ABC transporters control ATP release through cholesterol-dependent volume-regulated anion channel activity

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A list of materials included

Figure S1. The exonuclease inhibitor did not alter hypotonicity-induced calcium responses

Figure S2. M_βCD treatment reduces filipin fluorescence intensity

Figure S3. The cellular cholesterol level controls ATP release

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Figure S1. The exonuclease inhibitor did not alter hypotonicity-induced calcium responses *A* and *B*, HEK cells transfected transiently with RFP (*A*) or ABCG1 (*B*) were preincubated with an exonuclease inhibitor, 100 μ M ARL67156, and calcium responses were measured after hypotonic stimulation (final, 250 mmol/kg). The exonuclease inhibitor did not alter hypotonicity-induced calcium responses. Traces and quantifications of peak calcium responses (Δ F/F) with error bars are shown (n = 4). Data are mean ± standard deviation.



Figure S2. M_βCD treatment reduces filipin fluorescence intensity

HEK293 cells were treated with 5 mM methyl-beta cyclodextrin (M β CD) for one hour at 37°C for cholesterol depletion. Filipin staining was performed using the cell-based cholesterol detection assay kit (Cayman Chemical) before images were taken using a DeltaVision microscope equipped with an oil Plan Apo N 60x/1.42 NA objective (Olympus). *A*, representative images of filipin fluorscence in HEK cells treated with vehicle or M β CD *B*, filipin fluorescence intensity was quantified by line scanning of individual HEK cells (18 cells from 3 experiments each). M β CD treatment reduced filipin signal. Data are mean ± standard deviation; unpaired t test (*B*); ***p < 0.001.



Figure S3. The cellular cholesterol level controls ATP release

ATP release induced by hypotonicity (final, 250 mmol/kg) from HEK cells incubated with various concentrations of methyl-beta-cyclodextrin (M β CD) for cholesterol depletion or cholesterol mixed with M β CD for cholesterol repletion (n=6).). One-way ANOVA followed by Tukey's test with pairwise comparisons made to vehicle alone (0 mM); F(8,45) = 204.3, p < 0.001. Data are mean ± standard deviation; *p < 0.05, ***p < 0.001



