Extracellular Nucleotides Inhibit Insulin Receptor Signaling, Stimulate Autophagy and Control Lipoprotein Secretion

Cynthia Chatterjee, Daniel L. Sparks*

Atherosclerosis, Genetics and Cell Biology Group, University of Ottawa Heart Institute, Ottawa, Ontario, Canada

Abstract

Hyperglycemia is associated with abnormal plasma lipoprotein metabolism and with an elevation in circulating nucleotide levels. We evaluated how extracellular nucleotides may act to perturb hepatic lipoprotein secretion. Adenosine diphosphate (ADP) (>10 μ M) acts like a proteasomal inhibitor to stimulate apoB100 secretion and inhibit apoA-I secretion from human liver cells at 4 h and 24 h. ADP blocks apoA-I secretion by stimulating autophagy. The nucleotide increases cellular levels of the autophagosome marker, LC3-II, and increases co-localization of LC3 with apoA-I in punctate autophagosomes. ADP affects autophagy and apoA-I secretion through P2Y₁₃. Overexpression of P2Y₁₃ increases cellular LC3-II levels by ~50% and blocks induction of apoA-I secretion. Conversely, a siRNA-induced reduction in P2Y₁₃ protein expression of 50% causes a similar reduction in cellular LC3-II levels and a 3-fold stimulation in apoA-I secretion. P2Y₁₃ gene silencing blocks the effects of ADP on autophagy and apoA-I secretion. A reduction in P2Y₁₃ expression suppresses ERK1/2 phosphorylation, increases the phosphorylation of IR- β and protein kinase B (Akt) >3-fold, and blocks the inhibition of Akt phosphorylation by TNF α and ADP. Conversely, increasing P2Y₁₃ expression significantly inhibits insulin-induced phosphorylation of insulin receptor (IR- β) and Akt, similar to that observed after treatment with ADP. Nucleotides therefore act through P2Y₁₃, ERK1/2 and insulin receptor signaling to stimulate autophagy and affect hepatic lipoprotein secretion.

Citation: Chatterjee C, Sparks DL (2012) Extracellular Nucleotides Inhibit Insulin Receptor Signaling, Stimulate Autophagy and Control Lipoprotein Secretion. PLoS ONE 7(5): e36916. doi:10.1371/journal.pone.0036916

Editor: Claude Beaudoin, Blaise Pascal University, France

Received December 20, 2011; Accepted April 16, 2012; Published May 10, 2012

Copyright: © 2012 Chatterjee, Sparks. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Funding: This work was supported by a grant from the Heart and Stroke Foundation of Ontario (DLS). CC is the recipient of a Government of Ontario Graduate Scholarship. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

* E-mail: dsparks@ottawaheart.ca

Introduction

Chronic hyperglycemia in insulin resistance is known to increase the risk of cardiovascular disease and to be associated with elevated plasma apoB100 and low HDL levels [1,2]. Elevated blood glucose is also known to stimulate nucleotide secretion and purinergic signaling [3,4]. Under stress or injury, blood and vascular cells release nucleotides, such as ATP and ADP [5,6]. Extracellular nucleotide concentration in the bloodstream is normally in the nM-μM range [7,8], but can increase significantly in disease states [5,9,10]. Purinergic signaling events stimulate mitogen-activated protein kinase (MAPK) pathways and trigger the release of pro-inflammatory cytokines [6,11,12]. Extracellular nucleotides thereby directly impact the development of cardiovascular disease by promoting an "injury response" in circulating blood cells and vascular tissues [11–13].

Extracellular nucleotides affect hepatic lipoprotein metabolism through membrane G-protein coupled receptors (GPCR) [14,15]. Compounds that stimulate HDL secretion from the liver appear to act through an inhibition of nucleotide signaling. Niacin has been shown to act through GPCR pathways to stimulate the secretion of HDL [16,17] and niacin is thought to inhibit the cellular degradation of apoA-I through an inhibition of nucleotide signaling [18]. We have shown that linoleic acid phospholipids (i.e. DLPC) also act through nucleotide signaling pathways to stimulate HDL secretion [19]. These phospholipids uniquely affect

MAPK and protein kinase B (Akt) signaling [20] to block apoA-I degradation in liver cells [21].

Factors that stimulate or inhibit HDL secretion from the liver appear to have the opposite effect on the secretion of the LDL protein, apoB100. ApoB100 secretion from liver cells is regulated by protein folding and proteasomal degradation [22,23] and proteasomal inhibitors are known to stimulate the secretion of apoB100 [23]. Proteasomal inhibitors also stimulate cellular autophagic pathways [24,25]. Autophagy is an adaptive cellular "stress response" that promotes the lysosomal degradation of cytosolic components when a cell is stimulated by stressors, i.e. nutrient deprivation, extracellular signals, hormones, cytokines and pathogens [26,27]. Autophagy is designed to protect the cell by eliminating harmful cellular components through catabolism and recycling. Nucleotides act much like proteasomal inhibitors to stimulate apoB100 secretion and autophagy. The nucleotide, adenosine diphosphate (ADP), significantly increases apoB100 secretion from liver cells and increases the levels of the autophagy marker, microtubule-associated protein 1 light chain 3 (LC3-II). Autophagy has been shown to be associated with cardiovascular disease and studies suggest that excessive autophagy can lead to cardiac hypertrophy and heart failure [28,29]. Pharmacological intervention to regulate cellular autophagy may therefore have therapeutic value in the treatment of cardiovascular disease.

This study shows that ADP acts through the specific GPCR, $P2Y_{13}$, to stimulate autophagy and block HDL secretion. While stimulation in purinergic signaling would be expected to affect cellular autophagy through MAPK pathways [26,30], we now show that ADP also acts through $P2Y_{13}$ to block insulin receptor (IR- β) signaling and prevent the activation of Akt. The inhibition of insulin signaling pathways and Akt phosphorylation are known to stimulate autophagy [26,27]. ADP therefore stimulates autophagy and inhibits HDL secretion by both a stimulation of MAPK and inhibition of Akt. The study suggests that elevations in circulating nucleotide levels in hyperglycemic states may affect hepatic lipoprotein secretion through a stimulation in purinergic signaling and a coordinated regulation of both proteasomal and autophagic protein degradation.

Materials and Methods

Reagents

Dilinoleoylphosphatidylcholine (DLPC) was obtained from Avanti Polar Lipids (Alabaster, AL). Adenosine 5'-diphosphate sodium salt (ADP), adenosine triphosphate (ATP), chloroquine diphosphate salt as well as the PI3 kinase inhibitors, 3methyladenine (3-MA) and Wortmannin were purchased from Sigma-Aldrich (Oakville, ON). The antibody to P2Y13 was obtained from Abcam (Cambridge, MA). The LC3 polyclonal antibody was purchased from MBL International (Woburn, MA). Antibodies to phosphorylated ERK1/2 (p44/p42), phosphorylated Akt (Ser473), phosphorylated mTOR (Ser2448), phosphorylated IR- β (Tyr1345) and β -actin, as well as the mTOR inhibitor, rapamycin, were all obtained from Cell Signaling Technology (Danvers, MA). Human TNFa was purchased from Calbiochem (San Diego, CA). The monoclonal antibody to human apoA-I was purchased from Meridian Life Sciences, Inc (Saco, ME). The antibody to apoB (1D1) was obtained from Dr. Milne and Dr.



Figure 1. Extracellular nucleotides block the induction of apoA-I secretion. HepG2 cells were pre-treated with adenosine diphosphate (ADP) (10 to 100 μ M)) for 30 min. and then incubated with 12 μ M DLPC in serum-free DMEM media. (**A**) Conditioned media was collected after 24 h and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the media is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.01 vs ADP, **P<0.001 vs Control, ***P<0.001 vs DLPC. (**C**) Cell lysates were collected after 4 h treatment and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the cell lysate is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.01 vs ADP, **P<0.001 vs Control, ***P<0.001 vs DLPC. (**C**) Cell lysates were collected after 4 h treatment and apoA-I concentration in the cell lysate is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments.

doi:10.1371/journal.pone.0036916.g001



Figure 2. ADP stimulates apoB100 and apoE secretion. (**A&B**) HepG2 cells were incubated with 100 μ M adenosine diphosphate (ADP) for 24 h in serum-free DMEM media. Conditioned media was collected and immunoblotted for apoB100 (**A**) and apoE (**B**). Histograms represent band densitometry analysis of apoB100 or apoE, normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.01 vs Control. (**C&D**) HepG2 cells were treated with 100 μ M adenosine diphosphate (ADP) or 25 μ M ALLN (N-Acetyl-L-leucyl-L-leucyl-L-norleucinal) for 4 h in serum-free DMEM media. (**C**) Conditioned media was collected and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the media is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs Control. (**D**) ApoB100 concentration in the media was determined by Western blot and histograms represent band densitometry analysis of apoB100, normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs Control. (**D**) ApoB100 concentration in the media was determined by Western blot and histograms represent band densitometry analysis of apoB100, normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs Control. (**D**) ApoB100 concentration and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs Control. (**D**) ApoB100 concentration in the media was determined by Western blot and histograms represent band densitometry analysis of apoB100, normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs Control. (**D**) ApoB100 concentration and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs Control.

Marcel (University of Ottawa Heart Institute). Affinity purified peroxidase linked goat anti-mouse and anti-rabbit antibodies were purchased from GE Healthcare (UK). All Stars Negative control small interference RNA (siRNA) were purchased from Qiagen (Mississauga, ON) and human P2Y₁₃ siRNA were purchased from Thermo Scientific Dharmacon (Lafayette, CO). Human P2Y₁₃ plasmid was purchased from Origene (Rockville, MD). Inhibitors were of analytical grade and were solubilized in dimethyl sulfoxide (DMSO).

Cells and Cell Culture

Human hepatocarcinoma, HepG2, cells were regularly maintained in Dulbecco's modified Eagle medium (DMEM) (5 g/L glucose) containing 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin. Passages 4–10 were used and cells that were 80% confluent were treated with DLPC, nucleotides and/or inhibitors for the indicated times and concentrations under serumfree conditions. Cell viability was evaluated after all treatment conditions.

Preparation of DLPC Micelles

DLPC micelles were prepared in DMSO by sonication as previously described [20]. Purity of all phospholipids was >99%.

Knockdown of Human P2Y₁₃ by Small Interference RNA

HepG2 cells were transiently transfected with All Stars Negative control siRNA from Qiagen (Mississauga, ON) or two different siRNA sequences (ACCUUCAUCAUCUACCU- $P2Y_{13}$ CAAAUU or GACACUCAUGCUUCCUUUCAAUU) from Thermo Scientific Dharmacon (Lafayette, CO), by reverse transfection using Lipofectamine 2000 (Invitrogen, Carlsbad, CA) in 12-well plates. In brief, complexes were prepared per manufacturer's specifications with a Lipofectamine 2000-to-siRNA volume-to-mole ratio of 2:40 (µL:pmol) in 200 µL of Opti-MEM I Reduced Serum Media (Invitrogen, Carlsbad, CA). Lipofectamine-siRNA complexes were added to the cells immediately after the cells were seeded at a density of 500,000 cells/well in a volume of 1 mL of normal growth media containing 10% FBS in the absence of penicillin/streptomycin. The cells were treated with



Figure 3. ADP stimulates autophagy and increases cellular LC3-II. (A) HepG2 cells were treated with 100 µM adenosine diphosphate (ADP) for 4 h in serum-free DMEM media. Cell lysates were immunoblotted for LC3. Histograms represent band densitometry analysis of LC3-I and LC3-II,

PLoS ONE | www.plosone.org

normalized to β-actin and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs Control, **P<0.001 vs Control. (**B**) HepG2 cells were serum-starved (Control) or pretreated for 30 min. with 100 µM adenosine diphosphate (ADP) in serum-free DMEM media and then lysates were harvested at the indicated timepoints (0, 3, 6 & 9 h). (**Left panels**) Cell lysates were immunoblotted for LC3 and β-actin. (**Right panels**) Cell lysates were immunoblotted for p62 and β-actin. Blots are representative of 2 independent experiments. (**C**) HepG2 cells were serum-starved (Control) or treated with 100 µM ADP in serum-free DMEM media for 4 h. Cells were fixed and permeabilized and then LC3 and apoA-I were detected by indirect immunofluorescence using primary antibodies against human LC3 and apoA-I, and Alexa Fluor-conjugated secondary antibodies (Alexa Fluor 488 goat anti-rabbit Ab (green for LC3) and Alexa Fluor 647 anti-mouse Ab (red for apoA-I)) by confocal microscopy. Micrograph 100× and 40× images of representative cells from 2 independent experiments done in quadruplicate are shown. doi:10.1371/journal.pone.0036916.q003

ADP, TNF α or DLPC in serum-free DMEM 48 h after transfection. Cell media and lysate samples were harvested at the indicated timepoints for both immunoblot and ELISA analysis. Transfection of the control and test siRNA caused no cytotoxic effects.

Overexpression of Human P2Y₁₃ by Plasmid

The pCMV6 vector containing the full-length human $P2Y_{13}$ cDNA was purchased from Origene (Rockville, MD). HepG2 cells were transiently transfected with control plasmid or the pCMV6- $P2Y_{13}$ plasmid by reverse transfection using FuGENE HD (Roche Applied Science, Laval, QC). Complexes were prepared per manufacturer's instructions with a FuGENE HD-to-DNA volume-to-mass ratio of 6:2 (µl to µg) in 100 µL of Opti-MEM I Reduced



Figure 4. Increasing P2Y₁₃ expression stimulates autophagy and blocks apoA-I secretion. HepG2 cells were transfected with either a control pCMV plasmid (pCMV) or a pCMV plasmid expressing human P2Y₁₃ (pCMV-P2Y13). Cell lysates were collected 48 h after transfection and immunoblotted for P2Y₁₃ to measure protein overexpression (**Upper left panel**). Histograms represent band densitometry analysis of P2Y₁₃, normalized to β -actin (**inset A**) and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs pCMV. (**A**) Cell lysates were immunoblotted for LC3 and histograms represent band densitometry analysis of LC3-II, normalized to β -actin and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs pCMV. (**A**) Cell lysates were immunoblotted for LC3 and histograms represent band densitometry analysis of LC3-II, normalized to β -actin and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs pCMV. (**B**) Transfected cells were treated with 100 μ M ADP in serum-free DMEM media for 4 h, conditioned media was collected and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the media is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs pCMV. (**B**) Transfected and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the media is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs pCMV Control. (**C**) Transfected cells were treated with 12 μ M DLPC in serum-free DMEM media for 24 h, conditioned media was collected and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the media is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.01 vs pCMV-Control, **P<0.01 vs pCMV+DLPC. doi:10.1371/journal.pone.0036916.g004



Figure 5. Reducing P2Y₁₃ expression blocks autophagy and stimulates apoA-I secretion. HepG2 cells were transfected with either a negative control (si-ctrl) or a siRNA against human P2Y₁₃. Cell lysates were collected 48 h after transfection and immunoblotted for P2Y₁₃ to confirm protein knockdown (**Upper left panel**). (**A**) Cell lysates were immunoblotted for LC3 and histograms represent band densitometry analysis of LC3-II, normalized to β -actin, and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs si-Ctrl. (**Inset A**) Histograms represent band densitometry analysis of P2Y₁₃ normalized to β -actin and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs si-Ctrl. (**Inset A**) Histograms represent band densitometry analysis of P2Y₁₃ normalized to β -actin and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs si-Ctrl. (**B**) Transfected cells were treated with 100 µM ADP in serum-free DMEM media for 4 h, conditioned media was collected and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the media is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.05 vs si-Ctrl, **P<0.05 vs si-Ctrl. (**T** ransfected cells were treated with 12 µM DLPC in serum-free DMEM media for 24 h, conditioned media was collected and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the media is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments. *P<0.01 vs si-Ctrl, **P<0.01 vs si-Ctrl, **P<0.03 vs si-Ctrl, **P<0.03 vs si-Ctrl, **P<0.03 vs si-Ctrl, **P<0.04 vs si-Ctrl, **P<0.04 vs si-Ctrl, **P<0.05 vs si-Ctrl, **P<0.05 vs si-Ctrl, **P<0.05 vs si-Ctrl, **P<0.01 vs si-Ctrl, **P<0.01 vs si-Ctrl+DLPC.

Serum Media (Invitrogen, Carlsbad, CA). HepG2 cells were trypsinized and seeded in 12-well plates at a density of 500,000 cells/well in a volume of 1 mL in normal growth media containing 10% FBS in the absence of penicillin/streptomycin and then 50 μ L of the transfection complexes were immediately added to the suspended cells. The cells were treated with ADP, TNF α or DLPC in serum-free DMEM 48 h after transfection. Cell media and lysate samples were harvested at the indicated timepoints for both immunoblot and ELISA analysis. Transfection of the control and test plasmid caused no cytotoxic effects.

ApoA-I ELISA

ApoA-I concentration in conditioned media and cell lysate samples were analyzed by ELISA according to manufacturer's instructions as previously described [20]. 96-well plates were coated overnight with a mouse anti-human apoA-I monoclonal antibody (Meridian Life Sciences, Inc, Saco, ME). Wells were blocked with BSA and then samples/standards were incubated in the wells for 2 h, followed by a 1 h incubation with a horseradish peroxidase-linked goat anti-human apoA-I antibody. K-blue Max TMB substrate (Neogen, Inc) was added to each well, the reaction was stopped with 0.2 N HCl, and the absorbance was recorded at 450 nm. ApoA-I concentration in the conditioned media and cell lysate samples were normalized to total cell protein.

Immunoblot Analysis

After treatment for the indicated timepoints, cells were washed twice with ice-cold PBS. Cells were lysed in NP-40 lysis buffer (Biosource, Camarillo, CA) supplemented with 1 mM PMSF and 1× protease inhibitor cocktail (Sigma, Saint Louis, MO). Cell protein concentrations were determined by the BCA Protein Assay (Thermo Fisher Scientific, Waltham, MA). Cell lysate samples containing equal total protein (30 μ g) were separated by SDS-PAGE and analyzed by Western blot using specific antibodies to apoA-I, apoB100, P2Y₁₃, LC3, p62, p-ERK1/2, p-Akt, p-mTOR, p-IR- β , and β -actin. Blots were exposed using the Alpha Innotech FluorChemTM HD Imager and band intensities were quantified with the Alpha Ease FCTM software.



Figure 6. Extracellular nucleotides and P2Y₁₃ expression regulate ERK1/2 signaling. (**A**) HepG2 cells were pre-treated with 12 μ M DLPC for 30 min. and then incubated with and without ADP (100 μ M) for 0, 5, 15 and 30 min in DMEM serum-free media. Cell lysates were immunoblotted for phosphorylated ERK1/2. Histograms represent densitometry analysis of p-ERK1/2 normalized to β -actin and expressed as mean percent change \pm SD of 3 independent experiments. *P<0.01 vs Ctrl, **P<0.05 vs ADP 5 min. (**B**) HepG2 cells were transfected with either negative control (si-Ctrl) or P2Y₁₃ siRNA (si-P2Y₁₃) and incubated for 24 h. Cell lysates were immunoblotted for phosphorylated and total ERK1/2. Histograms represent band densitometry analysis of the ratio of phospho-ERK1/2 (p-ERK1/2) to total ERK1/2 (t-ERK1/2) and are expressed as mean \pm SD for 3 independent experiments. *P<0.05 vs. control siRNA. doi:10.1371/journal.pone.0036916.q006

Immunofluorescence and Colocalization

After treatment, the cells were fixed with 4% paraformaldehyde, permeabilized with 0.1% Triton X-100, and then blocked in 10% FBS for 30 min. The cells were then immunostained with 1:200 rabbit polyclonal anti-LC3 (MBL International, Woburn, MA) and 1:200 mouse anti-apoA-I antibodies (Meridian Life Sciences, Saco, ME) for 1 h and then with 1:1000 Alexa Fluor 488 goat antirabbit IgG (H+L) and 1:1000 Alexa Fluor 647 donkey anti-mouse IgG (H+L) (Invitrogen, Burlington, ON) for an additional 1 h. Images were acquired on an Olympus 1X80FV1000 confocal laser microscope using Olympus Fluoview FV1000 software and colocalization was quantified using FV10-ASW V2.1.

Statistical Analysis

Values are shown as Mean \pm SD for at least 3 independent experiments and P < 0.05 was considered significant. Differences between mean values were evaluated by one-way analysis of variance (ANOVA) on ranks by a pairwise multiple comparison using the Student-Newman-Keuls post-hoc test (SigmaStat; Systat Software, Inc., San Jose, CA).

Results

Extracellular nucleotides inhibit hepatic apoA-I secretion

Previous work has shown that membrane ATPases control the nucleotide-dependent endocytosis of HDL [14,31] and that inhibition of ADP production by membrane ATPases may promote HDL secretion [18,19]. To determine if ADP directly affects hepatic HDL secretion, experiments were undertaken to evaluate the effect of ADP on apoA-I secretion from liver cells. As

we have previously shown [19], dilinoleoylphosphatidylcholine (DLPC) (12 μ M) stimulates a ~4-fold increase in apoA-I secretion (accumulation in the media) from HepG2 cells over 24 h (Figure 1A). Pre-treatment of the cells with 20-100 µM ADP completely blocked the induction of apoA-I secretion by DLPC (Figure 1A). Treatment with ATP (100 μ M) had a lesser effect than ADP and only blocked the induction of apoA-I secretion by 60% (Figure S1). The effects of DLPC and ADP on apoA-I secretion were also evident after short time periods. ADP $(100 \ \mu M)$ was able to significantly reduce basal apoA-I secretion at 4 h (Figures 1B and 2C). DLPC stimulated the secretion of apoA-I after 4 h and ADP completely blocked this DLPCinduction of apoA-I secretion. Lower doses of ADP (25 and 50μ M) also inhibited apoA-I secretion at 4 h (not shown). ADP treatment had no effect on cellular apoA-I levels after 4 h (Figure 1C) or 24 h (not shown). ADP therefore significantly reduced apoA-I mass (media+lysate) at 4 h by 20% (ADP alone) to 30% (ADP+DLPC).

Adenosine diphosphate stimulates hepatic apoB100 and apoE secretion

Experiments were undertaken to determine if ADP might affect the secretion of other apolipoproteins. In contrast to that observed with apoA-I, ADP significantly increased apoB100 secretion (**Figure 2A**) and apoE secretion (**Figure 2B**) from HepG2 cells after 24 h. ApoB100 secretion is regulated by proteasomal degradative pathways [22,23,32] and therefore the effect of ADP on apolipoprotein secretion was compared to that of proteasomal inhibitors. ADP and ALLN (25 μ M) decreased apoA-I levels in the media to a similar extent at 4 h (**Figure 2C**), but increased the



Figure 7. Extracellular nucleotides and P2Y₁₃ expression regulate insulin receptor signaling. (A) HepG2 cells were pre-treated with 12 μ M DLPC for 30 min. and then incubated with and without ADP (100 μ M) for 0, 5, 15 and 30 min in DMEM serum-free media. Cell lysates were immunoblotted for phosphorylated Akt (Ser473). Histograms represent densitometry analysis of p-Akt normalized to β -actin and expressed as mean percent change \pm SD of 3 independent experiments. *P<0.01 vs Control, **P<0.001 vs DLPC 5 min. (B) HepG2 cells were pre-treated with adenosine diphosphate (ADP) (100 μ M) or TNF α (10 ng/ml) for 5 min. and then with human insulin (100 nM) for 5 min in DMEM serum-free media. Cell lysates were immunoblotted for phosphorylated Akt (Ser473). Histograms represent densitometry analysis of p-Akt normalized to β -actin and expressed as mean \pm SD for 3 independent experiments. *P<0.001 vs Chrl, **P<0.001 vs insulin alone. (**C&D**) HepG2 cells were transfected with either negative control (si-Ctrl) or P2Y₁₃ siRNA (si-P2Y₁₃) and incubated for 48 h. Cells were then treated with adenosine diphosphate (ADP) (100 μ M) or TNF α (10 ng/ml) for 5 min. in DMEM serum-free media. (C) Cell lysates were immunoblotted for phosphorylated for 48 h. Cells were then treated with adenosine diphosphate (ADP) (100 μ M) or TNF α (10 ng/ml) for 5 min. in DMEM serum-free media. (C) Cell lysates were immunoblotted for phosphorylated insulin receptor (IR- β) (Tyr1345). Histograms represent densitometry analysis of p-IR- β normalized to β -actin and expressed as mean \pm SD for 3 independent experiments. *P<0.01 vs si-Ctrl, **P<0.01 vs for 3 independent experiments. *P<0.01 vs clrl, **P<0.01 vs independent experiments. *P<0.01 vs si-Ctrl, **P<0.01 vs independent experiments. *P<0.01 vs is-Ct

P<0.001 vs si-Ctrl. (**D**) Cell lysates were also immunoblotted for phosphorylated Akt (Ser473). Histograms represent densitometry analysis of p-Akt normalized to β -actin and expressed as mean \pm SD for 3 independent experiments.*P<0.01 vs si-Ctrl, **P<0.05 vs. si-Ctrl, ***P<0.001 vs si-Ctrl. doi:10.1371/journal.pone.0036916.g007

concentration of apoB100 in the media (**Figure 2D**). The work suggests that ADP may regulate apolipoprotein secretion in a similar fashion to that observed with proteasomal inhibitors.

Adenosine diphosphate stimulates autophagy

Studies have shown that proteasomal inhibitors also act to stimulate autophagy [24,25] and therefore experiments were undertaken to determine if ADP affects autophagy and impacts the level of the autophagy marker, LC3. HepG2 cells were treated with 100 µM ADP for 4 h and then cell lysates were probed for LC3. Figure 3A shows that ADP significantly increased both LC3-I and LC3-II levels at 4 h. The data suggests that ADP stimulates autophagy and this view is confirmed by the data in Figure 3B. To determine if ADP uniquely affects the activation versus flux of LC3 in HepG2 cells, 9 h time course studies were undertaken. Figure 3B shows that serum starvation (control) increases LC3-II levels in HepG2 cell lysates at 3 h, which return to below basal levels by 9 h (left panel blots). ADP increases LC3-II levels more quickly than the control starvation (within the 30 min pretreatment) and maintains increased cellular LC3-II levels for 6 h (Figure 3B, left panel blots). Similar results were seen with another autophagy marker protein, p62 (Figure 3B, right panel blots).

Immunofluorescent experiments using laser confocal microscopy confirm the Western blot data and further showed that apoA-I and LC3 are colocalized within autophagosomes. **Figure 3C** (**and Figure S2**) illustrates 4 h confocal micrographic images of indirect immunofluorescent stained apoA-I and LC3 in fixed and permeabilized HepG2 cells. Control starvation conditions show the staining pattern for both LC3 (green) and apoA-I (red) and illustrate low levels of colocalization of the two proteins in the merged images (yellow-orange). Similar to that shown in **Figures 3A&B**, ADP increases cellular LC3 levels relative to control, and confocal images show LC3 to be localized to punctate autophagosomal structures within the cell (**Figure 3C**). When LC3 and apoA-I immunofluorescent images for ADP treated cells



Figure 8. Extracellular nucleotides act through P2Y₁₃ to stimulate autophagy. Elevations in blood glucose promote the secretion and accumulation of nucleotides in the circulation. Nucleotides act through P2Y₁₃ to activate mitogenic pathways, inhibit insulin receptor signaling and stimulate autophagic protein degradation. Enhanced purinergic signaling in insulin resistance may give rise to a chronic induction of cellular autophagy and a reduction in apoA-I secretion from the liver.

doi:10.1371/journal.pone.0036916.g008

are merged, the images show significantly (P<0.001) more yelloworange structures in ADP-treated vs control cells (23.8% \pm 5.2% vs 7.5% \pm 3.0%), which indicates that ADP significantly increases the colocalization of apoA-I with LC3. ADP-dependent increases in LC3-II levels were completely inhibited by treatment with 3methyladenine (3-MA) (5 mM) or U0126 (10 µM), while LC3-II levels were further increased by treatment with chloroquine (50 µM) or Wortmannin (10 µM) (**Figure S3**).

P2Y₁₃ expression affects autophagy and apoA-I secretion

Extracellular ADP affects cellular apoA-I metabolism through the GPCR, P2Y₁₃ [15] in hepatocytes and therefore the effect of P2Y₁₃ expression on autophagy and apoA-I secretion were evaluated. Transfecting HepG2 cells with a pCMV6-P2Y₁₃ plasmid promoted a ~50% increase in P2Y₁₃ expression (**Figure 4, Western blot and inset**) and a parallel increase in LC3-II levels (**Figure 4A**). Treatment of liver cells with a pCMV6-P2Y₁₃ plasmid blocked the basal secretion of apoA-I at 4 h, similar to that observed after treatment with ADP, while a combination of pCMV6-P2Y₁₃ and ADP had no additional effect on apoA-I secretion (**Figure 4B**). Increasing P2Y₁₃ expression in the liver cells significantly reduced the DLPC-induction of apoA-I secretion after 24 h (**Figure 4C**).

Reducing P2Y₁₃ expression had the opposite effect. Transfecting HepG2 cells with P2Y₁₃ siRNA promoted a >50% reduction in P2Y₁₃ protein expression (**Figure 5, Western blot and inset**) and caused a similar reduction in LC3-II levels (**Figure 5A**). P2Y₁₃ siRNA significantly stimulated apoA-I secretion at both 4 h and 24 h (**Figures 5B and 5C**). After a 4 h incubation, ADP was unable to block apoA-I secretion in cells treated with P2Y₁₃ siRNA, but conversely, increased apoA-I secretion relative to P2Y₁₃ siRNA treatment alone (**Figure 5B**). Treatment with DLPC (12 μ M) or P2Y₁₃ siRNA for 24 h stimulated apoA-I secretion by ~2.5-fold, while treatment with both DLPC and P2Y₁₃ siRNA promoted a ~10-fold stimulation in apoA-I secretion from HepG2 cells (**Figure 5C**). P2Y₁₃ siRNA completely blocked the effect of ADP on LC3-II (**Figure S4**).

Nucleotides and P2Y₁₃ regulate MAPK signaling

Cellular autophagy is regulated by the activation of MAPK (ERK1/2) and therefore the importance of ERK1/2 in ADPdependent purinergic signaling was investigated. The nucleotide, ADP, stimulates ERK1/2 phosphorylation over a 30 min period and DLPC blocked the activation of ERK1/2 by ADP (**Figure 6A**). P2Y₁₃ expression also affects ERK1/2 phosphorylation and a 50% reduction in cellular P2Y₁₃ expression caused a significant reduction in ERK1/2 phosphorylation after 24 h (**Figure 6B**). Cell viability was unaffected by the P2Y₁₃ siRNA.

Nucleotides and P2Y₁₃ regulate insulin receptor signaling

Cellular autophagy is activated by the inhibition of protein kinase B (Akt) [27]. DLPC stimulates the phosphorylation of Akt at 5 min. and ADP completely blocks the activation of Akt by DLPC (**Figure 7A**). ADP also inhibits the activation of Akt by insulin. **Figure 7B** shows that insulin stimulates Akt phosphorylation in HepG2 cells and that both tumor necrosis factor α (TNF α) and ADP inhibit insulin-induced Akt phosphorylation by ~50%. Reducing P2Y₁₃ expression by treatment with P2Y₁₃ siRNA appeared to inhibit cellular autophagic pathways by stimulating

the phosphorylation of Akt. A reduction in cellular P2Y₁₃ expression significantly increases the phosphorylation of IR- β and Akt (Figure 7C&D) by >3-fold. P2Y₁₃ gene silencing significantly augmented the insulin-induced phosphorylation of Akt (**Figure S5**) and blocked the inhibitory effect of $TNF\alpha$ and ADP on IR-β and Akt phosphorylation (Figure 7C&D and S5). Similar results were observed for the insulin-like growth factor receptor (IGF-1R) (not shown). P2Y₁₃ overexpression had the opposite effect. Transfection of HepG2 cells with pCMV6-P2Y₁₃ plasmid significantly reduced the phosphorylation of IR- β (**Figure S6**). Increasing $P2Y_{13}$ expression significantly inhibited the insulininduced phosphorylation of IR- β and Akt (**Figure S6**), similar to that observed after treatment of HepG2 cells with ADP (Figure 6C). P2Y₁₃ expression significantly affected Akt phosphorylation, but had no effect on mTOR (Figure S7). ADP actually increased mTOR phosphorylation, while the mTOR inhibitor, rapamycin, significantly reduced p-mTOR levels (Figure S8).

Discussion

Insulin resistance and hyperglycemia have been shown to perturb plasma lipoprotein metabolism and increase apoB100 levels, but decrease HDL [1,2]. Insulin resistance is consequently a well accepted risk factor for the development of cardiovascular disease [1]. High blood glucose levels stimulate ATP production and promote the release of nucleotides from circulating blood cells, endothelial cells and smooth muscle cells [3,4]. ATP is unstable in the circulation and is quickly converted to ADP [7,8]. Elevations in blood nucleotide levels can impact cardiovascular disease [4,5,13] and inhibition of ADP-dependent thrombosis with P2Y₁₂ receptor inhibitors has already shown significant cardiovascular therapeutic value [33,34]. Niacin has also been shown to have cardiovascular therapeutic value and this molecule appears to act through another G-protein coupled receptor [16] to block purinergic signaling [18] and atherogenesis [35].

Our studies show that HDL secretion is regulated similarly in primary human hepatocytes and HepG2 liver cells and that the linoleic acid phospholipid, dilinoleoylphosphatidylcholine (DLPC), can stimulate HDL/apoA-I secretion [19,20,36]. DLPC appears to act much like niacin to prevent purinergic signaling by inhibiting F1-ATPase and blocking the production of ADP [19]. This view has been confirmed by the present work, which shows that ADP is a potent antagonist to the induction of apoA-I secretion by DLPC. Figure 1 shows that an $[ADP] > 10 \,\mu M$ completely blocked the induction of apoA-I secretion by DLPC at 24 h and also at 4 h. Conversely, ADP can directly stimulate apoB100 and apoE secretion from human liver cells at 24 h (Figure 2A&B). The normal physiological concentration of ADP in the bloodstream has been reported to be $\sim 15 \,\mu M$ [8], but elevated blood glucose levels can increase nucleotide secretion and accumulation in the circulation [3,4]. High circulatory nucleotide levels would therefore be expected to block hepatic apoA-I secretion and stimulate apoB100 output. This is indeed similar to what is thought to occur in hyperglycemic, insulin resistant patients [1,37,38]. This may suggest that elevations in blood nucleotide levels may be partly causative to abnormal plasma lipoprotein levels.

Since apoB100 secretion is known to be regulated by proteasomal degradation, we evaluated the effect of proteasomal inhibitors on both apoB100 and apoA-I secretion. ADP and the proteasomal inhibitor, ALLN, appear very similar and both inhibit apoA-I secretion, but stimulate apoB100 secretion at 4 h (**Figure 2C&D**). ADP may therefore act similar to proteasomal

inhibitors to stimulate autophagy [24,25]. ADP stimulates autophagy in HepG2 cells and increases the level of autophagic markers, LC3-I and LC3-II, and p62 over a 6 h period (Figure 3). The ability of ADP to stimulate apoB100 secretion (Figure 2) and increase p62 levels may indicate that ADP blocks proteasomal degradation, since p62 levels are known to rise when the proteasome is inhibited [26,27]. Confocal studies confirmed the higher levels of LC3 after treatment with ADP. Micrographs showed that LC3-II was located in punctate autophagosomes within the liver cells and clearly showed that higher levels of LC3 and apoA-I were colocalized within autophagosomes in ADP treated cells (Figure 3C). The view that a stimulation in autophagy may inhibit apoA-I secretion is consistent with earlier work, which has shown that serum deprivation, a treatment wellknown to directly stimulate autophagy (Figure 3B and [26,27]), also inhibits apoA-I secretion of HepG2 cells by $\sim 50\%$ over the first hour [39].

Human liver cells contain two ADP-receptors, P2Y1 and P2Y13, but HDL metabolism is primarily affected by $P2Y_{13}$ [15,40,41]. ADP is a potent agonist to $P2Y_{13}$ and stimulates a rapid (10 min) endocytic recycling pathway for extracellular apoA-I [14,15,31]. This recycling pathway has been shown contribute to apoA-I lipidation, cholesterol efflux and apoE resecretion, but does not promote significant apoA-I degradation [42–44]. Conversely, it is known that apoA-I secretion is affected by cellular degradation [18,21] and activation of P2Y13 appears to stimulate degradation pathways. The direct effect of $P2Y_{13}$ on both autophagy and apoA-I secretion from human liver cells is clearly illustrated by increasing or silencing P2Y13 gene expression. P2Y13 overexpression significantly increased cellular LC3-II levels and decreased apoA-I secretion (Figure 4), similar to that seen with exogenous ADP. Conversely, P2Y₁₃ gene silencing with siRNA significantly decreased LC3-II levels, increased both basal apoA-I secretion and the DLPC induction in apoA-I secretion, and blocked the effects of ADP on autophagy and apoA-I secretion (Figure S4 and **Figure 5**). $P2Y_{13}$ expression in HepG2 cells therefore appears to regulate apoA-I secretion through cellular autophagic pathways.

In contrast to this work, studies in $P2Y_{13}$ -deficient mice have shown that reducing P2Y₁₃ expression caused a small reduction in plasma HDL levels in vivo [45,46]. This may be partly due to the fact that mice are not a human equivalent model for the study of HDL metabolism, since numerous liver-specific signaling pathways differ in mice [47]. Nucleotide signaling also differs significantly in rodents [40,48]. In studies with rat hepatocytes, ADP acts through $P2Y_2$ receptors to increase [IP3] and [Ca²⁺] [48]. This does not occur in human cells. Treatment of both primary human liver cells and HepG2 cells with ADP or UDP has no effect on cellular [IP3] and $[Ca^{2+}]$ [40]. In human liver cells, ADP stimulates MAPK and reduces [cAMP] [15,41], both of which are known to reflect activation of $P2Y_{13}$ [49,50]. We show that ADP stimulates ERK1/2 in human liver cells (Figure 6A), but ADP has the opposite effect in murine pancreatic cells and inhibits ERK1/2 phosphorylation [51]. ADP and P2Y₁₃-dependent signaling may therefore be very different in humans and rodents and this may explain why P2Y13-deficient mice show no major lipoprotein phenotype [45,46].

Nucleotide signaling through P2Y receptors is well known to activate MAPK pathways [49,50] and MAPK is a well-known activator of cellular autophagy [26]. DLPC may therefore stimulate apoA-I secretion by blocking MAPK activation and preventing the autophagic degradation of apoA-I. DLPC can block an ADP-dependent activation of ERK1/2 in HepG2 cells (**Figure 6A**), similar to that shown in neuronal cells, where DLPC blocked ERK1/2 activation by TNFα and hydrogen peroxide

10

[52]. Treatment of liver cells with the MEK1/2 inhibitor, U0126, completely blocked the ADP-dependent increase in LC3-II levels (**Figure S3**) and $P2Y_{13}$ -gene silencing also muted ERK1/2 activation (Figure 6B). Our data suggests that MAPK is not alone in regulating autophagy and apoA-I secretion. ADP can block the activation of Akt by DLPC and insulin (Figure 7A&B) and Akt is an established inhibitor of autophagy [26,30]. Therefore, while nucleotides may affect cellular autophagic pathways through MAPK pathways, the insulin signaling-dependent Akt pathways may also play an important role in the purinergic regulation of autophagy and HDL secretion. Akt can inhibit autophagy directly, and indirectly through mTOR [26]. Our work, however suggests that ADP stimulates autophagy through mTOR-independent pathways, since P2Y13 expression had no effect on mTOR phosphorylation, while ADP increased p-mTOR levels (Figures S7&S8).

If ADP-dependent signaling through P2Y₁₃ impacts Akt-dependent pathways, it follows that P2Y₁₃ expression would be expected to affect insulin receptor signaling. Consistent with this view, a reduction in P2Y₁₃ expression directly stimulates the phosphorylation of IR- β and Akt by >3-fold (**Figure 7**), increases insulin-dependent signaling (**Figure S5**) and completely blocks the inhibition of insulin receptor signaling by TNF α and ADP (**Figure 7**). Increasing P2Y₁₃ expression has the opposite effect and significantly inhibits insulin-induced phosphorylation of IR- β and Akt (**Figure S6**). Nucleotide signaling through P2Y₁₃ may therefore affect insulin receptor signaling. This work appears consistent with other studies showing that ADP acts through P2Y₁₃ to inhibit insulin secretion from pancreatic beta cells [51,53].

This work shows that lipoprotein secretion and insulin signaling pathways are affected by hepatic membrane purinergic receptor signaling. Elevations in blood glucose promote the synthesis and secretion of nucleotides from circulating blood cells and vascular tissues. Nucleotides that accumulate in the circulation can then act through purinergic receptors, i.e. P2Y₁₃, to stimulate mitogenic pathways and inhibit insulin receptor signaling. Enhanced purinergic signaling in insulin resistance may give rise to an inhibition of proteasomal degradation and a chronic induction of cellular autophagy (**Figure 8**). The net result is a stimulation of apoB100 secretion from the liver and a reduction in apoA-I secretion. This may partly explain the well-described lipoprotein phenotype associated with insulin resistance [1].

Supporting Information

Figure S1 ATP decrease apoA-I levels in the media at 24 h. (A) HepG2 cells were pre-treated with adenosine diphosphate (ADP) or adenosine triphosphate (ATP) (100 μ M) for 30 min. and then incubated with 12 μ M DLPC in serum-free DMEM media. Conditioned media was collected after 24 h treatment and apoA-I concentration was quantified by ELISA. ApoA-I concentration in the media is normalized to total cell protein and expressed as mean \pm SD of 3 independent experiments *P<0.01 vs DLPC, **P<0.001 vs DLPC. (TIF)

Figure S2 ADP stimulates autophagy. HepG2 cells were serum-starved (Control) (A) or treated with 100 μ M ADP (B) in serum-free DMEM media for 4 h. Cells were fixed and permeabilized and then apoA-I and LC3 were detected by indirect immunofluorescence using confocal microscopy. Original images of representative micrographs at 100 × magnification from 2 independent experiments performed in quadruplicate are shown.

(TIF)

Figure S3 Effect of autophagy inhibitors on cellular LC3-II. HepG2 cells were pre-treated with 50 μ M chloroquine, 5 mM 3-methyladenine (3-MA), 10 μ M wortmannin or 10 μ M U1026 for 30 min±100 μ M ADP for 4 h in serum-free DMEM media. Cell lysates were immunoblotted for LC3. Histograms represent band densitometry analysis of LC3-II normalized to β -actin and expressed as percent change ± SD of 3 independent experiments. *P<0.05 vs Control and **P<0.01 vs Control. (TIF)

Figure S4 P2Y₁₃ knockdown inhibits the ADP-dependent stimulation in autophagy. HepG2 cells were transfected with either negative control (si-Ctrl) or P2Y₁₃ siRNA (si-P2Y13) and incubated for 48 h. Cells were then incubated with ADP (100 μ M) for 4 h in DMEM serum-free media. Cell lysates were immunoblotted for LC3 and blots are representative of 3 independent experiments.

(TIF)

Figure S5 Reducing P2Y₁₃ expression augments insulin receptor signaling. HepG2 cells were transfected with either negative control (si-Ctrl) or P2Y₁₃ siRNA (si-P2Y13) and incubated for 48 h. Cells were then pre-incubated with ADP (100 µM) for 5 min. and then with human insulin (100 nM) for 5 min in DMEM serum-free media. Cell lysates were immunoblotted for phosphorylated Akt (Ser473). Histograms represent densitometry analysis of p-Akt normalized to β-actin and expressed as mean percent change ± SD for 2 independent experiments.*P<0.05 vs si-Ctrl, **P<0.01 vs si- P2Y13. (TIF)

Figure S6 P2Y₁₃ overexpression blocks insulin receptor signaling. HepG2 cells were transfected with either a control pCMV plasmid (pCMV) or a pCMV plasmid expressing human P2Y₁₃ (pCMV-P2Y13). Cell lysates were collected 48 h after transfection and immunoblotted for P2Y₁₃ to measure protein overexpression (**inset, panel A**). Cells were then treated with human insulin (100 nM) for 5 min in DMEM serum-free media. Cell lysates were immunoblotted for insulin receptor (p-IR- β) (**A**) and phosphorylated Akt (Ser473) (**B**). Histograms represent densitometry analysis normalized to β -actin and are expressed as mean percent change \pm SD for 3 independent experiments. (**A**)*P<0.001 vs pCMV and **P<0.01 vs pCMV+Insulin. (**B**) *P<0.001 vs pCMV and **P<0.05 vs pCMV+Insulin. (TIF)

Figure S7 Reducing P2Y₁₃ expression had no effect on the phosphorylation of mTOR. HepG2 cells were transfected with either a negative control (si-ctrl) or a siRNA against human P2Y₁₃. Cell lysates were collected 48 h after transfection and immunoblotted for phosphorylated Akt (Ser473) and phosphorylated mTOR (Ser2448). Histograms represent densitometry analysis of p-Akt and p-mTOR normalized to β -actin and expressed as mean \pm SD of 3 independent experiments. *P<0.001 vs si-Ctrl.

(TIF)

Figure S8 ADP increases mTOR phosphorylation. HepG2 cells were treated with 100 μ M ADP or 250 nM rapamycin for 4 h. Cell lysates were immunoblotted for phosphorylated mTOR (Ser2448). Histograms represent densitometry analysis of p-mTOR normalized to β -actin and expressed as mean percent change \pm SD of 3 independent experiments. *P<0.05 vs control.



Acknowledgments

We thank Dr. Nihar Pandey for his advice and technical assistance in the insulin signaling experiments.

References

- Grundy SM (1998) Hypertriglyceridemia, atherogenic dyslipidemia, and the metabolic syndrome. Am J Cardiol 81: 18B–25B.
- Adiels M, Olofsson SO, Taskinen MR, Boren J (2008) Overproduction of very low-density lipoproteins is the hallmark of the dyslipidemia in the metabolic syndrome. Arterioscler Thromb Vasc Biol 28: 1225–1236.
- Solini A, Iacobini C, Ricci C, Chiozzi P, Amadio L, et al. (2005) Purinergic modulation of mesangial extracellular matrix production: role in diabetic and other glomerular diseases. Kidney Int 67: 875–885.
- Nilsson J, Nilsson LM, Chen YW, Molkentin JD, Erlinge D, et al. (2006) High glucose activates nuclear factor of activated T cells in native vascular smooth muscle. Arterioscler Thromb Vasc Biol 26: 794–800.
- Di VF, Solini A (2002) P2 receptors: new potential players in atherosclerosis. Br J Pharmacol 135: 831–842.
- Di VF, Boeynaems JM, Robson SC (2009) Extracellular nucleotides as negative modulators of immunity. Curr Opin Pharmacol 9: 507–513.
- Brown PR, Parks RE Jr., Herod J (1973) Use of high-pressure liquid chromatography for monitoring nucleotide concentration in human blood: a preliminary study with stored blood cell suspensions. Clin Chem 19: 919–922.
- Harkness RA, Coade SB, Webster AD (1984) ATP, ADP and AMP in plasma from peripheral venous blood. Clin Chim Acta 143: 91–98.
- Dwyer KM, Deaglio S, Gao W, Friedman D, Strom TB, et al. (2007) CD39 and control of cellular immune responses. Purinergic Signal 3: 171–180.
- Erlinge D, Burnstock G (2008) P2 receptors in cardiovascular regulation and disease. Purinergic Signal 4: 1–20.
- Khakh BS, North RA (2006) P2X receptors as cell-surface ATP sensors in health and disease. Nature 442: 527–532.
- Trautmann A (2009) Extracellular ATP in the immune system: more than just a "danger signal". Sci Signal 2: pe6.
- Sellers MB, Tricoci P, Harrington RA (2009) A new generation of antiplatelet agents. Curr Opin Cardiol 24: 307–312.
- Martinez LO, Jacquet S, Esteve JP, Rolland C, Cabezon E, et al. (2003) Ectopic beta-chain of ATP synthase is an apolipoprotein A-I receptor in hepatic HDL endocytosis. Nature 421: 75–79.
- Jacquet S, Malaval C, Martinez LO, Sak K, Rolland C, et al. (2005) The nucleotide receptor P2Y13 is a key regulator of hepatic high-density lipoprotein (HDL) endocvtosis. Cell Mol Life Sci 62: 2508–2515.
- Tunaru S, Kero J, Schaub A, Wufka C, Blaukat A, et al. (2003) PUMA-G and HM74 are receptors for nicotinic acid and mediate its anti-lipolytic effect. Nat Med 9: 352–355.
- Li X, Millar JS, Brownell N, Briand F, Rader DJ (2010) Modulation of HDL metabolism by the niacin receptor GPR109A in mouse hepatocytes. Biochem Pharmacol 80: 1450–1457.
- Zhang LH, Kamanna VS, Zhang MC, Kashyap ML (2008) Niacin inhibits surface expression of ATP synthase {beta} chain in HepG2 cells: implications for raising HDL. J Lipid Res 49: 1195–1201.
- Pandey NR, Renwick J, Rabaa S, Misquith A, Kouri L, et al. (2009) An induction in hepatic HDL secretion associated with reduced ATPase expression. Am J Pathol 175: 1777–1787.
- Pandey NR, Renwick J, Misquith A, Sokoll K, Sparks DL (2008) Linoleic Acid-Enriched Phospholipids Act through Peroxisome Proliferator-Activated Receptors alpha To Stimulate Hepatic Apolipoprotein A-I Secretion. Biochemistry 47: 1579–1587.
- Hopewell S, Pandey NR, Misquith A, Twomey E, Sparks DL (2008) Phosphatidylinositol acts through mitogen-activated protein kinase to stimulate hepatic apolipoprotein A-I secretion. Metabolism 57: 1677–1684.
- Fisher EA, Zhou MY, Mitchell DM, Wu XJ, Omura S, et al. (1997) The degradation of apolipoprotein B100 is mediated by the ubiquitin-proteasome pathway and involves heat shock protein 70. J Biol Chem 272: 20427–20434.
- Adeli K, Macri J, Mohammadi A, Kito M, Urade R, et al. (1997) Apolipoprotein B is intracellularly associated with an ER-60 protease homologue in HepG2 cells. J Biol Chem 272: 22489–22494.
- Ding WX, Ni HM, Gao W, Yoshimori T, Stolz DB, et al. (2007) Linking of autophagy to ubiquitin-proteasome system is important for the regulation of endoplasmic reticulum stress and cell viability. Am J Pathol 171: 513–524.
- Zhu K, Dunner K Jr., McConkey DJ (2010) Proteasome inhibitors activate autophagy as a cytoprotective response in human prostate cancer cells. Oncogene 29: 451–462.
- Ravikumar B, Sarkar S, Davies JE, Futter M, Garcia-Arencibia M, et al. (2010) Regulation of mammalian autophagy in physiology and pathophysiology. Physiol Rev 90: 1383–1435.
- Kroemer G, Marino G, Levine B (2010) Autophagy and the integrated stress response. Mol Cell 40: 280–293.

Author Contributions

Conceived and designed the experiments: CC DLS. Performed the experiments: CC. Analyzed the data: CC DLS. Wrote the paper: CC DLS.

- De Meyer GR, Martinet W (2009) Autophagy in the cardiovascular system. Biochim Biophys Acta 1793: 1485–1495.
- Nemchenko A, Chiong M, Turer A, Lavandero S, Hill JA (2011) Autophagy as a therapeutic target in cardiovascular disease. J Mol Cell Cardiol 51: 584–593.
 Yang Z, Klionsky DJ (2010) Eaten alive: a history of macroautophagy. Nat Cell
- Fang Z, Khonsky DJ (2010) Eaten anve, a instory of macroautophagy, ivat Cen Biol 12: 814–822.
 Esbra AC, Vantouvout P, Champagne F, Targe F, Polland C, et al. (2006) Call
- Fabre AC, Vantourout P, Champagne E, Terce F, Rolland C, et al. (2006) Cell surface adenylate kinase activity regulates the F(1)-ATPase/P2Y (13)-mediated HDL endocytosis pathway on human hepatocytes. Cell Mol Life Sci 63: 2829–2837.
- 32. Rutledge AC, Qiu W, Zhang R, Kohen-Avramoglu R, Nemat-Gorgani N, et al. (2009) Mechanisms targeting apolipoprotein B100 to proteasomal degradation: evidence that degradation is initiated by BiP binding at the N terminus and the formation of a p97 complex at the C terminus. Arterioscler Thromb Vasc Biol 29: 579–585.
- Behan MW, Chew DP, Aylward PE (2010) The role of antiplatelet therapy in the secondary prevention of coronary artery disease. Curr Opin Cardiol 25: 321–328.
- 34. Cattaneo M (2010) New P2Y(12) inhibitors. Circulation 121: 171-179.
- Lukasova M, Malaval C, Gille A, Kero J, Offermanns S (2011) Nicotinic acid inhibits progression of atherosclerosis in mice through its receptor GPR109A expressed by immune cells. J Clin Invest 121: 1163–1173.
- Chatterjee C, Young EK, Pussegoda KA, Twomey EE, Pandey NR, et al. (2009) Hepatic high-density lipoprotein secretion regulates the mobilization of cellsurface hepatic lipase. Biochemistry 48: 5994–6001.
- Lewis GF, Rader DJ (2005) New insights into the regulation of HDL metabolism and reverse cholesterol transport. Circ Res 96: 1221–1232.
- Meshkani R, Adeli K (2009) Hepatic insulin resistance, metabolic syndrome and cardiovascular disease. Clin Biochem 42: 1331–1346.
- Ranganathan S, Kottke BA (1990) Rapid regulation of apolipoprotein A-I secretion in HepG2 cells by a factor associated with bovine high-density lipoproteins. Biochim Biophys Acta Lipids Lipid Metab 1046: 223–228.
- Schofl C, Ponczek M, Mader T, Waring M, Benecke H, et al. (1999) Regulation of cytosolic free calcium concentration by extracellular nucleotides in human hepatocytes. Am J Physiol 276: G164–G172.
- Malaval C, Laffargue M, Barbaras R, Rolland C, Peres C, et al. (2009) RhoA/ ROCK I signalling downstream of the P2Y13 ADP-receptor controls HDL endocytosis in human hepatocytes. Cell Signal 21: 120–127.
- Heeren J, Grewal T, Laatsch A, Rottke D, Rinninger F, et al. (2003) Recycling of apoprotein E is associated with cholesterol efflux and high density lipoprotein internalization. J Biol Chem 278: 14370–14378.
- Rohrer L, Cavelier C, Fuchs S, Schluter MA, Volker W, et al. (2006) Binding, internalization and transport of apolipoprotein A-I by vascular endothelial cells. Biochim Biophys Acta 1761: 186–194.
- Denis M, Landry YD, Zha X (2008) ATP-binding cassette A1-mediated lipidation of apolipoprotein A-I occurs at the plasma membrane and not in the endocytic compartments. J Biol Chem 283: 16178–16186.
- Fabre AC, Malaval C, Ben AA, Verdier C, Pons V, et al. (2010) P2Y13 receptor is critical for reverse cholesterol transport. Hepatology 52: 1477–1483.
- Blom D, Yamin TT, Champy MF, Selloum M, Bedu E, et al. (2010) Altered lipoprotein metabolism in P2Y(13) knockout mice. Biochim Biophys Acta 1801: 1349–1360. pp 1349–1360.
- Berthou L, Duverger N, Emmanuel F, Langouët S, Auwerx J, et al. (1996) Opposite regulation of human versus mouse apolipoprotein A-I by fibrates in human apolipoprotein A-I transgenic mice. J Clin Invest 97: 2408–2416.
- Dixon CJ, Hall JF, Boarder MR (2003) ADP stimulation of inositol phosphates in hepatocytes: role of conversion to ATP and stimulation of P2Y2 receptors. Br J Pharmacol 138: 272–278.
- Communi D, Gonzalez NS, Detheux M, Brezillon S, Lannoy V, et al. (2001) Identification of a novel human ADP receptor coupled to G(i). J Biol Chem 276: 41479–41485.
- Marteau F, Le PE, Communi D, Communi D, Labouret C, et al. (2003) Pharmacological characterization of the human P2Y13 receptor. Mol Pharmacol 64: 104–112.
- Tan C, Salchi A, Svensson S, Olde B, Erlinge D (2010) ADP receptor P2Y(13) induce apoptosis in pancreatic beta-cells. Cell Mol Life Sci 67: 445–453.
- Pandey NR, Sultan K, Twomey E, Sparks DL (2009) Phospholipids block nuclear factor-kappa B and tau phosphorylation and inhibit amyloid-beta secretion in human neuroblastoma cells. Neuroscience 164: 1744–1753.
- Amisten S, Meidute-Abaraviciene S, Tan C, Olde B, Lundquist I, et al. (2010) ADP mediates inhibition of insulin secretion by activation of P2Y13 receptors in mice. Diabetologia 53: 1927–1934.