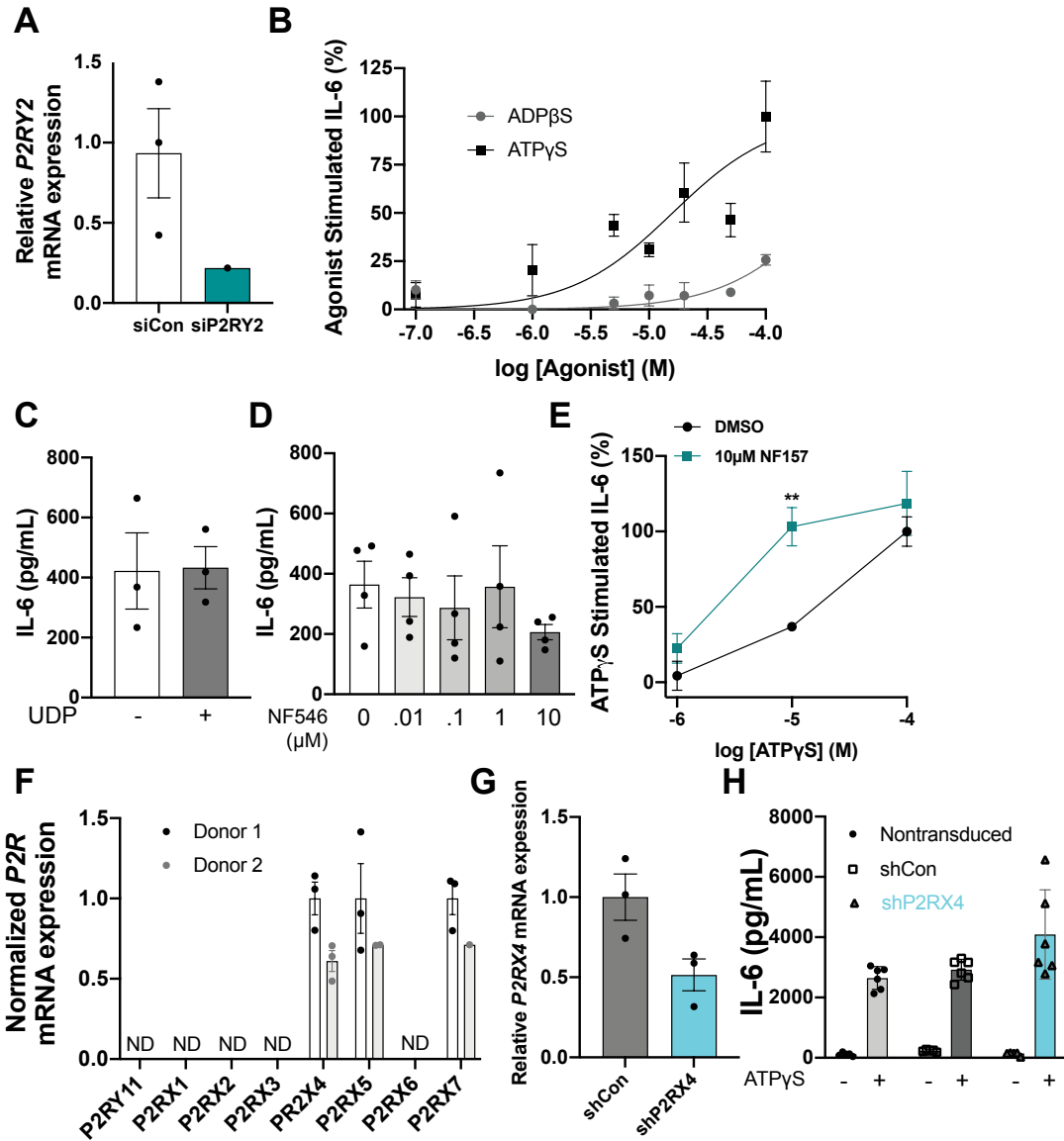
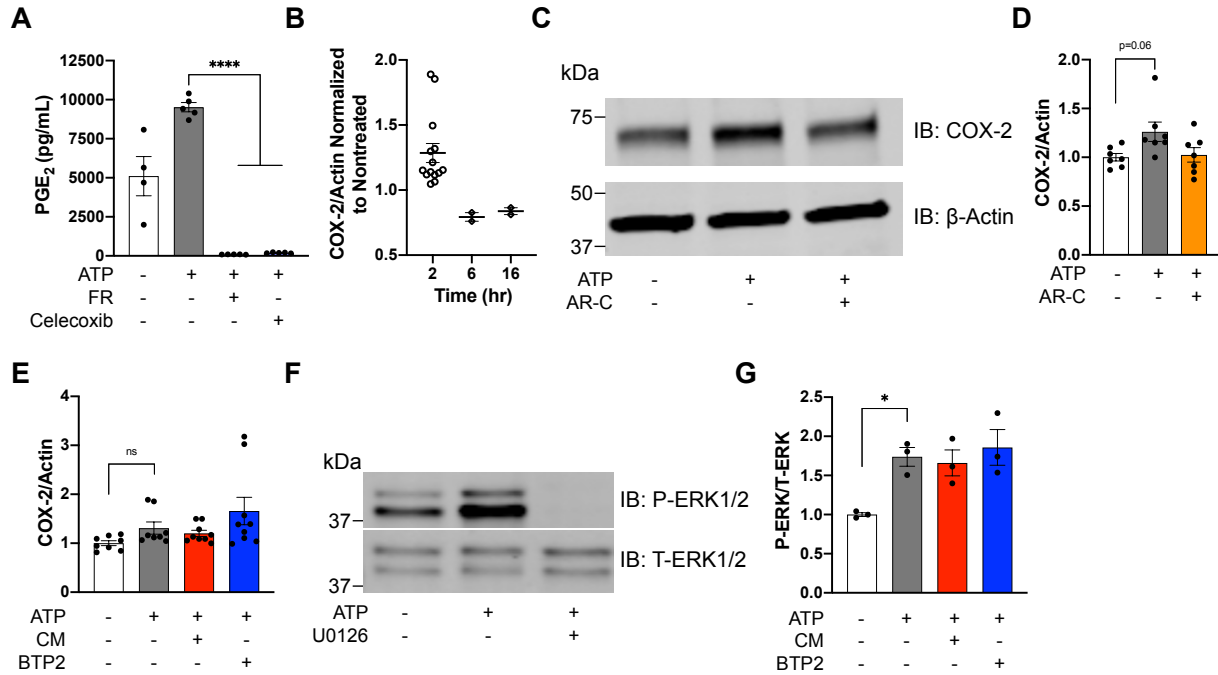


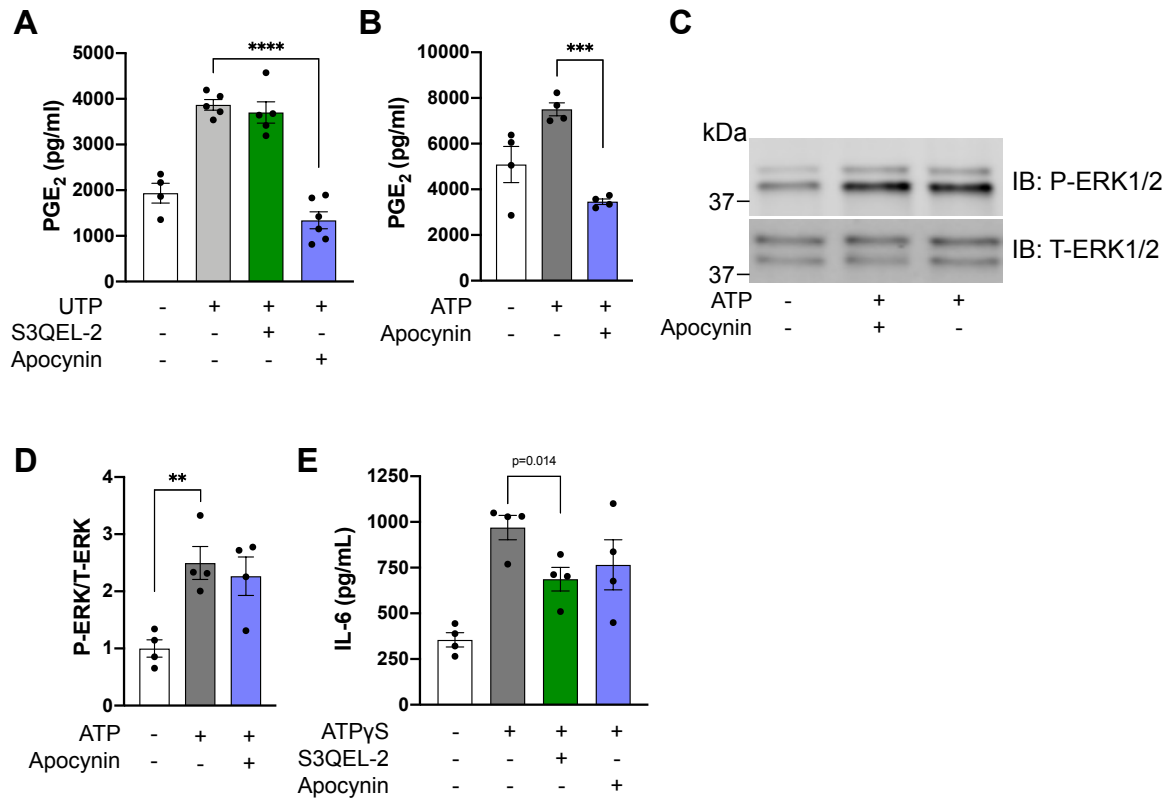
**Supplemental Figure 1. Extracellular ATP evokes  $Ca^{2+}$  influx through CRAC channels.** **A)** *siORAI1* decreases *ORAI1* mRNA expression. Expression was normalized to the three housekeeping genes *RPLP0*, *PPIA*, *HPRT1*. Data are mean  $\pm$  SEM of  $n = 2-3$  samples. **B)** *siSTIM1* decreases STIM1 protein expression. STIM1 protein was detected using a polyclonal antibody (See Methods). **C)** ATP-induced  $Ca^{2+}$  influx is inhibited by pretreatment (2hrs) with the ORAI1 inhibitor CM4620 (1  $\mu$ M). Cells were treated with ATP (50  $\mu$ M) in  $Ca^{2+}$ -free Ringer's solution to allow for store release followed by perfusion of Ringers containing 2 mM  $Ca^{2+}$ . Data are mean  $\pm$  SEM of  $n = 14-21$  cells. **D)** Quantification of the rate of  $Ca^{2+}$  influx in **C**. Rate of influx was calculated for the 24 seconds immediately following extracellular  $Ca^{2+}$  addition. Each data point is the mean  $Ca^{2+}$  influx rate for a given experiment (one dish) and graph is mean  $\pm$  SEM of  $n = 3$  independent tests. \* $p < 0.05$ .



**Supplemental Figure 2. P2Y receptor activation fails to elicit IL-6 release from AECs.** **A)** *siP2RY2* decreases *P2RY2* mRNA expression. Expression was normalized to the housekeeping gene *RPLP0*. Data are mean  $\pm$  SEM of  $n = 1-3$  samples. **B)** Dose-response for ATP $\gamma$ S versus ADP $\beta$ S induced IL-6 secretion measured in cell culture supernatant 16 hrs following agonist addition. Data are mean  $\pm$  SEM of  $n = 4-6$  samples. **C)** UDP (a P2Y<sub>6</sub> receptor agonist) does not elicit IL-6 secretion. Data are mean  $\pm$  SEM of  $n = 3$  samples. **D)** NF546 (P2Y<sub>11</sub> receptor agonist) fails to elicit IL-6 secretion. Data are mean  $\pm$  SEM of  $n = 4$  samples. **E)** NF157 (an antagonist of P2Y<sub>11</sub> and P2X<sub>1</sub> receptors) enhances low dose ATP $\gamma$ S-induced IL-6 secretion. Data are mean  $\pm$  SEM of  $n = 3$  samples. **F)** Two independent donors (NHBE cells) were examined for P2R expression via RT-qPCR. P2RX<sub>4,5,7</sub> transcripts were detected while P2RY<sub>11</sub>, P2RX<sub>1-3,6</sub> were not seen. **G)** *P2RX4* mRNA knockdown using shRNA quantified via RT-qPCR. Expression was normalized to the housekeeping gene *RPLP0*. Data are mean  $\pm$  SEM of  $n = 3$  samples. **H)** Knockdown of *P2RX4* does not affect ATP $\gamma$ S-evoked IL-6 synthesis. Data are mean  $\pm$  SEM of  $n = 5-6$  samples. \*\* $p < 0.01$



**Supplemental Figure 3. P2Y2 receptor-mediated PGE<sub>2</sub> synthesis requires COX-2 activity but not increase in COX-2 expression** **A**) PGE<sub>2</sub> synthesis induced by ATP (100 μM) is abolished with the COX-1 inhibitor FR122047 (1 μM) and the by the COX-2 inhibitor Celecoxib (1 μM). Cells were pretreated for 1 hr prior to agonist stimulation. Data are mean ± SEM of n = 4-5 samples. **B**) Time course for ATP (100 μM)-mediated COX-2 upregulation. Data are mean ± SEM of n = 2-14 samples. **C**) ATP (100 μM, 2hrs) induces only a small extent of COX-2 upregulation which is inhibited by AR-C 118925XX (10 μM). **D**) Summary of COX-2 upregulation using densitometry analysis. Data are mean ± SEM of n = 7 samples. **E**) CRAC channel inhibitors CM4620 (1 μM) and BTP2 (1 μM) do not alter ATP (100 μM) induced COX-2 upregulation (2hr stimulation duration). Data are mean ± SEM of n = 7-9 samples and densitometry was performed for quantification. **F**) Pretreatment of cells with the MEK1/2 inhibitor U0126 (20 μM) inhibits ATP (100 μM) induced ERK1/2 phosphorylation. **G**) CRAC channel inhibitors CM4620 (1 μM) and BTP2 (1 μM) do not alter ATP induced ERK1/2 activation (100 μM, 15min stimulation). Data are mean ± SEM of n = 3 samples and densitometry was performed for quantification. \*p<0.05, \*\*\*\*p<0.0001



**Supplemental Figure 4. Inhibition of NADPH oxidase occludes ATP- and UTP- induced synthesis of PGE<sub>2</sub>.** **A)** Apocynin (200μM), an inhibitor of NADPH oxidase, abrogates UTP (50μM) induced PGE<sub>2</sub> synthesis. By contrast, the mitochondrial complex III inhibitor S3QEL-2 (20μM) does not affect the UTP (50μM)- induced PGE<sub>2</sub> response. Data are mean ± SEM of n = 4-6 samples. **B)** Likewise, apocynin (200μM) abrogates ATP (50μM)-induced PGE<sub>2</sub> synthesis. Data from the first two bars are the same as the control responses shown in Figure 5B. Data are mean ± SEM of n = 4 samples. **C)** Pretreatment of cells with the apocynin (200μM) does not inhibit ATP (100μM)-induced ERK1/2 phosphorylation (15min stimulation). **D)** Densitometry to quantify experiments shown in C. Data are mean ± SEM of n = 4 samples. **E)** Inhibition of NADPH oxidase with apocynin (200μM) or the mitochondrial complex III inhibitor S3QEL-2 (20μM) have only modest effects on ATPγS (100μM) induced IL-6 synthesis. Data are mean ± SEM of n = 4 samples. \*\*p<0.01, \*\*\*p<0.001, \*\*\*\*p<0.0001.