OAB patients with pyuria, using a luciferin
315 luciferase assay. Basal ATP release from urothelium of OAB patients with pyuria was
316 significantly inhibited (by $67\pm 9\%$, $n=3$) by the P2 receptor antagonist suramin ($1\ \text{mM}$)
317 and almost abolished by the hemichannel and gap junction blocker carbenoxolone
318 (CBX, $50\ \mu\text{M}$; $n=3$), yet, was significantly potentiated (by $74\pm 13\%$, $n=3$) by the P2
319 receptor agonist UTP ($1\ \mu\text{M}$) (Figure 3A). The UTP-evoked potentiation of ATP was
320 subtly inhibited (by $\sim 30\ \text{nM}$, $n=3$) by co-incubation with CBX ($50\ \mu\text{M}$) and
321 significantly inhibited by co-incubation with suramin ($1\ \text{mM}$; $P<0.05$, $n=3$), suggesting
322 at least two mechanisms of ATP release (*i.e.* hemichannel-mediated and
323 downstream P2 receptor-evoked) (Figure 3A and 7A). Botulinum toxin-A (BTX-A; 20
324 units/ml) known to inhibit vesicular release of ATP, brefeldin-A (BFA; $20\ \mu\text{M}$) known
325 to inhibit vesicular trafficking, capsazepine ($3\ \mu\text{M}$) a blocker of stretch-activated TRP
326 channels, and 4,4'-diisothiocyano-2,2'-stilbenedisulfonic acid (DIDS, $100\ \mu\text{M}$) a

327 calcium-activated chloride channel blocker, did not significantly alter basal ATP
328 release (all n=3-4) (Figure 3A). Unfortunately, basal ATP release, as opposed to
329 stretch-evoked ATP release, from urothelium of non-OAB patients and OAB patients
330 without pyuria could not be fully investigated due to barely detectable levels of ATP
331 (see Figure 2A). However, the effects of UTP (1 μ M, n=3) were investigated in these
332 tissues and found not to significantly increase basal ATP concentration. Stretch-
333 evoked ATP release (*i.e.* release evoked by a hypotonic stimulus) from urothelium of
334 OAB patients with pyuria was significantly inhibited (by $72\pm 14\%$, n=5) by suramin (1
335 mM; $P<0.05$) and almost abolished by BTX-A (20 units/ml, n=3) and BFA (20 μ M,
336 n=3), but unaffected by capsazepine, CBX, DIDS or UTP (n=3-5) (Figure 3B).
337 Stimulated ATP release from urothelium of non-OAB patients and OAB patients
338 without pyuria, in addition to being abolished by suramin, BTX-A and BFA, was
339 significantly inhibited by capsazepine (by $67\pm 11\%$ [n=3] control and $83\pm 8\%$ [n=4]
340 OAB-pyuria; $P<0.05$).

341 Given that BTX-A abolished hypotonicity-evoked ATP release from urothelium
342 of OAB patients with pyuria (Figure 3B) we investigated whether vesicles were
343 evident in the urothelium. Quinacrine staining of wax-embedded and sliced biopsies
344 demonstrated ATP-containing vesicular structures throughout the urothelium and in
345 underlying tissue (Figure 4A). Hypotonic challenge of biopsies from OAB patients
346 with pyuria prior to wax embedding and slicing resulted in significantly less dense
347 quinacrine staining (3.9 ± 1.3 arbitrary units [AU], n=3, as compared to 0.9 ± 0.3 AU,
348 n=3), suggesting vesicle emptying following hypotonic stimulation (Figure 4B), which
349 was inhibited by the addition of BTX-A (1.3 ± 1.3 AU, n=3).

350

351 *Altered expression of P2 receptor mRNA in bladder urothelium of OAB patients.* To
352 quantify the relative abundance of P2 receptor mRNA in microdissected urothelium
353 of non-OAB patients, OAB patients without pyuria, and OAB patients with pyuria, we
354 calculated a ratio of the P2 receptor gene of interest to a constitutively expressed
355 housekeeping gene (GAPDH) using RT-PCR.

356 We failed to detect significant levels (*i.e.* >5 arbitrary units) of P2X₄ and P2Y₄
357 mRNA in urothelium from any experimental group. In contrast, significant amounts

358 of mRNA were detected for P2X₁, 2, 3, 5, 6 and 7, and P2Y_{1, 2, 6, 11, 12, 13, 14} in
359 urothelium of non-OAB controls; order of expression: P2Y₁₄>>P2X_{1, 3, 5, 6} and
360 7=P2Y_{1, 6, 11, 12 and 13}>P2X₂=P2Y₂ (Figure 5). Urothelium from OAB patients without
361 pyuria showed a significant increase in abundance of P2Y_{11 and 13} mRNA (by 200-fold
362 and 10-fold, respectively; n=6; P<0.01). Whereas, urothelium from OAB patients
363 with pyuria showed a significant increase in abundance of P2Y_{2 and 11} mRNA (100-fold
364 and 50-fold, respectively; n=6; P<0.01) (Figure 5B).

365

366 *Intracellular bacteria in shed urothelial cells of OAB patients with pyuria.* To
367 investigate whether intracellular bacteria are responsible for increased basal release
368 of ATP from urothelium from OAB patients with pyuria we stained cytopun fresh
369 urine samples with acridine orange and crystal violet. Biopsy tissue was not used in
370 this part of the investigation given its precious nature and the necessity of its use in
371 mechanistic luciferin luciferase studies. Planktonic bacteria were observed in 1 of 16
372 samples from non-OAB patients, whereas, intracellular bacteria were not observed
373 in any sedimentary cells (8±2 urothelial cells per sample, n=16) (Figure 6A).
374 Similarly, planktonic bacteria were observed in <10% of samples from OAB patients
375 without pyuria (n=33) and intracellular bacteria were not observed in any
376 sedimentary cells (Figure 6B). The number of sedimentary cells identified as
377 urothelial cells in urine samples from OAB patients without pyuria was 15±3 (n=33).
378 Planktonic bacteria were observed in 9 of 16 samples from OAB patients with pyuria,
379 and intracellular bacteria were observed in sedimentary urothelial cells from 13
380 samples (19±3 urothelial cells per sample, n=16). In the 13 samples, 52±9% of
381 urothelial cells were found to contain intracellular bacteria (Figure 6C). To confirm
382 that the cells containing intracellular bacteria were urothelial cells, the deposit was
383 fixed with PFA (4%) then further treated with anti-Uroplakin III (UP-III) and DAPI. In
384 all cases, those cells initially identified as urothelial cells by their morphology alone,
385 were confirmed as urothelial cells by positive UP-III immunofluorescence. Z-stack
386 images obtained by confocal microscopy further confirmed the intracellular
387 localization of bacteria (Figure 6D).

388

389 *Urinary AMP and adenosine levels are elevated in OAB.* Given that basal ATP release
390 is significantly greater from urothelium of OAB patients with pyuria we investigated
391 whether this was detectable by HPLC in MSU samples, as well as other nucleotides
392 (ADP, AMP, GTP, GDP, GMP, UTP, UDP and UMP) and nucleosides (adenosine,
393 guanosine and uridine) that may be altered (Table 1). Urinary AMP levels were
394 significantly greater in samples from OAB patients without pyuria ($14.4 \pm 8.3 \mu\text{M}$,
395 $n=17$, $P<0.05$) and OAB patients with pyuria ($8.5 \pm 2.3 \mu\text{M}$, $n=16$, $P<0.05$) than non-
396 OAB patients ($2.7 \pm 0.5 \mu\text{M}$, $n=11$). In addition, urinary adenosine levels were
397 significantly greater in samples from OAB patients without pyuria ($228 \pm 106 \mu\text{M}$,
398 $n=17$, $P<0.05$) than non-OAB patients ($61 \pm 58 \mu\text{M}$, $n=11$). Nucleotide and nucleoside
399 release from biopsy tissue was not investigated using HPLC given the necessity of the
400 tissue's use in mechanistic luciferin luciferase studies.

401

402

403 **Discussion:**

404 The main findings of this investigation revealed that basal release of ATP from the
405 urothelium is significantly greater for human OAB patients with pyuria ≥ 10 wbc μl^{-1}
406 than for OAB patients without pyuria (or with pyuria < 10 wbc μl^{-1}) or non-OAB
407 patients, which may account for the heightened symptoms seen in these patients
408 (36). More specifically, we present evidence that (1.) bacteria reside in some
409 urothelial cells of OAB patients with pyuria ≥ 10 wbc μl^{-1} , and (2.) a sequential
410 signaling mechanism occurs whereby (i) basal ATP release from the urothelium is via
411 hemichannels, (ii) ATP released through hemichannels acts in an autocrine/paracrine
412 manner by activating P2 receptors expressed throughout the urothelium (likely to be
413 the P2Y₂ subtype, shown here to be upregulated in OAB patients with pyuria), and
414 (iii) P2 receptor activation causes yet further ATP release from the urothelium via an
415 undetermined mechanism. Taken together, these findings lead to the proposal that
416 in a subset of OAB patients (*i.e.* OAB patients presenting with pyuria ≥ 10 wbc μl^{-1}),
417 there is heightened basal ATP release from, and increased P2 receptor expression in,
418 the urothelium originating from intracellular bacteria colonization, that culminates in
419 inappropriate sensory nerve excitation and the symptoms of OAB seen in these
420 patients (see Figure 7A).

421 In addition to reporting altered urothelial ATP release from OAB patients
422 presenting with pyuria, we also describe stretch-evoked ATP signaling in human
423 urothelium (which does not significantly differ between OAB patients without
424 pyuria, OAB patients with pyuria, and non-OAB patients). Specifically we present
425 evidence that a sequential signaling mechanism occurs whereby (i) urothelial cell
426 stretch (as occurs in bladder filling) evokes vesicular ATP release, (ii) ATP released
427 from vesicles acts in an autocrine/paracrine manner by activating P2 receptors (of
428 which a variety and abundance are expressed throughout the urothelium), and (iii)
429 P2 receptor activation causes yet further vesicular ATP release from the urothelium
430 to presumably activate P2X receptors on suburothelial sensory nerves and therefore
431 signal bladder fullness (see Figure 7B).

432

433 *Mechanisms of urothelial ATP release.*

434 It is well established that extracellular nucleotide signaling arising from the
435 urothelium is important in the regulation of bladder function (3, 8). However, to
436 date little work has been performed on human urothelium. This, in part, is because
437 of the reported fragility of the urothelial cell layer and superficial umbrella cells, and,
438 the logistic difficulty in obtaining tissue samples using the cold-cup biopsy technique
439 (normally performed under general anaesthesia). However, here we demonstrate
440 that full-thickness intact urothelium obtained using flexible cystoscopy, a routinely
441 used procedure using a fibre optic instrument and not requiring general anesthesia
442 of the donor, can be used successfully to study the physiology/pathophysiology of
443 the urothelium *in vitro*.

444 Using urothelium samples, obtained by flexible cystoscopy, we have
445 presented evidence for an autocrine/paracrine ATP signaling mechanism in human
446 urothelium that presumably culminates in sensory nerve excitation (see Figure 7).
447 Our proposed mechanisms of autocrine/paracrine ATP signaling are broadly in
448 keeping with others. Autocrine/paracrine signaling by ATP, and other nucleotides, in
449 epithelial tissues is a well documented phenomenon, especially in renal tissue where
450 basal ATP release and stretch-evoked ATP release (a function of tubular flow rate)
451 influence ion and water transport mechanisms, and may even influence intrarenal
452 blood flow (21, 35, 39, 60). With respect to the bladder, the group of Birder was the
453 first to propose that the abundance, and variety, of P2 receptors expressed
454 throughout the urothelial cell layer might serve to amplify stretch-evoked ATP
455 signaling (15). Non-neuronal cellular release mechanisms of ATP are not well
456 understood. Proposals to date involve a number of complementary pathways that
457 include transport via ATP-binding-cassette (ABC) proteins, connexin hemichannels,
458 large-diameter anion channels and exocytotic vesicular release (1, 31). Our data
459 using human urothelium demonstrate ATP release via hemichannels and vesicles
460 (see Figure 3 and 4). This investigation extends findings of earlier studies using
461 urothelium by demonstrating multiple, and different, release mechanisms for both
462 basal release (in OAB patients with pyuria) and stretch-evoked release (from all
463 experimental groups). However, disappointingly we report at least one unidentified
464 ATP release mechanism in OAB patients with pyuria that is insensitive to all
465 blockers/antagonists tried (see Figure 7). It is perhaps prudent to mention at this

466 juncture that the effect of bacteria on urothelial cell permeability has not be
467 investigated, which of course may also account for increased increased basal release
468 of ATP from the urothelium of patients with OAB and pyuria.

469 Current OAB therapies include antimuscarinic and/or BTX-A treatment. The
470 former is associated with severe side-effects and high rates of withdrawal, and the
471 latter is notoriously expensive. Our study tested the ability of suramin and BTX-A to
472 alter the ATP signaling cascades seen in urothelium from OAB patients. Suramin,
473 successfully inhibited the exaggerated basal ATP release unique to OAB patients with
474 pyuria, whereas BTX-A was without effect (see Figure 3A). However, stimulated ATP
475 release, similar in non-OAB patients and OAB patients alike, was inhibited by both
476 suramin and BTX-A (see Figure 3B). These findings suggest that BTX-A may not be a
477 suitable therapy for those suffering from OAB with pyuria, and that selective
478 targeting of urothelial ATP receptors may be an alternative pharmacological strategy
479 to currently used antimuscarinics in the treatment of OAB (irrespective of pyuria
480 status).

481

482 *P2 receptor expression in urothelium.*

483 Using real time-PCR, we molecularly characterized the P2 receptor subtypes
484 expressed in urothelium of non-OAB patients, OAB patients without pyuria, and OAB
485 patients with pyuria. We consistently demonstrated mRNA expression of P2X1-3 and
486 5-7 receptor subunits, but not P2X4, in samples from all 3 experimental groups. The
487 level of expression for each subtype did not significantly differ between
488 experimental groups. Given the ability of P2X receptors to form homomeric and
489 heteromeric ion channels, the possibility exists that up to 10 subtypes of P2X
490 receptor (P2X1, 2, 3, 5, 6, 7, 1/2, 1/5, 2/3 and 2/6) may be functionally expressed in
491 human urothelium. For P2Y receptors, we consistently demonstrated mRNA
492 expression for all subtypes (P2Y_{1, 2, 6 and 11-14}) with the exception of P2Y₄. However,
493 levels of P2Y_{2, 11 and 13} were significantly increased in OAB patients, with an increase in
494 P2Y₂ being specific to OAB patients with pyuria and P2Y₁₃ being specific to OAB
495 patients without pyuria (see Figure 5).

496 Our PCR findings demonstrating expression of almost all P2 receptors in
497 human urothelium are broadly in accordance with previous studies that collectively

498 reported all P2X (P2X1-7) and P2Y_{1, 2, 4, 6 and 11} localization/expression in native
499 urothelium of human, rat, mouse, rabbit, guinea-pig and cat and in a human
500 urothelial cell line (UROtsa cells) (4, 6, 13-15, 18, 19, 24, 40, 42, 50, 57, 61). That
501 others have not demonstrated P2Y₁₂₋₁₄ expression is perhaps due to the recent
502 discovery and cloning of these subtypes from human tissue. Our PCR investigation
503 also extends findings of earlier studies in which P2 receptor expression in the
504 urothelium is described for both human and feline IC patients, whereby P2X1 and 3
505 and P2Y₂ are decreased (6, 15).

506 Interestingly, stimulation of some P2 receptors (P2X_{2, 4 and 7}, and P2Y₆)
507 results in the release of key proinflammatory cytokines (*e.g.* IL-1 β , IL-6, IL-8 and
508 TNF α) (3, 47). IL-1, IL-6 and IL-8 have been shown to presage UTI symptoms (27, 53).
509 Here we present data suggesting that P2Y₆ expression may be increased (not quite
510 significant; Figure 5B) in the urothelium of OAB patients with pyuria, which may be
511 critical to the release of these cytokines, and amplify the cascade of events leading
512 to the heightened symptoms of OAB.

513

514 *Bacterial colonization of bladder urothelium.*

515 With our findings in mind, the most apposite series of animal experiments have been
516 reported by the Hultgren group (41, 59). Using a murine model of chronic urinary
517 infection, they demonstrated the ability of *E. coli* to colonize the superficial
518 urothelium forming intracellular bacterial colonies (IBCs). These colonies exhibit a
519 reduced susceptibility to antibiotics and host immune mechanisms. Electron
520 microscopy studies showed that colonies formed pod-like protrusions from the cell
521 wall, with the resident bacteria encased in a polysaccharide-rich matrix surrounded
522 by a protective shell. Eventually, bacteria detached from the pod and burst into the
523 bladder lumen where the escaped bacteria then infected fresh cells. However, in
524 this study we failed to identify bacteria residing within the urothelial cells lining the
525 bladder (*i.e.* in biopsy material) but instead bacteria were identified inside shed
526 urothelial cells found in urine samples. The possibility exists that the shed umbrella
527 cells were from the renal pelvis, ureters, urethra and bladder; although unlikely
528 given that the urine was obtained by CSU sampling.

529 Interestingly, Rosen and colleagues have published data that showed IBCs
530 forming in urothelial cells of patients with acute cystitis (60). UPEC
531 lipopolysaccharide (LPS) is an extremely potent activator of innate immune
532 responses acting via binding to CD14 and Toll like receptors in the bladder
533 urothelium to activate p38 MAP kinase, Ca²⁺ and cAMP signalling which in turn
534 triggers IL-6 and IL-8 production. (34, 56). Furthermore, ATP released from both
535 UPEC and infected cells can stimulate IL-8 production via P2 receptor signalling (4,
536 43, 55). IL-6 is associated with activation of the acute phase response, which
537 increases production of C-reactive protein from the liver and IL-8 acts as a
538 chemotactic factor for neutrophils (pyuria). The presence of pyuria in the absence of
539 infection (determined by failure to isolate $\geq 10^5$ colony forming units) in a subset of
540 OAB patients, which suggests bladder infection and IBC, is a relatively recent
541 discovery (44).

542

543 *Urinary nucleotides and nucleosides as biomarkers of OAB.*

544 Previous studies have utilized the luciferin luciferase assay to investigate the
545 potential for urinary ATP concentration to serve as a suitable biomarker for UTIs.
546 These studies report ATP concentrations of 5-25 nM in MSU samples collected from
547 patients testing negative for UTIs, and 112-140 nM for patients with positive culture
548 (26, 38), suggesting that ATP could be a biomarker for urinary infection. However, a
549 more recent study looking at ATP concentrations in the urine of OAB patients found
550 similar low levels of ATP in both OAB and non-OAB control groups (1.5 and 1.4 nM
551 [normalized to creatinine concentrations], respectively) (33). Using a novel HPLC
552 technique we found significantly higher levels of AMP in MSU samples from OAB
553 patients (irrespective of pyuric status) than in samples from non-OAB patients. In
554 addition, we detected elevated adenosine in OAB patients without pyuria. Whereas
555 these data are currently too preliminary to propose biomarkers for OAB, it may be
556 useful in determining additional signaling mechanisms responsible for the symptoms
557 of OAB.

558

559 *Should pyuria inform the diagnosis and treatment of OAB?*

560 Data presented here raise the question as to whether OAB with pyuria should be
561 classified and treated differently to OAB without pyuria. Considerable overlap
562 already exists between OAB and chronic prostatitis or “chronic pelvic pain
563 syndrome” and IC, also called “painful bladder syndrome”, and in all cases the
564 diagnosis, rightly or wrongly, relies on exclusion of urinary infection by routine
565 culture methods (54). Our findings suggest that OAB with pyuria has a subclinical
566 UTI component (intracellular bacteria in the urothelium), not dissimilar to recent
567 findings for IC yet without pain and inflammation (29). That significant basal ATP
568 release is novel to OAB with pyuria suggests treatment could be designed
569 accordingly. Further multi-centre investigations are required to fully answer this
570 question; which may need to wait until routine UTI detection methodologies are
571 updated.

572

573 **Summary:**

574 These data provide a better understanding of OAB aetiology and, nucleotide release
575 and signaling in bladder epithelium (in health and disease). The former may prompt
576 clinical re-classification of OAB, to aid successful diagnosis and treatment in the
577 future. The later may help identify alternative, efficacious, and acceptable
578 therapeutic treatments for the unpleasant symptoms of OAB. Conceivably, efficient
579 therapeutic strategies may be evolved to target nucleotide signaling by intravesicular
580 delivery methods to alleviate OAB symptoms and/or treat intracellular bacteria of
581 OAB with pyuria.

582

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589

590 **Competing Interests:**

591 The author(s) declare to have no competing interests.

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594 **Table 1:**

Compound	Non-OAB	OAB	OAB
		without pyuria	with pyuria
ATP	1.4±0.8 nM	4.2±3.6 nM	2.0±0.8 nM
ADP	19.0±8.5µM	11.7±2.5µM	24.2±6.5 µM
AMP	2.7±0.5µM	14.4±8.3 µM*	8.5±2.2 µM*
adenosine	61.0±58.5µM	228±86 µM*	113±42.3 µM
GTP	44.9±13.9 µM	157±92.7µM	89.8±42.3 µM
GDP	128±54.0 µM	83.2±20.0 µM	138±47.3 µM
GMP	5.5±2.0µM	20.7±10.3 µM	15.9±6.7 µM
guanosine	478±174 µM	225±81.5µM	527±176 µM
UTP	9.9±2.4 µM	22.9±14.5 µM	8.6±2.6 µM
UDP	30.6±13.9µM	20.2±6.3µM	46.5±19.3 µM
UMP	1.4±0.6 mM	734±393 µM	667± 261 µM
uridine	7.3±3.4µM	25.0±8.1 µM	10.0±4.8 µM

595

596 *Urinary AMP levels are elevated in OAB patients presenting with pyuria.* HPLC was
597 used to measure the concentration of nucleotides and nucleosides in urine samples
598 from non-OAB patients, OAB patients without pyuria or with pyuria <10 wbc µl⁻¹
599 (OAB without pyuria), or with pyuria ≥10 wbc µl⁻¹ (OAB with pyuria). Creatinine levels
600 were similar in all 3 patient groups (non-OAB, 119±25 mg/dl; OAB without pyuria,
601 132±20 mg/dl; OAB with pyuria, 121±14 mg/dl). Data are mean±SEM (n=11 for non-
602 OAB, n=16 for OAB without pyuria, n=17 for OAB with pyuria). * denotes statistical
603 difference from non-OAB patients (*P*<0.05).

604

605 **Appendix:**606 *Human P2 receptor primers and mRNA accession numbers.* Primers were designed

607 using Primer 3 Web software (Whitehead Institute for Biomedical Research,

608 Cambridge, USA) and manufactured by Eurofins-MWG (Ebersberg, Germany).

Gene	Accession No.	5'to 3' Sequence	Position
P2X1	NM_002558	S: CGCCTTCTCTTCGAGTATGA	471-491
		AS: AGATAACGCCACCTTCTTATTAC	538 – 514
P2X2	NM_170682	S: GCCTACGGGATCCGCATT	958 – 975
		AS: TGGTGGGAATCAGGCTGAAC	1024– 1005
P2X3	NM_002559	S: GCTGGACCATCGGGATCA	135 – 152
		AS: GAAAACCCACCTACAAAGTAGGA	205 – 182
P2X4	NM_002560	S: CCTCTGCTTGCCAGGTA CT C	1108– 1128
		AS: CCAGGAGATACGTTGTGCTCAA	1176– 1155
P2X5	NM_002561	S: CTGCCTGTCGCTGTT CGA	311 – 328
		AS: GCAGGCCACCTTCTTGTT	378 – 360
P2X6	AF065385	S: AGGCCAGTGTGTGGTGT TCA	488 – 507
		AS: TCTCCACTGGGCACCAACTC	555 – 536
P2X7	NM_002562	S: TCTTCGTGATGACAACTTTCTCAA	401 – 425
		AS: GTCCTGCGGGTGGGATACT	476 – 458
P2Y ₁	NM_002563	S: CGTGCTGGTGTGGCTCATT	1352 – 1370
		AS: GGACCCCGGTACCTGAGTAGA	1419 – 1399
P2Y ₂	NM_176072	S:GAACTGACATGCAGAGGATAGAAGAT	1495 – 1520
		AS: GCCGGCGTGGACTCTGT	1567 – 1551
P2Y ₄	NM_002565	S: CCGTCTGTGCCATGACA	725 – 742
		AS: TGACCCCGAGCTGAAGT	793 – 776
P2Y ₆	NM_176797	S: GCCGGCGACCACATGA	1171 – 1186
		AS: GACCCTGCCTCTGCCATTT	1227 – 1209
P2Y ₁₁	NM_002566	S: CTGGAGCGCTTCTCTTAC	511 – 530
		AS: GGTAGCGTTGAGGCTGATG	586 – 567
P2Y ₁₂	NM_022788	S: AGGTCCTCTCCACTGCTCTA	318 – 339
		AS: CATCGCCAGGCCATTTGT	385 – 368
P2Y ₁₃	NM_023914	S: GAGACTCGGATAGTACAGCTGGTA	223 – 248
		AS: GCAGGATGCCGGTCAAGA	291 – 274
P2Y ₁₄	NM_014879	S: TTCCTTTCAAGATCCTTGGTGACT	433 – 456
		AS: GCAGAGACCCTGCACACAAA	505 – 486

Figure legends:

Figure 1. *Bladder biopsies obtained using flexible cystoscopy have full-thickness urothelium.* Human bladder biopsies were obtained using a flexible cystoscope under local or general anaesthetic. Biopsies were placed in 10% formalin for 48 h prior to dehydration with alcohol and xylene, and paraffin wax embedding. Sections (6 μm thick) were stained with H&E to investigate urothelium integrity. (A) Representative microphotograph of a biopsy section from an OAB patient that presented with pyuria of $<10 \text{ wbc } \mu\text{l}^{-1}$. Full-thickness urothelium is evident in the sample (between arrowheads). (B) Representative microphotograph of a biopsy section from an OAB patient that presented with pyuria of $\geq 10 \text{ wbc } \mu\text{l}^{-1}$. Full-thickness urothelium is evident in the sample (as in A); umbrella cells lining the luminal membrane are clearly visible (arrowheads). Scale bars equal 200 μm .

Figure 2. *Greater ATP release from urothelium of OAB patients with pyuria of $\geq 10 \text{ wbc } \mu\text{l}^{-1}$.* ATP release from microdissected urothelium was measured using a luciferin luciferase assay. ATP release was first measured at rest (classified here as 'basal' release) and then after addition of a hypotonic solution (to cause cell stretch and mimic bladder filling; classified here as 'stimulated' release). Data for stimulated ATP release is presented following subtraction of basal release values. (A) Basal ATP release from urothelium of OAB patients with pyuria $\geq 10 \text{ wbc } \mu\text{l}^{-1}$ (OAB + pyuria; n=15) was significantly greater than basal ATP release from urothelium of OAB patients without pyuria, or with pyuria $<10 \text{ wbc } \mu\text{l}^{-1}$ (OAB - pyuria; n=9), or non-OAB patients ($P<0.05$; n=9). (B) Stimulated ATP release from the urothelium was not significantly different between experimental groups (non-OAB patients, n=9; OAB patients without pyuria, n=9; OAB patients with pyuria, n=15). (C) The rate at which the concentration of ATP decreased, following stimulation and in the continued presence of hypotonic solution, was greatest from urothelium of non-OAB patients (n=9) and OAB patients with pyuria (n=15). The decreased rate of stimulated ATP degradation seen from urothelium of OAB patients without pyuria (n=9), was similar to that from urothelium of non-OAB patients in the presence of the ATPase inhibitor ARL 67156 (100 μM , n=5). Data shown are mean \pm SEM. * denotes significant difference from non-OAB, where $P<0.05$. In (A) and (B) data were compared by using one-way ANOVA and Dunnett's post-hoc test against non-OAB (the identified control); in (C) data were compared by using two-way ANOVA (*time after stimulation* being the repeated measures factor, and the between factor being the *patient conditions*) and Dunnett's post-hoc test against non-OAB (identified control).

Figure 3. *Mechanisms of ATP release from urothelium of OAB patients with pyuria.* ATP release from microdissected urothelium of OAB patients with pyuria $\geq 10 \text{ wbc } \mu\text{l}^{-1}$ (OAB + pyuria) was measured using a luciferin luciferase assay. ATP release was first measured at rest ('basal') or after addition of a hypotonic solution ('stimulated'). Data for stimulated ATP release is represented following subtraction of basal release values. Urothelium was pre-treated

with the stated drug for 5 min prior to measuring ATP release (basal or stimulated). (A) Basal ATP release from urothelium was significantly attenuated by both suramin (1 mM, n=3) and carbenoxolone (CBX, 50 μ M, n=3), and significantly potentiated by UTP (1 μ M, n=3) ($P<0.05$ in all cases). The potentiating effects of UTP were inhibited by co-treatment with either CBX or suramin. (B) Stimulated ATP release from the urothelium was significantly attenuated by suramin (n=5), botulinum toxin-A (BTX-A, 20 units/ml, n=3) or brefeldin-A (BFA, 20 μ M, n=3) ($P<0.05$ in all cases). Co-treatment of either BFA or suramin with UTP (n=3 in both cases) had no further effect. Data shown are mean \pm SEM. * denotes $P<0.05$. In (A) and (B) data were compared by using one-way ANOVA and Dunnett's post-hoc test against non-OAB (the identified control).

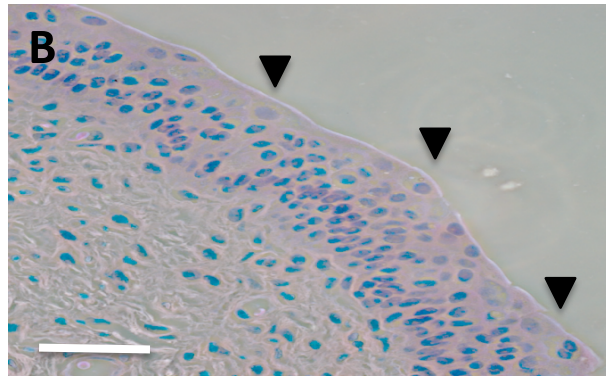
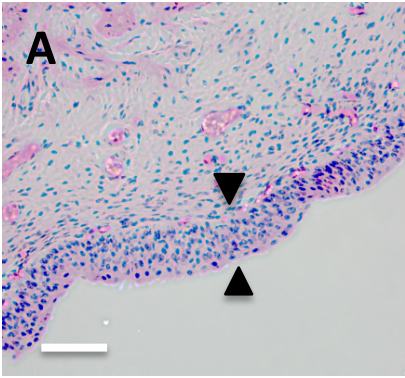
Figure 4. *Urothelial cell stretch evokes vesicular release of ATP.* Human bladder biopsies were obtained using a flexible cystoscope under local anaesthetic. Biopsies were placed in 10% formalin for 48 h prior to dehydration with alcohol and xylene, and paraffin wax embedding. In some cases, biopsies were challenged with a hypotonic solution (to cause cell stretch and mimic bladder filling) prior to fixation. Sections (6 μ m thick) were stained with DAPI and quinacrine to investigate localisation of ATP-containing vesicles. (A) Representative microphotograph of a biopsy section from an OAB patient with pyuria ≥ 10 wbc μ l⁻¹; i) DAPI staining (blue), ii) quinacrine staining (green), iii) composite image of DAPI and quinacrine staining, iv) bright field image. ATP containing vesicles are evident throughout the urothelium. (B) Representative microphotograph of a biopsy section from an OAB patient with pyuria ≥ 10 wbc μ l⁻¹, challenged with a hypotonic solution prior to fixation; i) - iv) as above. ATP containing vesicles are less evident throughout the urothelium. Scale bars equal 100 μ m.

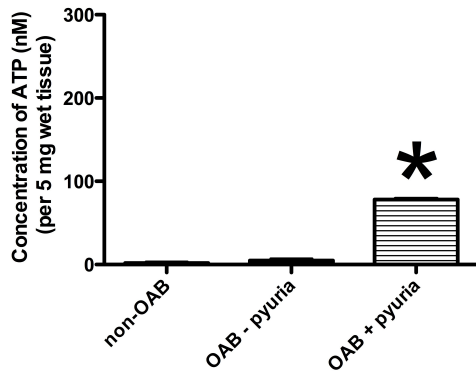
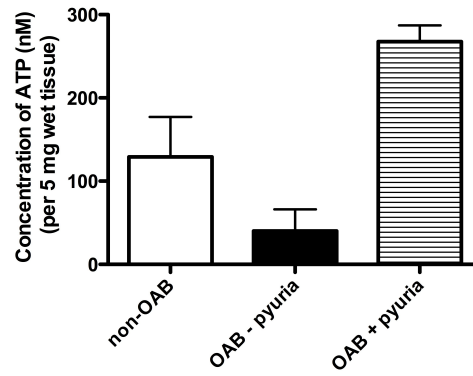
Figure 5. *Altered P2Y receptor expression in urothelium of OAB patients.* RT-PCR was performed on microdissected urothelium of non-OAB patients, OAB patients without pyuria, or with pyuria <10 wbc μ l⁻¹ (OAB – pyuria), and OAB patients with pyuria ≥ 10 wbc μ l⁻¹ (OAB + pyuria). The ratio of the P2 receptor gene of interest to a constitutively expressed housekeeping gene (GAPDH) was calculated. (A) Ionotropic P2X receptor mRNA levels were similar in urothelium from the three experimental groups (n=6). (B) Urothelium from OAB patients without pyuria showed a significant increase in abundance of metabotropic P2Y_{11 and 13} mRNA (n=6; $P<0.01$). Whereas, urothelium from OAB patients with pyuria showed a significant increase in abundance of metabotropic P2Y_{2 and 11} mRNA (n=6; $P<0.01$). Data shown are mean \pm SEM. * denotes significant difference from non-OAB, where $P<0.01$. In (A) and (B), data were compared by using one-way ANOVA and Dunnett's post-hoc test against basal or stimulated, respectively (the identified controls).

Figure 6. *Intracellular bacteria identified in shed urothelial cells from OAB patients with pyuria ≥ 10 wbc μl^{-1} .* Fresh urine was spun onto slides using a cytospin and the deposit stained with acridine orange (which fluoresces green in the presence of viable organism DNA). In order to differentiate extracellular from intracellular bacteria, a crystal violet counter-stain was added to quench the fluorescence of extracellular microorganisms. Images were acquired using an upright fluorescence microscope. Intracellular bacteria were not observed in sedimentary cells from the urine of non-OAB patients (see A for a representative image) or OAB patients without pyuria, or with pyuria < 10 wbc μl^{-1} (OAB – pyuria; see B for a representative image). (C) Representative microphotograph of shed urothelial cell colonized by intracellular bacteria (arrow) from an OAB patient with pyuria ≥ 10 wbc μl^{-1} (OAB + pyuria). Bacterial colonization of urothelial cells was seen in 81% of urine samples tested (n=16), and of those, $52 \pm 9\%$ of urothelial cells contained bacteria (19 ± 3 urothelial cells per sample). Further confirmation of the intracellular localisation of bacteria and cell type was obtained by treatment with anti-uroplakin III (red; a marker of urothelial cells) and DAPI (blue; marker of DNA). Images were acquired with a confocal microscope and Z-series processed using Volocity software. (D) Representative 2D microphotograph of three shed urothelial cells from urine of an OAB patient with pyuria (Di), whereby the intracellular localisation of bacteria is evident in the side projections (Dii and Diii; dashed line shows origin of side-projection). Scale bars equal 10 μm .

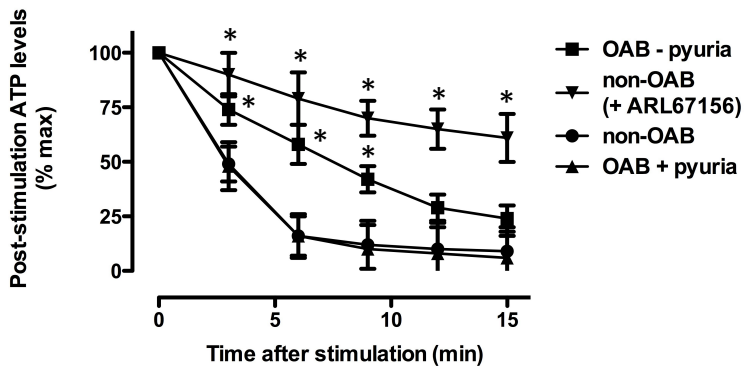
Figure 7. *Proposed mechanisms of ATP release from urothelium of OAB patients with pyuria.* In our investigations, basal release of ATP was ~ 15 -fold greater from urothelium of OAB patients with pyuria, nearing levels normally associated with stretch and bladder fullness. Histology demonstrates that bacteria reside in urothelial cells, and RT-PCR suggest that levels of P2Y₂ and P2Y₁₁ are increased. (A) When the bladder is at rest there is a substantial release of ATP (basal release) from the urothelium, which in our experiments is abolished by CBX, suggesting release via hemichannels (1.). We propose that the released ATP acts in a paracrine/autocrine fashion to activate P2 receptors expressed throughout the urothelium (including the upregulated P2Y₂ and P2Y₁₁ subtypes) (2.). P2 receptor activation in turn evokes further ATP release (in our experiments attenuated by suramin and potentiated by UTP) via an undetermined mechanism (3.). Presumably, released ATP reaches levels able to activate P2 receptors expressed on suburothelial sensory nerves, resulting in inappropriate signalling normally associated with bladder fullness (4.). We hypothesise that the presence of intracellular bacteria (IB) is responsible for increased P2 receptor expression and hemichannel mediated ATP release. (B) When the urothelium is stretched (*i.e.* when the bladder is full) there is a substantial release of ATP (stimulated release), which in our experiments is abolished by BTX-A or BFA, suggesting release from vesicles (1.). We propose that the released ATP acts in a paracrine/autocrine fashion to activate P2 receptors expressed throughout the urothelium (2.). P2 receptor activation in turn evokes further ATP release (in our experiments attenuated by suramin) again from vesicles (3.). Presumably, released ATP reaches

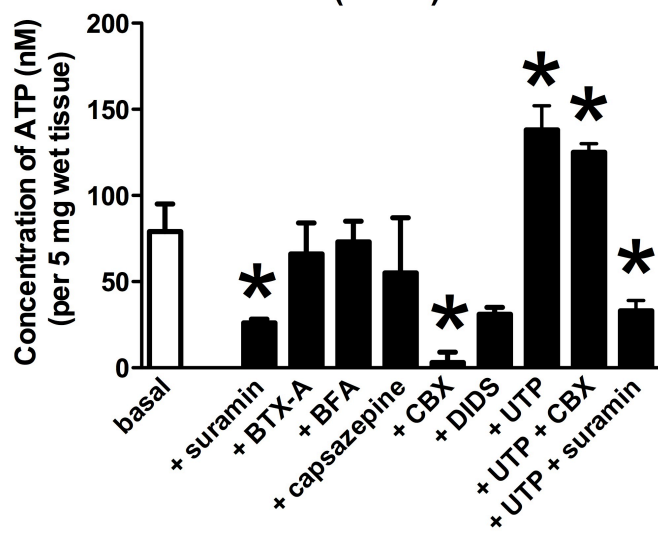
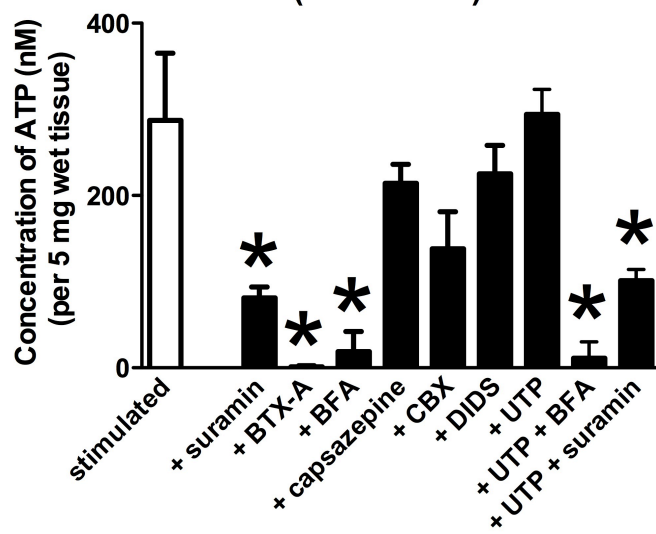
levels able to activate P2 receptors expressed on suburothelial sensory nerves, resulting in signalling of bladder fullness (4.). The presence of intracellular bacteria (IB) does not appear to alter stimulated ATP release from urothelium.

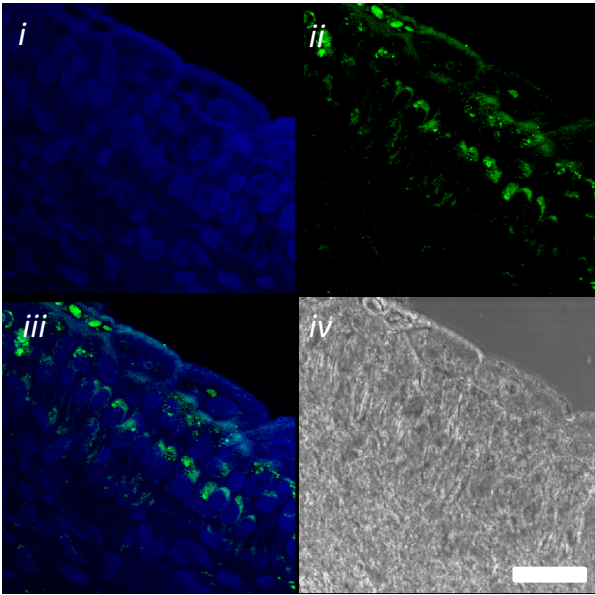
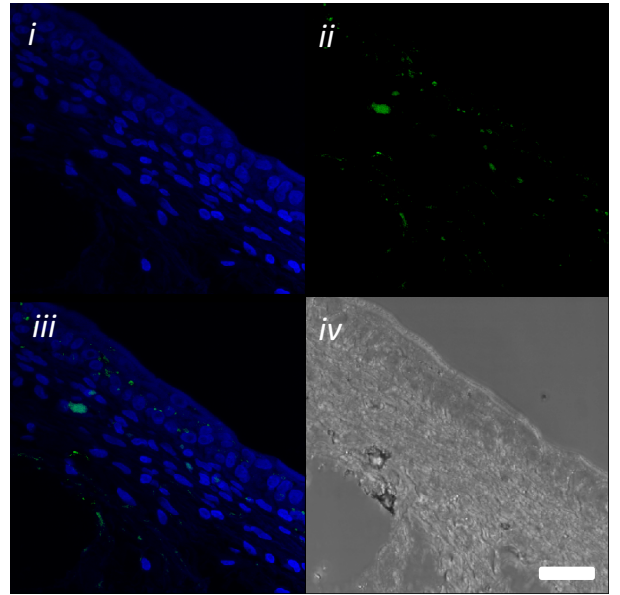


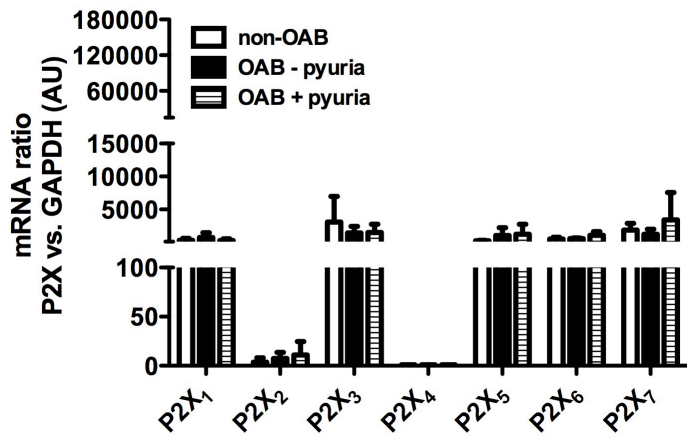
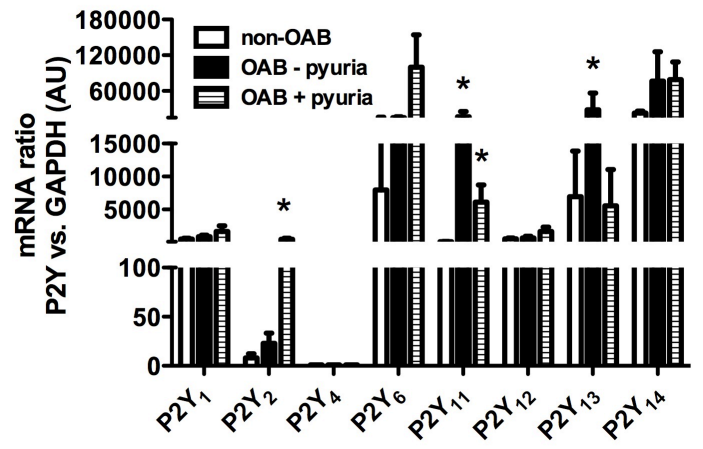
AATP release at rest
(basal)**B**Stretch-evoked ATP release
(stimulated)**C**

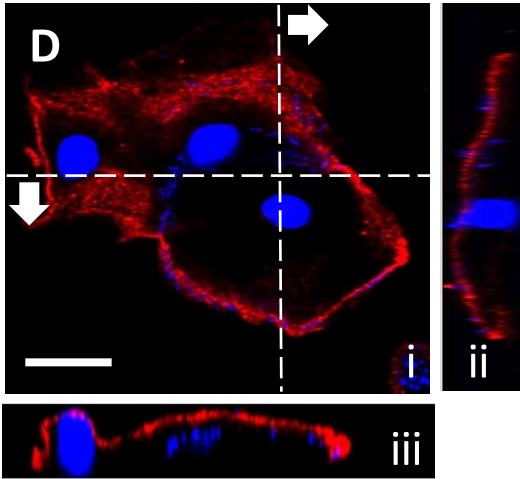
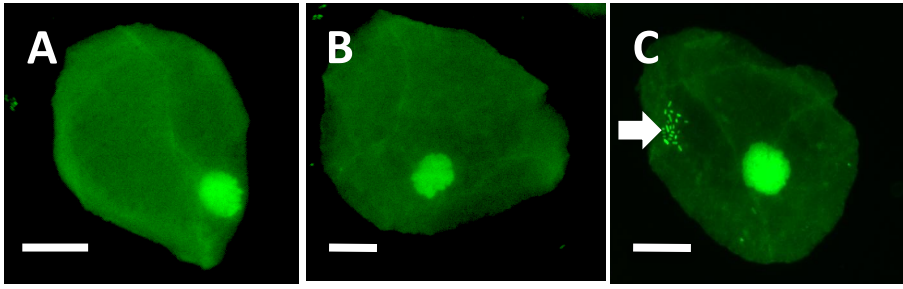
ATP decay (post stimulation)

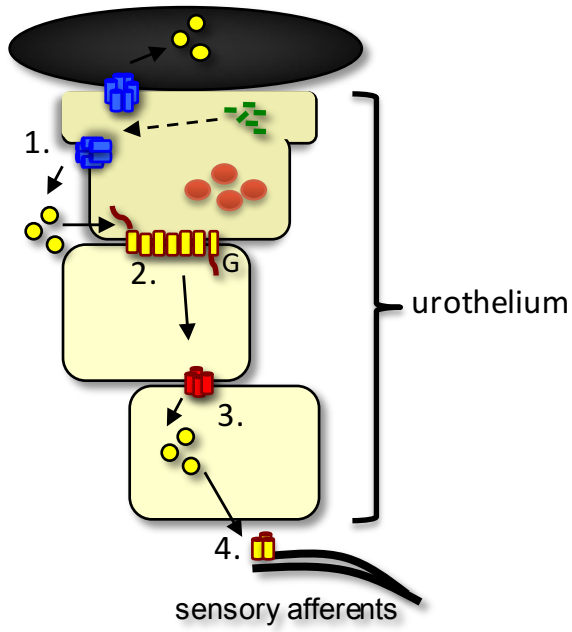
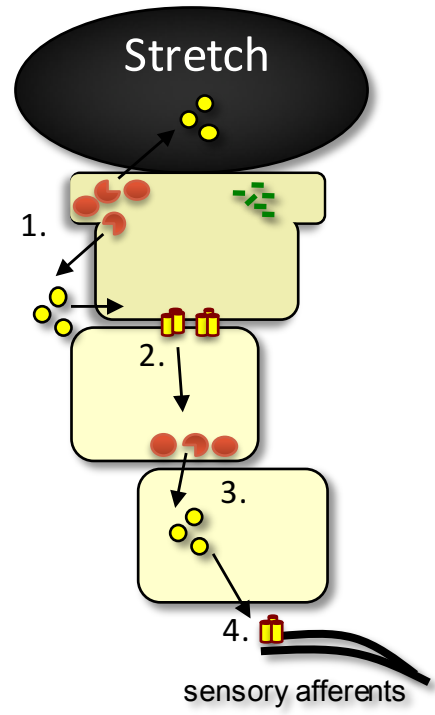









A**ATP release at rest
(basal)****B****Stretch-evoked ATP release
(stimulated)**

A**B**

A**B**



A**B**

Key: , intracellular bacteria; , hemichannel; , ATP; , P2Y receptor; , undetermined ATP channel; , P2X receptor; , ATP-containing vesicle.