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Statins perturb G $\beta\gamma$ signaling and cell behavior in a G γ subtype dependent manner

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Inhibition of Gβγ signaling by Statins

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List of nonstandard abbreviations

ARPE-19	- Adult Retinal Pigment Epithelial-19
CHD	- Coronary Heart Disease
CT	- C-terminus
DAG	- Diacylglycerol
DFBS	- Dialyzed fetal bovine serum
DMEM/F-12	- Dulbecco's modified eagle medium/nutrient mixture F-12
FBS	- Fetal bovine serum
FPP	- Farnesyl pyrophosphate
G proteins	- Guanine nucleotide-binding proteins
GGPP	- Geranylgeranyl pyrophosphate

GPCR	- G protein coupled receptor
GRK	- G protein receptor kinases
G γ	- G protein γ
IM	- Internal membranes
IP3	- Inositol 1, 4, 5-triphosphate
MEM	- Minimum essential medium
Nb80	- Nanobody-80
PI3K	- Phosphatidylinositol-4, 5-bisphosphate 3-kinase
PIP2	- Phosphatidylinositol 4, 5-bisphosphate
PLC β	- Phospholipase C β
PM	- Plasma membrane
RPMI 1640	- Roswell Park Memorial Institute medium
SEM	- Standard error of mean
TIRF	- Total internal reflection
vLE	- Leading edge velocity
vTE	- Trailing edge velocity
α 2-AR	- α 2-adrenergic receptor
β 1-AR	- β 1-adrenergic receptor
mCh	- mCherry

Abstract

Guanine nucleotide-binding proteins (G proteins) facilitate transduction of external signals to the cell interior, regulate a majority of eukaryotic signaling, and thus have become crucial disease drivers. G proteins largely function at the inner leaflet of the plasma membrane (PM) using covalently attached lipid anchors. Both small monomeric and heterotrimeric G proteins are primarily prenylated, either with a 15-carbon farnesyl or a 20-carbon geranylgeranyl polyunsaturated lipid. The mevalonate (HMG-CoA reductase) pathway synthesizes lipids for G protein prenylation. It is also the source of the precursor lipids for many biomolecules including cholesterol. Consequently, the rate-limiting enzymes of the mevalonate pathway are major targets for cholesterol-lowering medications and anti-cancer drug development. Although prenylated G protein γ ($G\gamma$) is essential for G protein coupled receptor (GPCR) mediated signaling, how mevalonate pathway inhibitors, statins, influence subcellular distribution of $G\beta\gamma$ dimer and $G\alpha\beta\gamma$ heterotrimer, as well as their signaling upon GPCR activation, is poorly understood. The present study shows that clinically used statins not only significantly disrupt PM localization of $G\beta\gamma$, but also perturb GPCR-G protein signaling and associated cell behaviors. The results also demonstrate that the efficiency of prenylation-inhibition by statins is $G\gamma$ subtype-dependent and is more effective towards farnesylated $G\gamma$ types. Since $G\gamma$ is required for $G\beta\gamma$ signaling and it shows a cell and tissue-specific subtype distribution, the present study can help understand the mechanisms underlying clinical outcomes of statins in patients. This work also reveals the potential of statins as clinically usable drugs to control selected GPCR-G protein signaling.

Keywords: Statins, AKT, Alpha-adrenergic receptors, Complement component 5a receptors, Cell migration, Cholesterol, Desensitization, G protein-coupled receptor kinases (GPKs), G protein-coupled receptors (GPCRs), GTP-binding proteins (G proteins), HMG-CoA reductase, Receptor internalization

Introduction

As a major source of morbidity and mortality in developed countries, cardiovascular diseases, specifically Coronary Heart Diseases (CHD), have been on the spotlight for years. The majority of the patients with cardiovascular diseases are diagnosed with atherosclerosis, which is associated with elevated serum cholesterol levels (Gordon and Kannel, 1971; Sytkowski et al., 1990). Although, a fraction of serum cholesterol comes from dietary cholesterol, it is primarily regulated through hepatic cholesterol biosynthesis by the mevalonate pathway (Hajar, 2011) (Fig. 1A). The rate-limiting step of cholesterol biosynthesis in the liver is the conversion of 5-hydroxy-3-methylglutaryl-coenzyme A (HMG-CoA) to mevalonic acid (Rodwell et al., 1976). The enzyme HMG-CoA reductase catalyzes this process (Goldstein and Brown, 1990). Inhibitors of HMG-CoA reductase, also called as statins, are heavily prescribed to reduce serum cholesterol. Initially, statins were identified as secondary metabolites in fungi, and mevastatin is the first natural statin identified (Alberts, 1988; Endo et al., 1976). All statins contain a region that resembles the HMG moiety and thus bind to the active site of the HMG-CoA reductase enzyme (Istvan and Deisenhofer, 2001). As a result, statins reversibly inhibit the enzymatic activity of HMG-CoA reductase and reduce mevalonic acid production (Istvan and Deisenhofer, 2001).

A number of side-effects associated with statin-use including myopathic effects, short-term memory loss, abnormal liver function, glucose intolerance, hyperglycemia, and increased risk for diabetes have been reported (Davies et al., 2016; Golomb and Evans, 2008; Ramkumar et al., 2016). Up to 25% of statin users report muscle symptoms, cramping, soreness and fatigue, thereby forcing many patients to either switch to lower doses, lower potency statins, or to withdraw (Carris et al., 2017). Statins are also associated with ocular adverse effects including blurred vision, visual impairment, visual field defect, reduced visual acuity, myopia, hypermetropia, presbyopia, and astigmatism (Machan et al., 2012; Mizranita and Pratisto, 2015). While one retrospective study indicated that all statins are associated with ocular side effects, atorvastatin showed higher incidences of muscle and liver problems (Mizranita and Pratisto, 2015). Higher occurrences of age-related cataracts in patients on statin therapy have been observed (Machan et al.,

2012). Furthermore, chronic cholesterol depletion using statins has impaired human serotonin_{1A} receptor function (Shrivastava et al., 2010).

Besides cholesterol, mevalonate pathway synthesizes several important lipid precursors such as polyisoprenoids and sterols (Miziorko, 2011). Isoprenoid lipids-farnesyl pyrophosphate (FPP) and geranylgeranyl pyrophosphate (GGPP)- provide membrane anchoring abilities to many proteins and thus are required for numerous functions including cytoskeletal reorganization, differentiation and proliferation (Buhaescu and Izzedine, 2007; Casey, 1992; Chan et al., 2003). FPP and GGPP are precursor-lipids for G proteins prenylation. GPCRs and G proteins govern numerous biological processes and have become therapeutic targets in areas including cancer, cardiac dysfunction, diabetes, central nervous system disorders, obesity, inflammation, vision, and pain. G protein mediated signaling process consists of GPCRs, heterotrimeric G proteins; G α , G $\beta\gamma$, and their effectors. G α and G $\beta\gamma$ act as switches that transmit information from cell surface receptors to intracellular effectors. N- and C-termini (NT and CT) of G α and G γ respectively are post-translationally modified with lipid anchors, facilitating heterotrimer formation, and their anchoring to the plasma membrane (PM). N-myristoylation and N-palmitoylation mediate G α -PM interactions (Chen and Manning, 2001). We showed that distinct PM-interacting properties of 12 mammalian G γ types are controlled by their CT sequences (Senarath et al., 2018). This sequence comprises a polybasic pre-CaaX region and a CaaX motif (Cook et al., 2006; Watson et al., 1994; Wedegaertner et al., 1995). CaaX motif Cys is prenylated either with a 15-Carbon farnesyl or 20-Carbon geranylgeranyl polyunsaturated lipid anchors (Wedegaertner et al., 1995). Type of prenylation is primarily dictated by the CaaX motif sequence of each G γ subtype (Wedegaertner et al., 1995) and lipid anchors for prenylation are synthesized by the HMG-CoA pathway (Lacher et al., 2017). Not only G γ , but many other Ras superfamily G proteins also use prenylation for membrane targeting. How statins perturb Ras members has been extensively studied (Cordle et al., 2005; Dai et al., 2007; Morck et al., 2009). Anti-cancer effects of statins are assigned to their ability to down regulate oncogenic Ras (Chan et al., 2003; Greenwood et al., 2006; Wong et al., 2002). Several studies have suggested that statins reduce isoprenylation of G γ and subsequently attenuate GPCR signaling (Muhlhauser et al., 2006; Schmechel et al., 2009). However, considering the

necessity of G γ for G $\beta\gamma$ signaling, and their diverse cell and tissue-wide expression, molecular details including whether statins universally and equally inhibit prenylation of G γ -members and attenuate GPCR-G protein signaling are not sufficiently explored.

Materials and Methods

Materials

HeLa, ARPE-19, and RAW 264.7 mouse macrophage cell lines were purchased from American Type Culture Collection (Rockville, MD, USA). Cell culture media; minimum essential medium (MEM), Dulbecco's modified eagle medium/nutrient mixture F-12 (DMEM/F-12), and Roswell Park Memorial Institute medium (RPMI 1640) were obtained from Gibco-Thermo Fisher Scientific (Waltham, MA, USA). Fetal bovine serum (FBS) and dialyzed fetal bovine serum (DFBS) were purchased from Atlanta biologicals (Hall County, Georgia, USA). The reagents; Lipofectamine 2000 (ThermoFisher), Fluvastatin, Atorvastatin, Lovastatin and Gallein (Cayman Chemical, Ann Arbor, MI), Norepinephrine and isoproterenol hydrochloride (Sigma-Aldrich, St. Louis, MO 63178, USA), C5a (ANASPEC, Fremont, CA, USA), Fluo-4 AM (Molecular probes, Eugene, Oregon), 11-*cis*-retinal (National Eye Institute, Bethesda, MD, USA).

DNA constructs

Blue opsin-mTurquoise, α 2-AR-CFP, Akt-PH-mCh, GFP-G γ 9, GFP-G γ 3, G β 1-YFP, YFP-G γ 2, G γ 4, G γ 5, GFP-Rac1, and untagged K-Ras and M-Ras were kindly provided by Professor N. Gautam's lab, Washington University, St. Louis, Missouri. mCh was amplified from Akt-PH-mCh with NheI and HindIII restriction sites and performed the restriction digestion of both Ras isoforms (vector: untagged K-Ras and M-Ras) and amplified mCh (insert). Digested insert and vectors were ligated and chemically transformed to generate N-terminally mCh tagged K-Ras and M-Ras constructs. Nb80 was kindly provided by Dr. Roshanak Irannejad, University of California, San Francisco, California. To generate the Nb80-mCh construct, GRK2-mCh was digested with BamHI and XbaI to remove GRK2. PCR product of Nb80 was then inserted into remaining mCh vector backbone finally producing Nb80-mCh. Venus-GRK3ct was kindly provided by Dr. Nevin Lambert. mCh-G γ 3(C72A) construct was generated by substituting Cys 72 residue in mCh-G γ 3 with an Ala residue employing the overlap PCR method.

Cell culture and Transfections

HeLa cells were cultured in MEM /10% DFBS/ 1% penicillin–streptomycin medium in a humidified 5% CO₂ incubator at 37°C. RAW 264.7 cells were cultured in RPMI 1640 /10% DFBS with 1% penicillin–streptomycin and ARPE-19 cells were cultured in DMEM/F-12/10% FBS/ with 1% penicillin–streptomycin. At 70-80% confluency, the growth medium was aspirated, and cells were incubated with versene-EDTA (CellGro) for 2 min at 37°C. After the incubation, cells were lifted and centrifuged at 1000g for 3 minutes. Versene-EDTA was then aspirated, and the cell pellet was resuspended in the growth medium and seeded on glass-bottomed dishes (In Vitro Scientific) with a 1×10^5 /ml cell density. The following day, after seeding, Lipofectamine 2000 mediated transfection was performed according to the manufacturer's instructions. After 3.5 hours of transfection, cells were replenished with the growth medium containing either dimethyl sulfoxide (DMSO) or respective statins. The next day, live cell imaging was performed.

Live cell imaging, confocal microscopy and data analysis

An imaging system composed of a Nikon Ti-R/B inverted TIRF microscope, a Yokogawa CSU-X1 spinning disk unit (5000 rpm), an Andor FRAP-PA (fluorescence recovery after photo-bleaching and photo-activation) module and iXon ULTRA 897BV back illuminated deep-cooled EMCCD camera was used for live cell imaging and well as optogenetics experiments. Grayscale confocal images were captured on to 512 x 512 pixel CCD with 16 X 16 μm^2 pixel size. Pixel depths of the active area and gain register were respectively 180,000 e- and 800,000 e-. Images were acquired with 50 ms exposure with 4-frame averaging. The EM gain was set for 300. Four, 40–100 mW 445 nm, 488 nm, 515 nm, and 594 nm solid-state lasers and an acousto-optical tunable filter (AOTF) were used for excitation, optical activation (OA) as well as photobleaching. Live cell imaging was performed using 60X, 1.4 NA (numerical aperture) oil objective. Emission filters; 478 \pm 20 nm (Cyan Fluorescent Protein-CFP, mTurquoise), 488 \pm 20 nm (enhanced Green Fluorescence Protein-eGFP), 542 \pm 30 nm (Yellow Fluorescent Protein-YFP, and Venus) and 630 \pm 30 nm (mCherry) were used respectively with 445 nm (\sim 40 μW), 488 nm (\sim 30 μW), 515 nm (\sim 20 μW) and 595

nm (~60 μ W) excitation lasers. The power listed was measured at the focal plane of the 60X objective using 1 cm^2 sensor (Ophir PD300). For optical activation of blue opsin using the FRAPPA raster scanning, selected regions of cells were exposed to 445 nm (0.2-1.0 μ W at the objective) light. Before OA, cells were incubated with 50 μ M 11-cis-retinal for 3-5 min in dark. Digital image analysis was performed using Andor iQ 3.1 software and color values were normalized using in-built Look Up Table tools maintaining a approximately constant setting for each fluorescence sensors. Fluorescence intensities obtained from regions of interest (PM, IMs, and cytosol) were baseline normalized using the intensity of the background. Normalized data were then plotted using Origin pro (OriginLab Corporation). Results of all quantitative assays (G $\beta\gamma$ translocation, PIP2 hydrolysis, and calcium responses) in time-curves were expressed with the standard error of the mean (SEM) from n number of cells (indicated in the figure legends).

Intracellular Ca^{2+} measurements

To measure cytosolic Ca^{2+} , RAW 264.7 cells were cultured on glass-bottomed dishes with RPMI 1640/DFBS at 37 $^{\circ}\text{C}$ with 5% CO_2 as described in cell culture and transfections. Experiments were performed two days after seeding cells on glass-bottomed dishes. The day after the seeding, cells were treated with 10 μ M fluvastatin and incubated overnight in the 37 $^{\circ}\text{C}$ incubator with 5% CO_2 . The following day, cells were washed twice with 1% (V/V) HBSS containing NaHCO_3 and Ca^{2+} (pH 7.2) supplemented with 10 μ M fluvastatin. Washed cells were then incubated for 30 min at room temperature with a fluorescent calcium indicator, Fluo-4 AM (2.28 μ M) and supplemented with 10 μ M fluvastatin in dark conditions. Before start imaging, cells were again washed twice with 1% (V/V) HBSS containing NaHCO_3 , Ca^{2+} and 10 μ M fluvastatin to remove excess Fluo-4 AM. Processed cells were imaged to capture fluorescent intensity of Fluo-4 AM (488 nm) at 1 s intervals using 488 nm excitation-515 nm emission after addition of 20 μ M c5a to activate endogenous c5a receptors. The control cells were treated with the vehicle solvent; DMSO, and imaged the same way as fluvastatin-treated cells.

Western blot analysis

After 12-hour incubation, either with carrier solvent (DMSO) or fluvastatin (2 μ M), RAW 264.7 cells were exposed to 12.5 μ M c5a for 30 mins to activate endogenous c5aR. After the incubation, cells were lysed and the whole cell lysate was subjected to SDS polyacrylamide gel electrophoresis to separate proteins. Proteins were transferred to a PVDF membrane. After blocking with 5% non-fat milk, the membrane was incubated with the primary antibodies specific for p-Akt, t-Akt and β -Actin (Cell Signaling: #4060S, Santa Cruz: sc-5298, sc-47778) overnight at 4 $^{\circ}$ C. The next day, the membrane was incubated with the appropriate HRP-conjugated secondary antibodies (Bio-Rad: #1705047, #1705046) at room temperature for 2 hours. After incubating with chemiluminescent detection solutions, the membrane was exposed to an X-ray film. The protein band intensities on X-ray films were quantified and normalized to the level of β -Actin. Experiments were performed in triplicates with independent biological replicates, and quantification and statistical analysis were performed.

Transwell invasion assay

Transwell invasion chambers (with 8 μ m membrane pores) were coated with 1x Matrigel (100 μ l). The next day, RAW 264.7 cells pre-incubated either with DMSO, Fluvastatin (2 μ M), Rho-A inhibitors (EHop-016 – 5 μ M, EHT1864- 10 μ M, GSK269962- 12.5 μ M) or Wortmannin (50 nM) for 24 hours were transferred on top of Matrigel-coated invasion chambers in serum free RPMI (5×10^4 cells/well). In Transwell chambers also, cells were treated with either DMSO or the above inhibitors. To asymmetrically activate endogenous c5aR, RPMI containing 10% DFBS with 12.5 μ M c5a was added to the lower chambers. After incubation for 24 hours at 37 $^{\circ}$ C and 5% CO₂, the invaded cells on the lower surface of the insert membrane were fixed and stained using the Fisher HealthCare™ PROTOCOL™ Hema 3™ Fixative and Solutions (FisherScientific) according to the manufacturer's protocol. The number of invaded cells were counted under a light microscope at 200X total magnification. Every experiment was performed in triplicate and all experiments were repeated four times over different days.

Statistics and reproducibility

Results of all quantitative assays ($G\beta\gamma$ translocation, PIP3 generation, cell migration, and Ca^{2+} response) were calculated as the mean \pm standard error of the mean (SEM) or standard deviation (SD) with the listed number of cells in the figure legends from multiple independent experiments (>3). $P < 0.05$ was considered as the threshold for the statistical significance. Comparison of distinct cellular conditions such as $G\gamma 9$ and $G\gamma 3$ cells and (or) exposed to several statins were performed using one or two-way ANOVA to assess statistical significance. Statistical analysis of data including Ca^{2+} and G protein translocation responses for control and statin-treated cells were performed using two-tailed unpaired t-test. $P < 0.05$ was considered significant. Both Ca^{2+} and Western blot analysis data were presented as mean \pm standard deviation (SD).

Results

Statins disrupt PM localization and signaling of endogenous Gβγ in HeLa cells:

The goal was to examine if statins interrupt the PM localization of endogenous Gβγ and their signaling activation by GPCRs in HeLa cells. Cells were transfected with Gβ1-YFP by incubating cells with transfection medium containing lipofectamine 2000 reagent for 3.5 hours, and subsequently cells were transferred to the regular culture medium, additionally containing statins, i.e. fluvastatin (2 μM), lovastatin (2 μM) or atorvastatin (20 μM) for 12 hours. These concentrations were selected considering their IC50 values (Martin-Navarro et al., 2013). In place of statins, control cells were treated with DMSO, the vehicle solvent of statins. Confocal time-lapse images of cells were captured using a 60X objective with a 1.49 NA. Subcellular distribution of GFP-Gβ1 was captured using 488 nm excitation and 515 nm emission. Cells treated with fluvastatin (+Flu) exhibited primarily a cytosolic Gβ1 distribution (Fig.1B, bottom), while control cells showed primarily a PM and a somewhat internal membrane (IM) distribution (Fig. 1B, top). Cytosolic GFP-Gβ1 distribution indicates that the associated Gγ lacks the prenyl modification at the Cys residue of the CaaX motif. Prenylated Gβγ usually resides either on the PM or IMs. To obtain a numerical value to quantify the extent of inhibition of PM localization, the ratio between fluorescence intensities of Gβ1-YFP on the PM and cytosol ($F_{PM/Cytosol}$) was calculated. A line profile of fluorescence intensities across the PM to the cytosol was measured (Fig 1B, cartoon and plot) and average fluorescence intensities on the PM and cytosol were calculated. Calculations showed that $F_{PM/Cytosol} \leq 1$ for cells with completely inhibited PM localization of Gβγ and $F_{PM/Cytosol} > 1$ for cells with a partial to no inhibition. Deviation of $F_{PM/Cytosol}$ of a treated cell from its control represents the extent of the presence of cytosolic Gβγ, indicating the extent of presumptive prenylation inhibition. For Gβ1 distribution, $F_{PM/Cytosol}$ was 1.04 ± 0.09 for +Flu cells (Fig. 1B, bar graph). The images show that in the +Flu cell, a distinguishable PM region is lacking since fluorescence in the cytosol and the PM are nearly similar. The plot shows the steep drop in the intensity (red curve) at the edge of the cell to the level of the extracellular background. Overall, contrary to the

control cell, +Flu data suggest strong prenylation inhibition by fluvastatin. Next, the effect of fluvastatin on availability of G protein heterotrimers to be activated by GPCR was examined using GPCR activation induced G β 1 γ translocation. We have previously shown that, upon GPCR activation G β γ translocate from the PM to IMs (Ajith Karunarathne et al., 2012) (Fig. 1C, cartoon). Control HeLa cells exhibited a robust translocation of G β 1-YFP upon activation of endogenous α 2-ARs with 100 μ M norepinephrine (Fig. 1C, top panel and plot). However, +Flu cells exhibited 42.85 % attenuated translocation (Fig. 1C, bottom panel and plot). This suggests it is likely that fluvastatin disrupts endogenous G γ prenylation and thereby attenuates heterotrimer formation as well as G β γ translocation.

Fluvastatin-induced inhibition of G γ PM localization is G γ identity dependent:

The goal was to examine whether fluvastatin equally and universally inhibits PM localization of all G γ types. This is crucial since G γ shows a cell and tissue type specific distribution (Cali et al., 1992). Compared to the control HeLa cells, which exhibited GFP-G γ 9 distribution on the PM, +Flu cells exhibited a completely cytosolic GFP-G γ 9 distribution (Fig. 2A, left images and bar graph). Upon activation of endogenous α 2-AR, control cells exhibited a robust translocation of GFP-G γ 9 to IMs indicating G protein activation and heterotrimer dissociation (Fig. 2A, +NE-top images and plot). On the contrary, fluvastatin treated cells exhibited only a minor increase in their already cytosolic GFP-G γ 9 after α 2-AR activation (Fig. 2A, +NE, +Flu images and plot). Like G γ 9, control HeLa cells expressing geranylgeranylated GFP-G γ 3 also showed a prominent PM distribution (Fig. 2B, bottom left and bar graph). Even after 12-hour exposure to fluvastatin, G γ 3 was primarily seen on the PM and only a minor cytosolic distribution was observed (Fig. 2B, middle left and bar graph). This nearly unperturbed PM-bound GFP-G γ 3 fluorescence in fluvastatin treated cells indicated that perhaps a considerable population of G γ 3 is still prenylated. Activation of endogenous α 2-AR receptors in both +Flu and control cells exhibited GFP-G γ 3 translocation from the PM to IMs, however exhibited a 1/3 lower magnitude in +Flu cells (Fig. 2B, plot). These data suggest that fluvastatin is likely to attenuate prenylations of G γ 3 to a level that induces a detectable

reduction of heterotrimer activation and free G $\beta\gamma$ translocation. To validate that the remaining PM-bound fraction of G γ observed after fluvastatin treatment is still prenylated, subcellular distribution of a prenylation deficient G γ 3 mutant was examined. We generated G γ 3(C72A) mutant to eliminate its prenylation, and in HeLa cells, this mutant exhibited completely a cytosolic distribution (Supplemental Figure 1A). This G γ distribution was similar to the G γ 9 distribution observed in +Flu cells. Therefore, we anticipated that membrane bound G γ are prenylated and cytosolic G γ lacks prenylation.

To confirm that statins inhibit membrane attachment of G γ by inhibiting mevalonate synthesis, HeLa cells expressing either GFP-G γ 3 or GFP-G γ 9 were exposed to fluvastatin (2 μ M) for 12 hours, in the presence and absence of mevalonate (0.5 mM). Contrary to control cells exposed to fluvastatin alone, G γ 9 and G γ 3 in +Flu and mevalonate treated cells showed completely a PM distribution (Fig. 2A and 2B, middle and bottom). These cells also exhibited translocation like untreated cells upon endogenous α 2-AR activation (Fig. 2A and 2B, plots). However, +Flu cells exhibited a 10 folds lower extent of translocation for G γ 9 and ~2 fold lower for G γ 3 than the respective control cells. Additionally, G γ 9 distribution in cells exposed to farnesyl transferase inhibitor, lonafarnib, is similar to that of fluvastatin treated cells (Supplemental Figure 1E). These data collectively show that fluvastatin induced inhibition of G γ membrane-anchoring is due to inhibition of mevalonate and subsequent reduction of isoprenyl pyrophosphate synthesis. This data also clearly shows that the observed lack of G γ PM attachment is not due to a non-specific effect of statins on the cell cycle.

Since G γ 9 and G γ 3 are respectively considered farnesylated and geranylgeranylated based on their CaaX motif sequences, we examined whether the PM interaction of other farnesylated as well as geranylgeranylated G γ types are similarly disrupted by fluvastatin. Like G γ 9, G γ 1 is also identified as a farnesylated protein and +Flu cells exhibited a completely cytosolic distribution of YFP-G γ 1, while control cells showed a prominent PM distribution (Fig. 3A). Except G γ 1, 9 and 11, other G γ types are considered primarily geranylgeranylated. Fluvastatin treated G γ 2 expressing cells exhibited a partial inhibition of YFP-G γ 2 membrane attachment (Fig 3B), which is similar to that of G γ 3. Interestingly, compared to G γ 1,

fluvastatin also induced a more pronounced cytosolic distribution of G γ 4 and G γ 5 (Fig. 3C and D). Nevertheless, control cells expressing these G γ sub-types always exhibited primarily PM distributions (Fig 3A-D). To further support our observation that the PM-localization of farnesylated G γ types are more effectively inhibited by fluvastatin than geranylgeranylated G γ s, the extent of observed inhibitions in +Flu cells across G γ types were normalized to their untreated controls (Supplemental Figure 1B). Out of the four G γ types compared, the highest inhibition of membrane anchorage is observed in G γ 1, while G γ 2, 4 and 5 are less affected by fluvastatin. Overall, the data suggest that presumptively fluvastatin inhibits farnesylation more effectively than geranylgeranylation.

Heterotrimeric G proteins shuttle between the PM and IMs, as heterotrimers at the GPCR-inactive state and as free G $\beta\gamma$ (Ajith Karunarathne et al., 2012). Regardless of the type of G γ in the heterotrimer, inactive-state shuttling allows heterotrimers to maintain a pool at IMs, while the majority of heterotrimers reside at the PM. To examine whether fluvastatin treatment alters relative PM-IM distribution of heterotrimers, HeLa cells co-transfected with the endoplasmic reticulum (ER) marker, CFP-KDEL and either GFP- G γ 9 or GFP- G γ 3 were exposed to fluvastatin. Unlike control cells which show heterotrimers on the PM and in IMs, both G γ types were not observed in IMs in +Flu cells. However, CFP-KDEL expression remained unchanged, indicating an intact ER (Supplemental Figure 1C). This indicates that PM is the primary location for heterotrimers and we anticipate that when cells have a surplus heterotrimers, they are docked in IMs. Since fluvastatin treatment is likely to reduce heterotrimer availability, as indicated by universal reduction of G $\beta\gamma$ translocation (Fig. 2), we hypothesize that +Flu cells can accommodate the available heterotrimers at the PM. Additionally the effect of statins on G α distribution was examined by expressing G α o-CFP, G β 1-mCh and G γ 9-YFP in the same cell. Although G β 1 and G γ 9 showed primarily a cytosolic distribution, G α stayed PM-bound in most cells. (Supplemental Figure 1D).

The mitigation-potentials of G protein-membrane interactions of different statins are diverse:

How different statins perturb G protein-membrane interactions presumptively by reducing G protein prenylation was comparatively examined. In place of 2 μ M fluvastatin, HeLa cells were exposed to either 2 μ M lovastatin or 20 μ M atorvastatin and their effect on GFP-G γ 9 or GFP-G γ 3 distribution in HeLa cells were examined (Martin-Navarro et al., 2013) (Fig. 4A, top panel and plot). A two-way ANOVA shows that G γ 3 and G γ 9 responses to statins significantly differ, $F(1,12) = 41.2$, $p < 0.05$, such that the average inhibition was significantly higher for G γ 9 ($M(\text{mean}) = 51.2\%$, $\text{SEM} = 10.8$) than for G γ 3 ($M = 22.8\%$, $\text{SEM} = 5.4$). The inhibitory effect on G γ membrane interactions of drugs were also significant, $F(2, 12) = 63.2$, $p > 0.05$ such that +Flu effect was significantly higher ($M = 74.6\%$, $\text{SEM} = 10.5$), compared to Lov ($M = 28.1\%$, $\text{SEM} = 7.5$) and Ator ($M = 19.0\%$, $\text{SEM} = 3.4$). The interaction effect was significant too, $F(2, 12) = 8.5$, $p < 0.05$, where Flu on G γ 9 ($M = 92.0\%$, $\text{SEM} = 1.0$) and Flu on G γ 3 ($M = 49.0\%$, $\text{SEM} = 1.0$) show the highest effect while Ator on G γ 3 ($M = 17.0\%$, $\text{SEM} = 3.7$) and Lov on G γ 3 ($M = 11.0\%$, $\text{SEM} = 1.0$) exhibit the lowest effect. Collectively, this analysis shows that G γ 9 and G γ 3 responded to drug treatments differently. Also, at the same confidence level, three drugs influenced each G γ types differentially. Although, the difference between influences of lovastatin and atorvastatin on each G γ types are not significant, they both differ from fluvastatin, i.e. the presumed G γ prenylation inhibition potential of both atorvastatin and lovastatin is significantly lower than fluvastatin.

Whether these statins also attenuate α 2-AR activation induced GFP-G γ 3 or G γ 9 translocation from the PM to IMs was examined (Supplemental Figure 1E). A one-way ANOVA shows that the inhibitory effects of statins on G γ 9 translocations were significant, $F(3, 65) = 30.53$, $p < 0.05$. The inhibitory effects of +Flu and +Lov were significant, where ($M = 1.01$, $\text{SEM} = 0.06$) and ($M = 1.04$, $\text{SEM} = 0.21$) respectively, while the effect of +Ator was not significant ($M = 1.40$, $\text{SEM} = 0.21$). A one-way ANOVA with $F(3, 86) = 9.74$, $p < 0.05$ for the inhibitory effects of statins on G γ 3 translocation was significant. +Flu on G γ 3 translocation ($M = 1.16$, $\text{SEM} = 0.11$) and +Ator ($M = 1.22$, $\text{SEM} = 0.12$) were significant while the effect

of +Lov ($M = 1.36$, $SEM = 0.24$) was not significant. These data collectively suggest that different statins possess distinct abilities to attenuate prenylation of $G\gamma$ and thereby the extents of $G\beta\gamma$ translocation.

Next, the ability of fluvastatin, lovastatin and atorvastatin to attenuate prenylation of Ras superfamily members was examined. The majority of the Ras superfamily proteins are prenylated and interact with lipid membranes where they regulate designated biological functions such as cell proliferation and differentiation (Wang and Casey, 2016). The action of the Ras superfamily proteins are tightly regulated and their aberrant behaviors are found in many cancers (Fernandez-Medarde and Santos, 2011). Together with chemotherapeutic agents, statins are prescribed to accelerate tumor cell apoptosis, since they diminish HMG-CoA pathway metabolites, FPP and GGPP, which are required for Ras protein prenylation (Hindler et al., 2006; Yu et al., 2018). HeLa cells incubated with fluvastatin exhibited a near-complete cytosolic distribution of mCh-KRas as well as mCh-MRas respectively indicating their lack of prenyl anchors (Fig. 4B). This inhibition was quantified using the method in Fig. 1B. Control cells incubated with vehicle solvent exhibited primarily PM bound KRas and MRas. At $p=0.05$, mean $F_{PM/Cytosol}$ ratios of control and +Flu are different where $F_{PM/Cytosol}(\text{control}) \gg 1 \geq F_{PM/Cytosol}(+\text{Flu})$. These results were not surprising since both KRas and MRas are considered as farnesylated (Berndt et al., 2011). Interestingly, HeLa cells exposed to lovastatin and atorvastatin exhibited primarily a PM distribution of KRas and MRas (Fig. 4B, bar graph right). Compared to the near-complete inhibition of PM interaction exhibited by fluvastatin, atorvastatin showed ~40% inhibition of MRas and near zero inhibition of KRas while lovastatin exhibited minor inhibition of both Ras members. Collectively these data suggest that, fluvastatin, lovastatin and atorvastatin exhibit distinct prenylation inhibitions of the same Ras protein.

Additionally, prenylation inhibition of small G-protein Rac1 by fluvastatin, lovastatin and atorvastatin was examined in HeLa cells expressing GFP-Rac1. Rac1 being a member of Rho-GTPases, plays a major role in cell migration by contributing to lamellipodia formation (Cotton and Claing, 2009). While the control cells treated with DMSO showed a clear Rac1 distribution on the PM (Fig. 4C, top right image and pie-chart) fluvastatin treated cells exhibited a nuclear localization (Fig. 4C, bottom left image and pie-chart). The extent of inhibition of PM localization of Rac1 by lovastatin and atorvastatin was

significantly lower than fluvastatin. A one-way ANOVA shows that three statins inhibit Rac1 localization at the PM to significantly different extents, $F(2, 6) = 117.5$, $p < 0.01$. Fluvastatin treated cells showed the highest mean inhibition of 99.2 % (SD = 0.5); while lower inhibitions were exhibited by lovastatin (77.0 % \pm 7.7), and atorvastatin (25.0% \pm 7.2). Additionally, confirming that nuclear-localized Rac1 is unprenylated, cells incubated with lonafarnib (10 μ M), a farnesyl transferase inhibitor, also exhibited nuclear localized Rac1-GFP (Supplemental Figure 2A). Moreover, G γ 9 in lonafarnib treated cells were cytosolic as well.

Ramifications of G $\beta\gamma$ signaling in statin-treated cells:

The majority of G $\beta\gamma$ effectors are PM bound or interact with the PM during activation. The goal was to examine whether fluvastatin disrupts G $\beta\gamma$ effector-signaling significantly. Whether the signaling of G $\beta\gamma$ composed of specific G γ types is more susceptible to be perturbed by fluvastatin was also examined.

(a) *G $\beta\gamma$ induced phospholipase C β (PLC β) activation and subsequent calcium mobilization:*

Compared to G α_q GTP, G $\beta\gamma$ activates PLC β to a lesser extent, however induces phosphatidylinositol 4, 5-bisphosphate (PIP2) hydrolysis, generating diacylglycerol (DAG) and inositol 1, 4, 5-triphosphate (IP3) (Park et al., 1993). We previously demonstrated that upon activation Gi-coupled α_2 -AR using norepinephrine, activates PLC β and induces calcium mobilization (Gupta et al., 2017). Since our data show that statins reduce concentration of PM-interacting G $\beta\gamma$ at near-complete to partial levels, we examined the effect of statins on G $\beta\gamma$ -governed calcium mobilization response. RAW 264.7 cells incubated with either 10 μ M fluvastatin, 20 μ M atorvastatin or 10 μ M lovastatin for 12 hours were pre-treated with the small molecule calcium indicator fluo-4 AM (2.28 μ M). Calcium imaging was performed as described before (Siripurapu et al., 2017). After baseline fluo-4 fluorescence was captured, endogenous c5a receptors were activated with 20 μ M c5a and fluo4 imaging was continued. A one-way ANOVA that yielded $F(3, 36) = 157.3$, $p < 0.05$ shows that compared to the control cells in which 55.3 % (SD = 5.7) cells showed c5A receptor activation induced calcium responses, fluvastatin treated cells showed a significant inhibition of

the response with a mean difference of 40.2 % (SEM = 2.0) cells (Fig.5A, images and bar graph). Not only the % number of cells with calcium increase (15.1% \pm 5.3) was reduced in fluvastatin treated cells, the relative response magnitude was also reduced (Fig. 5A, plot). Lovastatin was able to show a minor inhibition of calcium response while inhibition by atorvastatin was insignificant.

(b) *Gβγ mediated activation of PI3K, subsequent PIP3 generation and cell migration:* Subsequently examined was whether presumptive partial prenylation inhibition of geranylgeranylated Gγ types in RAW 264.7 cells by statins is translated into a reduction in signaling. We have demonstrated that farnesylated Gγ types, including Gγ9, attenuate Gβγ-effectors PLCβ and phosphatidylinositol-4, 5-bisphosphate 3-kinase (PI3K) activations, while geranylgeranylated Gγ types enhance their signaling (Senarath et al., 2018). We also showed that endogenously expressed Gγ3 in RAW cells are required for Gi pathway activation mediated PIP3 production and PLCβ activation during cell migration (Senarath et al., 2018; Siripurapu et al., 2017). In migratory cells, lamellipodia formation at the leading edge is driven by PIP3 produced as a result of Gβγ mediated PI3K activation (Kolsch et al., 2008). In order to interact with Gβγ, p110 subunit of PI3K translocates from the cytosol to the PM and subsequently catalyzes PIP3 production (Braselmann et al., 1997; Brock et al., 2003). We employed the translocation of fluorescently tagged PIP3 sensor (Akt-PH-mCh) from cytosol to the PM to measure PIP3 production. RAW 264.7 cells were transfected with blue opsin- mTurquoise (a Gi-coupled GPCR), Akt-PH-mCh and either GFP-GPI (WT) or GFP-Gγ3. Cells were exposed to either fluvastatin (2 μM), lovastatin (2 μM) or atorvastatin (20 μM) for 12 hours after transfection. Both WT-control and Gγ3-control RAW264.7 cells were treated with DMSO. Cells were supplemented with 11-*cis*-retinal (50 μM) to generate light sensing blue opsin and imaging of GFP using the 488 nm light activated opsin. Plots (Fig. 5B) and two-way ANOVA show that PIP3 responses observed in WT or Gγ3 cells upon blue opsin activation were not significantly different ($F(1, 57) = -0.026, p = 1$). Also, selected drugs to inhibit PIP3 production in WT or Gγ3 cells were not significantly different ($F(4, 57) = 0.754, p = 0.56$). However, all three drugs exhibited significant differences in their PIP3 response

both in WT and Gr3 cells ($F(4, 57) = 5.40$ $p < 0.01$) such that compared to WT-control and Gr3-control cells, fluvastatin treated cells showed respectively a ~73% and a ~82% reductions in PIP3 production. Similarly, lovastatin showed a 38% and 58%, and atorvastatin exhibited 13% and 79% reductions in WT and Gr3-control cells respectively.

We next examined if the observed reduction of PIP3 generation in fluvastatin treated RAW 264.7 cells also impairs optically-directed migration through localized activation of blue opsin to activate Gi-pathway. Blue opsin in localized regions of RAW 264.7 cells (Fig. 6A-blue box) pre-reconstituted with 11-*cis*-retinal was activated using blue light, and cells exhibited directional migration towards the optical stimuli. Compared to the migration of the leading edge of control cells ($6.8 \mu\text{M} \pm 0.8$), which also exhibited a robust lamellipodia formation and PIP3 production (Fig. 6A-yellow arrow), cells exposed to 10 μM fluvastatin exhibited significantly reduced ($p < 0.05$) migration ($2.2 \mu\text{M} \pm 0.2$, (Fig. 6A, bar graph). Fluvastatin treated cells showed a significantly attenuated PIP3 generation, ascertaining the data obtained for PIP3 generation after global activation of opsins (Fig. 5B- yellow arrow). Trailing edge migration of fluvastatin treated cells also exhibited a significantly reduced ($p < 0.05$) migration compared to the control cells.

Cell migration and invasion plays a critical role in cancer cell progression and subsequent cancer metastasis (Kirui et al., 2010). The role of G $\beta\gamma$ signaling in cell migration and invasion is thoroughly investigated and well documented (Kim et al., 2012; Kirui et al., 2010; Tang et al., 2011; Vazquez-Prado et al., 2016). In addition to PI3K, G $\beta\gamma$ signaling also contributes to full activation of PIP3-dependent Rac exchanger 1 (P-Rex1) and thus to Rac1 mediated cell mobility (Dbouk et al., 2012; Kim et al., 2012). Blockade of G $\beta\gamma$ is found to diminish cancer metastasis (Tang et al., 2011). Therefore, the ability of fluvastatin to perturb cell invasion was tested by employing Transwell invasion assay for control, as well as fluvastatin (2 μM), Rho-A inhibitors (EHop-016 – 5 μM , EHT1864- 10 μM , GSK269962- 12.5 μM) or wortmannin (50 nM) treated cells for 24 hours. Compared to the number of cells invaded in control, a one-way ANOVA with $F(3, 9) = 24$, $p < 0.01$ shows that all three treatments significantly inhibited number of

cells invaded (Fig. 6B). Fluvastatin, RhoA inhibitors and wortmannin exhibited, respectively ~11-fold, 75-fold and 6-fold reductions in their invasion. The higher reduction in cell invasion by fluvastatin compared to wortmannin can be attributed to the ability of fluvastatin to attenuate PM localization of both G $\beta\gamma$ as well as Rho GTPases such as Rac1, which facilitate cell migration.

(c) *Fluvastatin exposure attenuates downstream signaling of G $\beta\gamma$:* Akt/PKB pathway is one of the major signaling networks activated by G $\beta\gamma$. G $\beta\gamma$ induced phosphorylation of the PI3K and subsequent PIP3 generation recruits Akt to the PM. Activated Akt triggers numerous downstream responses including cell survival, proliferation, growth and cell migration (Hemmings and Restuccia, 2015). Since our data shows that fluvastatin treatment diminishes G $\beta\gamma$ -mediated PI3K/PIP3 signaling and Akt links G $\beta\gamma$ with cell migration and invasion, Western blot analysis was employed to examine the effect of fluvastatin on Akt phosphorylation. Raw264.7 cells were incubated either with 2 μ M fluvastatin or DMSO, for 24 hours. Cells were lysed and extracted proteins were separated using SDS-PAGE. Western blot analysis was performed using standard protocols and total Akt, phospho-Akt and actin was detected using specific primary and secondary antibodies. The phospho-Akt level was normalized to expression levels of both total-Akt and β -actin. Interestingly the data revealed a two-fold reduction in Akt phosphorylation compared to the DMSO-treated control cells (Fig. 6C, Supplemental Figure 2B). This data further confirms the ability of fluvastatin to inhibit G $\beta\gamma$ signaling and G $\beta\gamma$ -mediated downstream cellular events.

Fluvastatin suppresses internalization of activated β 1 adrenergic receptors:

Desensitization of activated GPCRs protects cells from excessive signaling and is achieved by the receptor internalization (van Koppen and Jakobs, 2004). G protein receptor kinases (GRKs) help physical removal of activated GPCR from the PM by phosphorylating the serine and tyrosine residues in the third cytosolic loop and the CT of the receptor (Evron et al., 2012; van Koppen and Jakobs, 2004). Especially, GRK2 (β -adrenergic receptor kinase 2) phosphorylates agonist-activated β -adrenergic receptors (β 1-AR), promoting

GPCR- β -arrestin interactions and subsequent desensitization (Lefkowitz, 1998). GRK2 activity requires binding of G $\beta\gamma$ (Haga and Haga, 1990; Pitcher et al., 1992). Disruption of G $\beta\gamma$ binding domain of GRK2 eliminates this activation-dependent receptor phosphorylation (Carman et al., 2000). Recently, nanobody-80 (Nb80), a conformation-specific single domain antibody for activated β 1-AR was employed to monitor active-state β 1-AR. Upon activation of β 1-AR with its agonist, it has been shown that cytosolic Nb80-GFP translocates to the PM and binds to the agonist bound activated β 1-AR. When GRK2 induces internalization of activated β 1-AR, bound Nb80 also internalizes and this is indicated by the accumulation of Nb80-GFP fluorescence in IMs (Irannejad et al., 2017) (Fig. 7A, cartoon). Since G $\beta\gamma$ plays a key role in GRK2 mediated receptor internalization, we examined the internalization of activated β 1-AR by monitoring subcellular location of Nb80-GFP in the presence and absence (control) of fluvastatin. To confirm G $\beta\gamma$ involvement, additional experiments were performed in the presence of gallein (G $\beta\gamma$ inhibitor) (Casey et al., 2010), and in cells expressing venus-GRK3ct (to sequester G $\beta\gamma$) (Hollins et al., 2009). HeLa cells expressing β 1-AR-CFP and Nb80-GFP were pre-incubated with 5 μ M fluvastatin or DMSO for 12 hours. Using 20 μ M isoproterenol, β 1-AR was activated and the Nb80-GFP recruitment to IMs was monitored (Fig. 7A, images and plot). A one-way ANOVA shows that compared to control cells with the mean normalized IM fluorescence of 2.061 ± 0.003 , fluvastatin show a significantly lower Nb80 recruitment to IMs (1.523 ± 0.002 , ~50%) at $p < 0.05$. Even more significant reductions (~80%) were observed in gallein treated cells (1.159 ± 0.06) and in GRK3ct expressing cells (1.216 ± 0.002) compared to control cells. Since, the extent of Nb80-recruitment inhibition by fluvastatin is between that of the control and gallein treated as well as GRK3ct cells, fluvastatin mediated G $\beta\gamma$ inhibition in Raw 264.7 cells appears to be partial.

Fluvastatin significantly impairs heterotrimer activation and G $\beta\gamma$ translocation from the PM to IMs in retinal pigment epithelial cells:

Use of statins is associated with ocular diseases including vision impairment, blurred vision and cataract development (Machan et al., 2012; Mizranita and Pratisto, 2015). Visual transduction requires the

activation of G protein heterotrimers by light-activated opsins and cells in the retina primarily employ farnesylated G γ types (G γ 1 and G γ 9) in heterotrimers. Since statins greatly perturb the PM localization of the farnesylated G γ types, we examined whether statins reduce G $\beta\gamma$ translocation (which is considered as a measure of heterotrimer activation (Senarath et al., 2016)) in Adult Retinal Pigment Epithelial-19 (ARPE-19) cells. G γ subtype expression profile of ARPE-19 cells was determined using quantitative real time PCR (qRT-PCR). Data show that the major G γ subtypes endogenously expressed in ARPE19 cells are G γ 11 and G γ 5 (Fig. 7B, bar graph). Since G γ 11 is farnesylated and possesses a low PM affinity (Ajith Karunaratne et al., 2012), GPCR activation induced translocation of G $\beta\gamma$ in fluvastatin-exposed ARPE-19 was examined. ARPE19 cells were transfected with G β 1-mCh and α 2AR-CFP, while exposing to 2 μ M fluvastatin. Control cells treated with DMSO exhibited a prominent distribution of G β 1-mCh on the PM and IMs, while G β 1 distribution was completely cytosolic in cells exposed to 2 μ M fluvastatin (Fig. 7B, top panel). Similarly, inhibition of membrane distribution of G γ 9 was also examined by transfecting ARPE19 cells with mCh-G γ 9. Like G β 1, G γ 9 also exhibited a complete cytosolic distribution in cells exposed to fluvastatin while control cells showed a membrane distribution (Fig. 7B, bottom panel). Next, the effect of fluvastatin on heterotrimer activation in ARPE19 cells was investigated by monitoring mCh-G γ 9 as well as G β 1-mCh translocation from the PM to IMs after activating the transfected α 2AR. Analogous to the observations in HeLa cells, both G β 1 and G γ 9 exhibited a robust translocation from PM to the IM in control cells while fluvastatin treated ARPE-19 cells failed to exhibit detectable translocations of either G β 1 or G γ 9 (Figure 7B, plots).

Discussion

Statins are globally used as effective lipid-lowering medications, primarily to reduce the risk of cardiovascular diseases and associated mortality. However, several side effects such as myopathy, ocular adverse effects including blurred vision and visual impairment, cognitive effects such as short term memory loss, abnormal liver function, glucose intolerance and increased risk for diabetes are reported in patients treated with statins (Davies et al., 2016; Golomb and Evans, 2008; Mizranita and Pratisto, 2015). Being a competitive inhibitor of the enzyme, HMG-CoA reductase; a crucial enzyme of the mevalonate (or cholesterol synthesis) pathway, statins effectively reduce not only the production of cholesterol but also other lipids including isoprenoids (Miziorko, 2011). Isoprenoids are precursors for the synthesis of prenyl pyrophosphates, which are required for the prenylation of a large cohort of G-proteins. The present study draws a clear link between the use of statins and the perturbation of GPCR and heterotrimeric G-protein signaling, providing potential explanations for some of the side-effects related to statin usage (Golomb and Evans, 2008).

Membrane anchorage is a crucial factor for the functioning of Ras superfamily and heterotrimeric G-proteins. Being able to modulate and prevent the activity of Ras superfamily G-proteins, statins exert several pleiotropic side effects such as reduction of cell proliferation that consequently has expanded its usage as a chemotherapeutic agent (Liao and Laufs, 2005). The data presented in this study exhibit that statins perturb the PM localization of $G\gamma$ and thereby $G\beta\gamma$ in a $G\gamma$ subtype dependent manner, as a result of the reduction of prenyl anchors required for $G\gamma$ prenylation (Hildebrandt, 2011; Wedegaertner et al., 1995). Interestingly, data show that $G\gamma$ types that are being prenylated with geranylgeranyl anchors are only partially susceptible to statin-mediated likely inhibition of prenylation. This differential inhibition is also characterized by near-complete inhibition of PM-localization of farnesylated $G\gamma$, attenuating activation of heterotrimers with farnesylated $G\gamma$. The effectiveness of statins to inhibit farnesylation is also evident from

the near-complete inhibition of primarily farnesylation-sensitive Ras family proteins such as KRas, NRas and HRas.

It has been suggested that the CT sequence that determines the type of prenylation is not just limited to CaaX motif (Senarath et al., 2018). Also suggested was the ability of proteins that usually prefer farnesylation can undergo geranylgeranylation, in the presence of farnesyl transferase inhibitors (Whyte et al., 1997). We hypothesized that, G γ types are not completely farnesylated or geranylgeranylated. Although nine out of twelve G γ types are primarily identified as geranylgeranylated proteins, to a certain degree they are also farnesylated as well. It appears that the major anchoring type of a G γ is recognized as the type of prenylation. Our experimental data show varying degrees of likely inhibition G γ geranylgeranylation upon exposure to statins supporting this hypothesis. For instance, extents of the presence of cytosolic G γ indicates a minor G γ 2, G γ 3, G γ 4 and G γ 5 prenylation inhibitions. This also suggests that a considerable fraction in G γ 4 as well as G γ 5 are likely to be farnesylated. A larger reduction of NE-induced G γ 3 translocation compared to that of G β 1 by fluvastatin was observed, although their subcellular distribution data shows a relatively a lesser inhibition of G γ 3 prenylation by fluvastatin. We believe that this is due to partially cytosolic YFP-G β 1 interfering with the detection of YFP-G β 1 translocation. The observed differential effects of statins on farnesylated and geranylgeranylated G γ , likely indicating distinct prenylation inhibition patterns, can be partly due to the higher incidence of GGPP production. This is in addition to the conversion of FPP to GGPP by GGPP synthase. Examination of cholesterol synthesis pathway indicates that, compared to FPP, a cell can have a relatively higher population of GGPP through the direct conversion of isopentenyl-PP to GGPP. Alternatively, in addition to being utilized to produce farnesylated proteins, FPP is also consumed to produce other lipids in the mevalonate pathway, including GGPP and cholesterol. Collectively, these differences can explain why G γ types differentially susceptible to prenylation inhibition by statins (Fig. 1A). Interestingly, the near-complete inhibition of both G γ 1 and G γ 9 suggests that, in the absence of farnesylation, these G γ types cannot effectively undergo geranylgeranylation.

Translocation of G $\beta\gamma$ in cells expressing different G γ subtypes upon the activation of GPCR indicates activation G proteins (Ajith Karunarathne et al., 2012). Attenuated-translocation observed in statin-exposed cells in a G γ -dependent manner, thus signifies the reduction of heterotrimer activation. Therefore, it is likely that heterotrimeric G protein signaling is similarly perturbed in patients treated with statins. G γ types show cell and tissue-specific subtype distributions (Cali et al., 1992). Our data therefore suggest that G protein signaling perturbation by statins can differentially influence GPCR-G protein controlled-processes in a tissue and organ function specific manner. Due to the likely-extensive inhibition of G γ farnesylation by statins, pronounced after effects in tissues and organs that predominantly employ farnesylated G γ types can be anticipated. For instance, photoreceptor cells in the retina of the eye express G γ 1 and G γ 9 as the major G γ (Pearing et al., 2013). This unique G γ expression can be understood by examining the visual transduction in which GTP bound G α -transducin is the major signal transducer, generated by the activated opsin to trigger cGMP signaling. Here, cells appear to employ G $\beta\gamma$ with farnesylated G γ due to their low PM affinity to prevent unnecessary as well as potentially deleterious G $\beta\gamma$ signaling for photoreceptor cells. If statins significantly reduce farnesylation, consequently that can reduce transducin-containing heterotrimer formation as well, hampering cGMP pathway activation. They also reported recovery from the effects upon statin discontinuation (Leuschen et al., 2013). These observations are consistent with the ability of statin to temporarily inhibit HMG-CoA. Data from ARPE19 cells with primarily farnesylated endogenous G γ showed significant inhibition of PM localization and GPCR activation-induced G $\beta\gamma$ translocation upon exposure to fluvastatin. This also indicates the likely after-effects of statins on the vision system. Based on these facts and data, we propose that the dominant expression of farnesylated G γ types in photoreceptor cells is linked to the occurrence of ocular disorders in the patients using statins.

Several studies have shown interference of immune responses by statins due to the attenuated production of downstream metabolites in HMG-CoA pathway including FPP and GGPPs. Similar to G γ , these lipids are required for PM-anchoring of small GTPases such as RhoA, Rac, and Ras. Our data show an impairment of their PM-localization in cells exposed to statins. Our data also show that statins induce

perturbation of G $\beta\gamma$ mediated cell migration and invasion, which are critical steps of immune responses and metastasis. Prolonged exposure to statins including fluvastatin has been shown to negatively impact cognitive function (Golomb and Evans, 2008; Ramkumar et al., 2016). Out of 12 gamma subtypes, G γ 2 and G γ 3 are predominant in the brain (Yim et al., 2017), the partial inhibition of their membrane localization by fluvastatin and the consequent perturbation of associated signaling can partially be responsible for the development of cognitive effects in patients using statins. Our previous findings show that Gi-pathway governed PIP3 production is G γ subtype dependent and G γ 3 is the foremost G γ that supports PIP3 production (Senarath et al., 2018). Partially delocalized G γ 3 from the PM to the cytosol upon fluvastatin treatment thus explains the attenuation of downstream signaling of G $\beta\gamma$ including Akt phosphorylation, cell migration and invasion.

Cardiac muscle activities are strongly regulated by GPCRs including β -adrenergic, acetylcholine and angiotensin receptors, thus they have become therapeutic targets for cardiovascular diseases (Bathgate-Siryk et al., 2014). Agonist activated cardiac β -adrenergic receptors undergo desensitization upon phosphorylation. This process is controlled by G $\beta\gamma$ and GRKs, attenuating GPCR signaling (Reiter and Lefkowitz, 2006). Upregulation of GRK isoforms such as GRK2 and GRK5 markedly cause chronic heart failures due to reduction of β -adrenergic receptors in cardiac muscles (Lymperopoulos et al., 2012). Patients treated with statins have exhibited substantially lower cardiac problems, including reduced damages from ischemia-reperfusion injury (IRI). Excessive β -adrenergic receptor internalization is a leading cause of IRI (Tilley and Rockman, 2006). G $\beta\gamma$ triggers GRK2-induced β adrenergic receptor phosphorylation, marking it for internalization (Cahill et al., 2017; Ribas et al., 2007), thus G $\beta\gamma$ signaling is deleterious during a myocardial infarction (Tilley and Rockman, 2006). Our data not only show that statins reduce G $\beta\gamma$ signaling, but also exhibit a significant reduction in β -adrenergic receptor desensitization. Our findings therefore suggest added beneficial effects of statin on cardiac function, in addition to their ability to lower cholesterol bio-synthesis.

In summary, we show for the first time that statins perturb GPCR-G protein signaling in a $G\gamma$ type dependent manner. Our data conclusively show that, fluvastatin inhibits prenylation of farnesylation-sensitive $G\gamma$ strongly. This also suggests that, functions in cells and tissues predominantly expressing these $G\gamma$ types are more susceptible to be interfered by statins. Additionally, we show that several crucial $G\beta\gamma$ -controlled cellular processes including chemokine cell migration and internalization of β -adrenergic receptors are significantly attenuated by statins. Our findings also indicate that numerous GPCR-G protein governed cellular functions, whether beneficial or not, are likely to be differentially perturbed by statins, and thus require further investigations.

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Authorship Contributions

Participated in research design: Karunarathne, Tennakoon and Kankanamge

Conducted experiments: Tennakoon, Kankanamge and Senarath

Performed data analysis: Tennakoon, Kankanamge, Senarath and Fasih

Wrote or contributed to the writing of the manuscript: Tennakoon, Karunarathne, Kankanmge, Senarath and Fasih

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Footnotes

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Legends for Figures

Figure 1.

G protein prenylation and fluvastatin induced inhibition of Gβ1γ localization on the PM. (A) Major steps of the hepatic cholesterol biosynthesis/mevalonate pathway. By inhibiting the rate-limiting step enzyme; HMG-CoA reductase, statins reduce biosynthesis of many lipids including cholesterol. (B) Quantification of statin-mediated inhibition of G protein PM localization in living cells. Images of HeLa cells expressing Gβ1-YFP. Cells treated with 2 μM fluvastatin exhibited a complete cytosolic distribution of Gβ1, while the control cells showed primarily PM and IM-localized Gβ1-YFP. Cartoon shows how the line-profile data was obtained to calculate the G proteins distribution ratio ($F_{PM/Cytosol}$) in control and fluvastatin treated cells. Bar graph shows a $F_{PM/Cytosol} < 1.0$ for fluvastatin treated cells since Gβ1 is cytosolic, likely due to lack of prenylation (error bars: SEM, n=12 cells, $p < 0.05$). (C) Left: GPCR activation induced translocation of Gβγ dimer from the PM to IMs. In GPCR-inactive state, G protein heterotrimers reside on the PM and upon activation, heterotrimers dissociate, generating Gα-GTP and free Gβγ. The resultant Gβγ then can translocate from the PM to IMs in a Gγ-type dependent manner. Middle: Images show translocation of Gβ1 in control HeLa cells upon activation of endogenous α2-AR with 100 μM norepinephrine (NE). Fluvastatin treated cells exhibited ~50% attenuated translocation. Right: The plot shows the translocation dynamics of Gβ1. Yellow arrow indicate translocated Gβ1 on IMs (scale: 5 μm, error bars: SEM, n=20 cells, two tailed t-test was performed after the time point the response reached equilibrium and the data are statistically significant at 0.0001 level).

Figure 2.

Fluvastatin differentially attenuates the PM localization of Gγ9 and Gγ3. (A) Images of control cells pretreated with the carrier solvent (DMSO) exhibited a clear distribution of GFP-Gγ9 on the PM, indicating prenylated Gγ9. Fluvastatin treated cells showed near-complete cytosolic GFP-Gγ9, suggesting a complete

inhibition of G γ 9 prenylation. +Flu + Mevalonate cells show that supplementation of mevalonate abrogates fluvastatin action. Bar graph show that of +Flu-cells with $F_{\text{PM/Cytosol}} \leq 1$, exhibiting lack of GFP fluorescence on the PM while both Control and +Flu + Mevalonate cells with $F_{\text{PM/Cytosol}} > 1$ indicating the PM-bound GFP fluorescence. Images (left) and the plot show endogenous α 2-AR activation mediated G γ 9 translocation only in Control and +Flu + Mevalonate cells. t-statistics show that at $p < 0.05$, mean $F_{\text{PM/Cytosol}}$ are not significantly different for control and +Flu + Mevalonate cells. In contrast, translocation in fluvastatin treated cells was significantly attenuated, indicating lack of PM-bound heterotrimers (error bars: SEM, n=12 cells, two tailed t-test was performed after the time point the response reached equilibrium and the data are statistically significant at 0.05 level). (B) Partial prenylation inhibition of GFP-G γ 3 in HeLa cells. Similar to GFP-G γ 9 expressing cells, Control and +Flu + Mevalonate cells showed that GFP-G γ 3 is on the PM. In contrast to G γ 9, +Flu-cells showed GFP-G γ 3 distribution only on the PM while no G γ 3 residing in IMs is observed. Upon endogenous α 2-AR activation, cell in all three conditions exhibited translocation. The plot (right) shows a $\sim 1/3$ -magnitude lower translocation in +Flu cells compared to control and +Flu + Mevalonate cells. This indicates that fluvastatin can attenuate G $\beta\gamma$ 3 activation. Yellow arrows indicate G $\beta\gamma$ translocated to IMs (scale: 5 μm , error bars: SEM, n=15 cells, two tailed t-test was performed after the time point the response reached equilibrium and the data are statistically significant at 0.05 level).

Figure 3.

Fluvastatin has distinct inhibition abilities of membrane localization of G γ 1, G γ 2, G γ 4, and G γ 5.

HeLa cells expressing (A) YFP-G γ 1 (B), YFP-G γ 2 (C), YFP-G γ 4 (D) and YFP-G γ 5 (E), pretreated overnight with 2 μM fluvastatin during transfection. Control cells showed a clear localization of G γ on the PM and in IMs. Fluvastatin treated cells showed a complete cytosolic distribution of G γ 1. G γ 2, G γ 4, and G γ 5 transfected cells exposed to fluvastatin showed only a partial presence of YFP in the cytosol, similar

to GFP-G γ 3 cells exposed to fluvastatin. Bar graphs show $F_{\text{PM/Cytosol}}$ in control and +Flu cells in which G γ 1, G γ 2 and G γ 4 show a significant increase in cytosolic YFP compared control cells. However, no significant difference in $F_{\text{PM/Cytosol}}$ of G γ 5 cells was observed between control and +Flu conditions. (scale: 5 μ m, error bars: SD, 10 < n < 15 cells, p < 0.05).

Figure 4.

Different statins show distinct inhibitions of the PM localization of the same G protein. (A) Compared to fluvastatin treated cells, HeLa cells expressing GFP-G γ 9 and GFP-G γ 3 treated with either lovastatin or atorvastatin only exhibited a minor inhibition of the G γ -PM localization. The plot shows percentage of cells showed at least partial membrane localization inhibition in the GFP-G γ expressing cell populations. Symbols ■, ★ and ● on the plot indicate complete, moderate, and minor inhibitions respectively (10 < n < 30 cells). A two-way ANOVA shows that G γ 3 and G γ 9 significantly differ, $F(1,12) = 41.2$, $p < 0.05$, such that the average inhibition was significantly higher for G γ 9 ($M = 51.2\%$, $SEM = 10.8$) than for G γ 3 ($M = 22.8\%$, $SEM = 5.4$). The inhibitory effect on G γ membrane interactions of drugs were also significant, $F(2, 12) = 63.2$, $p > 0.05$ such that +Flu effect was significantly higher ($M = 74.6\%$, $SEM = 10.5$), compared to Ator ($M = 28.1\%$, $SEM = 7.5$) and Lov ($M = 19.0\%$, $SEM = 3.4$). (B) HeLa cells expressing mCh-KRas and mCh-MRas pre-treated with either DMSO (control) or fluvastatin. Compared to control cells, both KRas and MRas in +Flu cells showed a complete cytosolic fluorescence distribution, indicating the near-complete inhibition of their farnesylation. Bar graph (middle) with $F_{\text{PM/Cytosol}}$ indicates significant differences in inhibitions between control and +Flu cells in both KRas and MRas cells. Bar graph (right) with % number of cell with PM localization. +Flu shows effective inhibitions of both Ras members while atorvastatin shows ~40% inhibition of MRas and near zero inhibition of KRas. + Lov exhibited minor inhibition of both Ras members (scale: 5 μ m, error bars: SEM, n=17 cells, p < 0.05). (C) HeLa cells expressing GFP-Rac1 showed a clear PM localization

while in +Flu cells showed GFP fluorescence primarily in the nucleus. Interestingly, +Lov and +Ator treatments induced bi-phasic responses, i.e. only a fraction of cell population exhibited a complete nuclear localization of Rac1. (scale: 5 μ m, error bars: SEM, $n \geq 15$ cells, $p < 0.05$)

Figure 5.

Fluvastatin perturbs G β γ signaling in RAW 264.7 macrophages. (A) Upon activation of endogenous c5a receptor with 20 μ M c5a, 55% of control (DMSO treated) RAW 264.7 cells exhibited increase in fluo-4 fluorescence (green), indicating cellular calcium increase. On the contrary, only ~11% of 10 μ M fluvastatin treated cells showed weaker calcium responses. The plot compares representative calcium responses by a control and +Flu cell. Bar graph shows that, compared to control cells, +Flu, +Ator and +Lov treated cells exhibited varying levels of calcium response inhibitions. However, responses observed +Ator cells were not significantly different from that of control cells (scale bar: 10 μ m, number of fields ≥ 6 , $n < 30$ cells in each field, error bars: SD, $p < 0.05$). (B) RAW 264.7 cells expressing blue opsin-mTurquoise (Gi-coupled GPCR), Akt-PH-mCh (PIP3 sensor), and either GFP-GPI or GFP-G γ 3 pre-incubated with 50 μ M 11-*cis*-retinal. Both WT Control and GFP-G γ 3 cells showed PIP3 generation, indicated by the translocation of cytosolic Akt-PH-mCh to the PM (Yellow arrow). Fluvastatin exhibited the highest inhibition in both WT Control and GFP-G γ 3 expressing cells. Plots and a two-way ANOVA showed that, the PIP3 production inhibition by all three drugs are statistically significant. (scale bar: 5 μ m, error bars: SEM, $n \geq 15$ cells, $p < 0.05$).

Figure 6.

Fluvastatin attenuates Gi-pathway governed cell migration, invasion and reduces Akt phosphorylation. (A) A RAW 264.7 cell expressing blue opsin-mTurquoise and Akt-PH-mCh and incubated with 50 μ M 11-*cis*-retinal exhibited a directional migration towards the optical stimuli upon localized optical activation of blue opsin (blue box). This migration is accompanied with lamellipodia and

clear generation of PIP3 at the leading edge. In contrast, cells treated with 10 μ M fluvastatin exhibited a significant reduction ($p < 0.05$) in cell migration, lamellipodia formation, and PIP3 generation (error bars: SD, $n > 10$ cells, $p < 0.05$). (B) Bar graph shows invasion of RAW 264.7 cells in the transwell invasion assay, induced by endogenous c5a receptor activation using 12.5 μ M c5a. Compared to control (DMSO) cells, fluvastatin, RhoA inhibitors (EHop-016, EHT1864, and GSK269962), and PI3K inhibitor (wortmannin) treated cells exhibited significant reductions in cell invasion ($n=2$). Compared to wortmannin treated cells, fluvastatin treated cells exhibited ~50% more inhibition, indicating that inhibition through fluvastatin can be due to prenylation inhibition of both G γ as well as Ras family proteins (error bars: SD, $n < 3$ independent experiments, $p < 0.05$). (C) Examination of Akt phosphorylation in cells exposed to fluvastatin using Western blot analysis. Images and bar graph show ~ 40% less Akt phosphorylation in +Flu cells, compared to control cells. Protein expressions were normalized to both total Akt and β -Actin. (error bars: SD, $n =$ three independent experiments, $p < 0.05$).

Figure 7.

Fluvastatin (A) attenuates internalization of activated β 1-adrenergic receptors in HeLa cells, and

(B) impairs G $\beta\gamma$ -mediated signaling in ARPE-19 cells. (A) The schematic representation of Nb80

translocation to activated β 1-AR and its subsequent internalization with the receptor. Images show the Nb80-GFP movement in HeLa cells expressing β 1-AR-CFP upon receptor activation. After activation of β 1-AR with 20 μ M isoproterenol, control cells showed a robust recruitment of Nb80 (both GFP and mCh tagged) to the IMs with the internalized and active β 1-AR (top image panel, yellow arrows). A comparatively reduced Nb80 recruitment was observed in +Flu cells (second panel right, white arrow). The plot shows ~ 50% reduced mean GFP fluorescence of Nb80 in the IMs in fluvastatin treated cells indicating attenuated β 1-AR desensitization ($n = 20$ cells from 4 independent experiments). Both 10 μ M gallein (G $\beta\gamma$ inhibitor) treated cells (third panel right, white arrow) and membrane targeted venus-GRK3ct (mask G $\beta\gamma$) expressing cells showed ~80% lower Nb80 accumulation in IMs (white arrows), indicating

the involvement of G $\beta\gamma$ in the pathway. Both Nb80-GFP and -mCh versions exhibited similar internalization in control cells (red and black curves respectively). (B) Real time PCR quantification of relative expression levels of the twelve G γ subtypes in ARPE-19 cells. Farnesylated G γ 11 showed the highest expression. Images show ARPE-19 cells expressing blue opsin-mTurquoise either with G β 1-mCh or mCh-G γ 9 and incubated with 50 μ M 11-*cis*-retinal. G β 1 and G γ 9 in control cells were primarily found on the PM (white arrows) and upon blue opsin activation both proteins exhibited robust translocations from the PM to IMs (yellow arrows). Likely due to prenylation inhibition, both G β 1 as well as G γ 9 distribution in +Flu cells were completely cytosolic and their translocations upon blue opsin activation were not detected. Collectively, these data indicate the lack of prenylation of endogenous G γ types as well as transfected G γ 9 upon fluvastatin treatment. (scale: 5 μ m, error bars: SEM, n = 16 cells, p < 0.05).

Figures

Figure 1

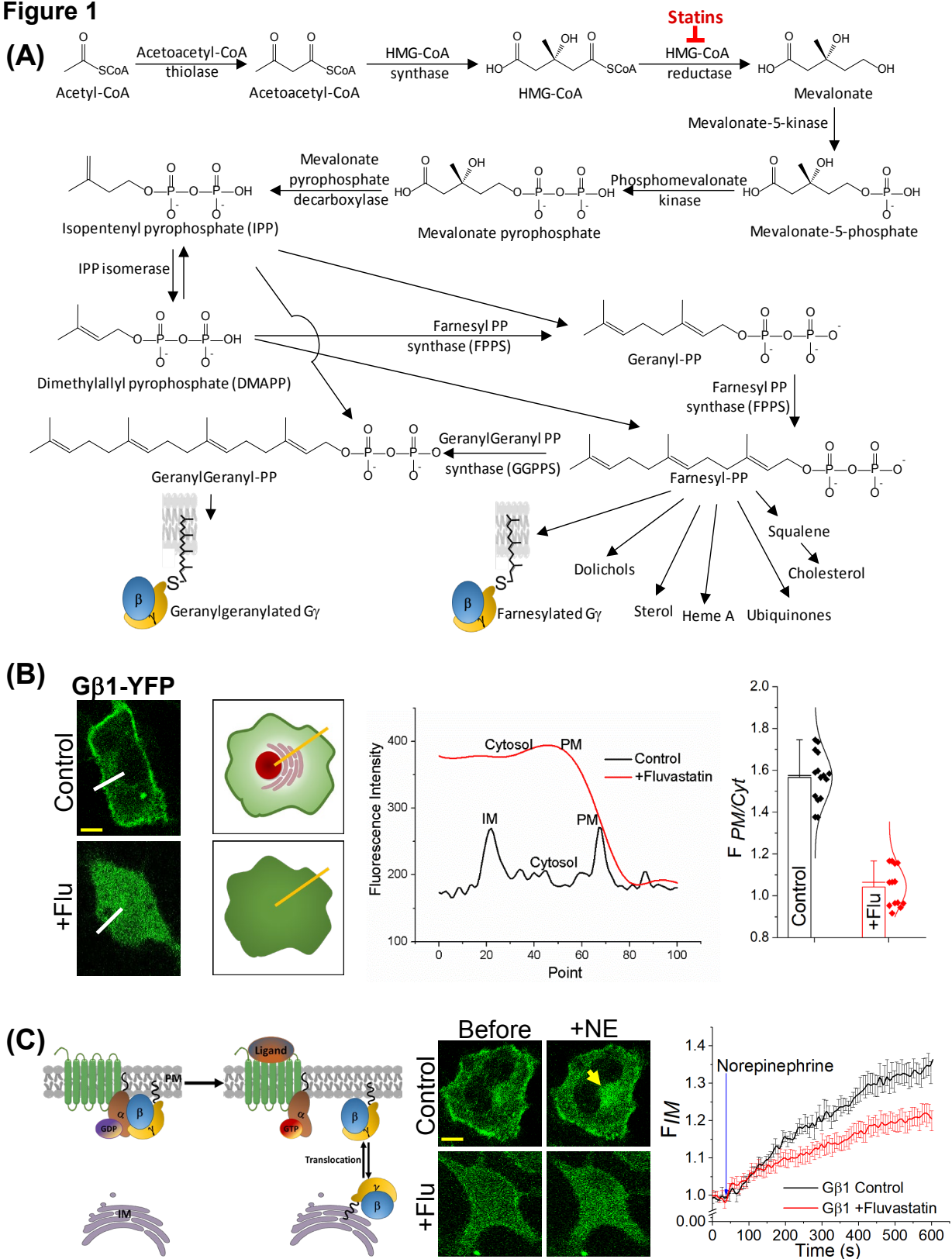


Figure 2

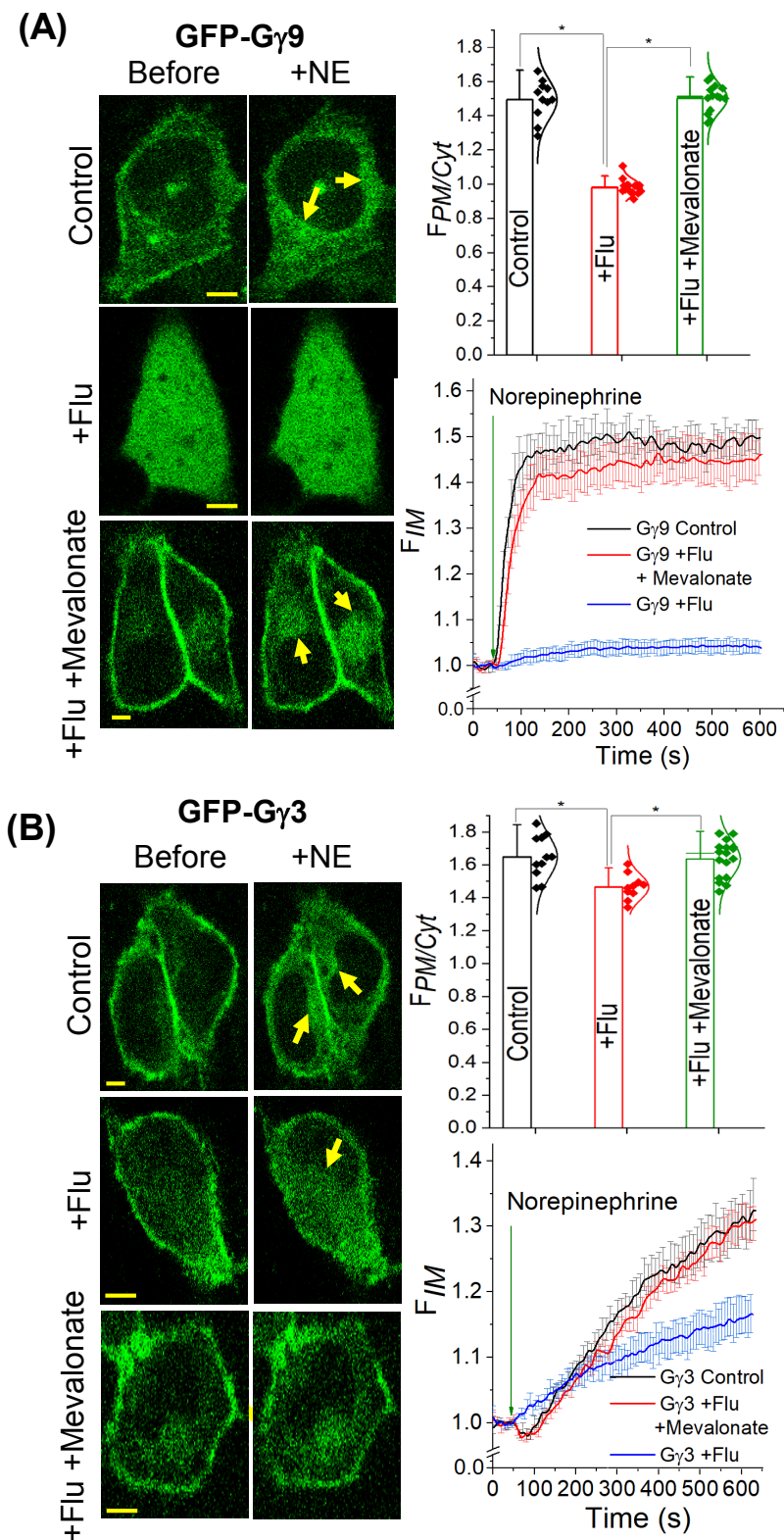


Figure 3

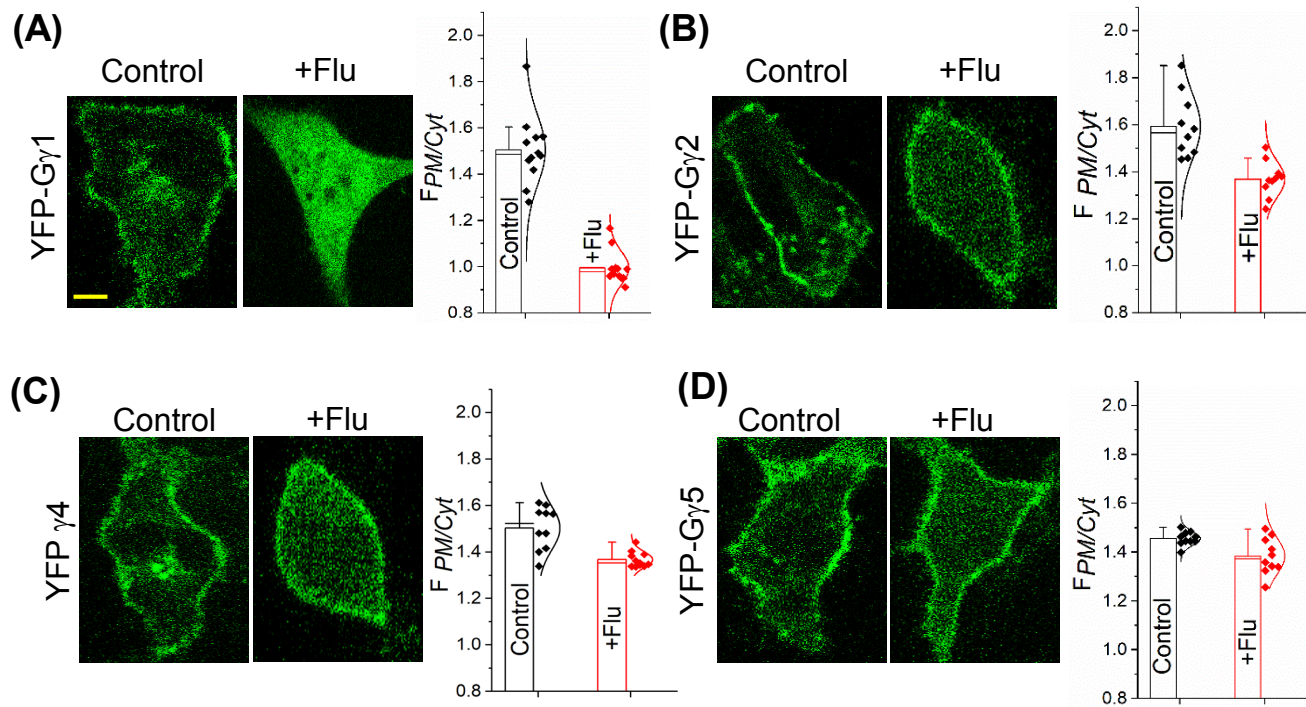


Figure 4

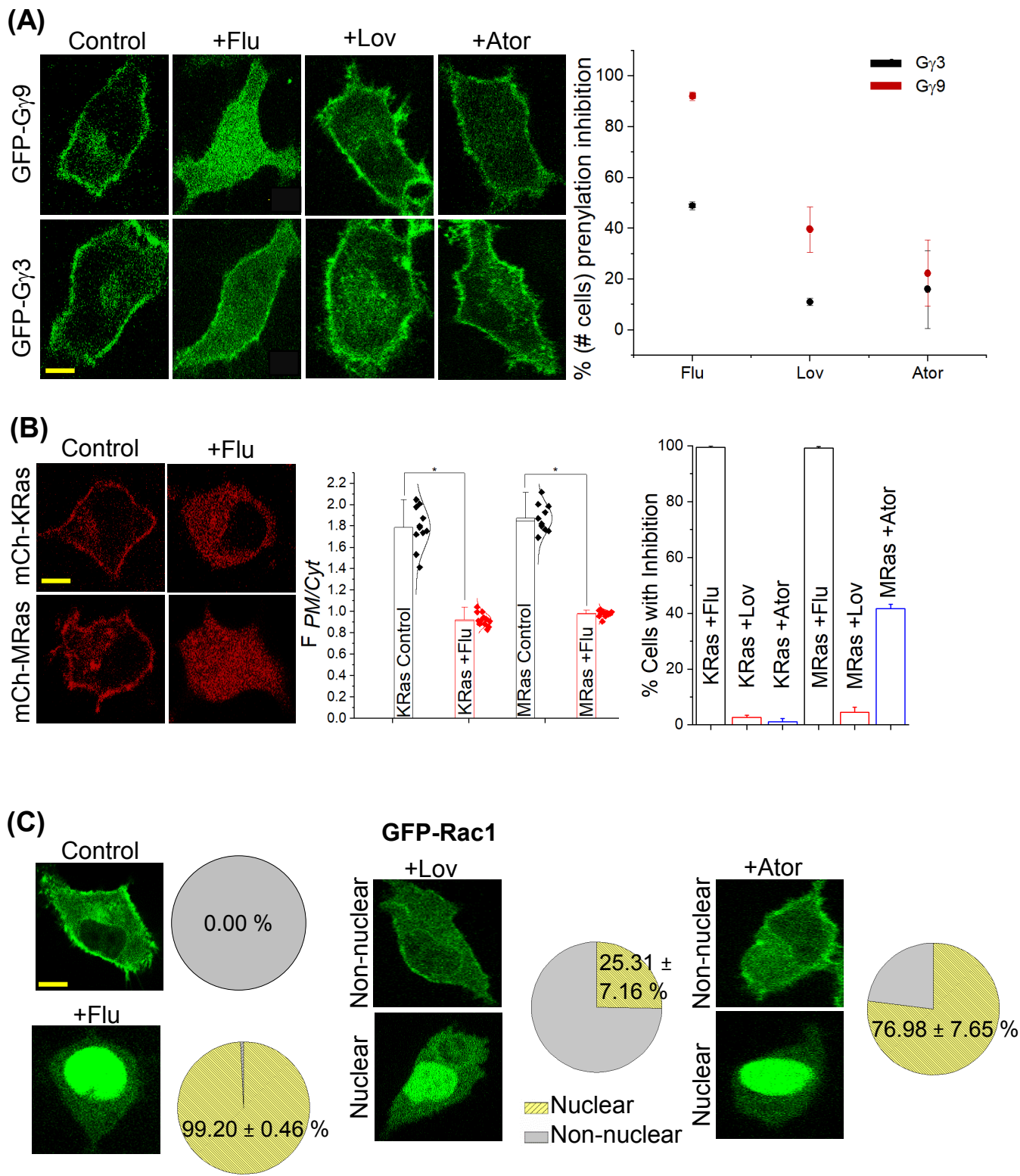
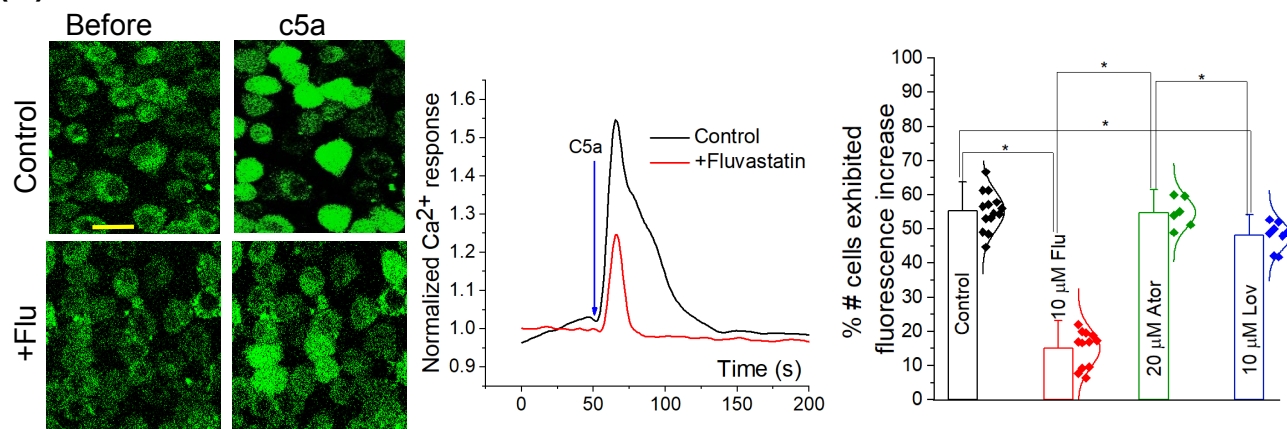


Figure 5

(A) RAW 264.7



(B)

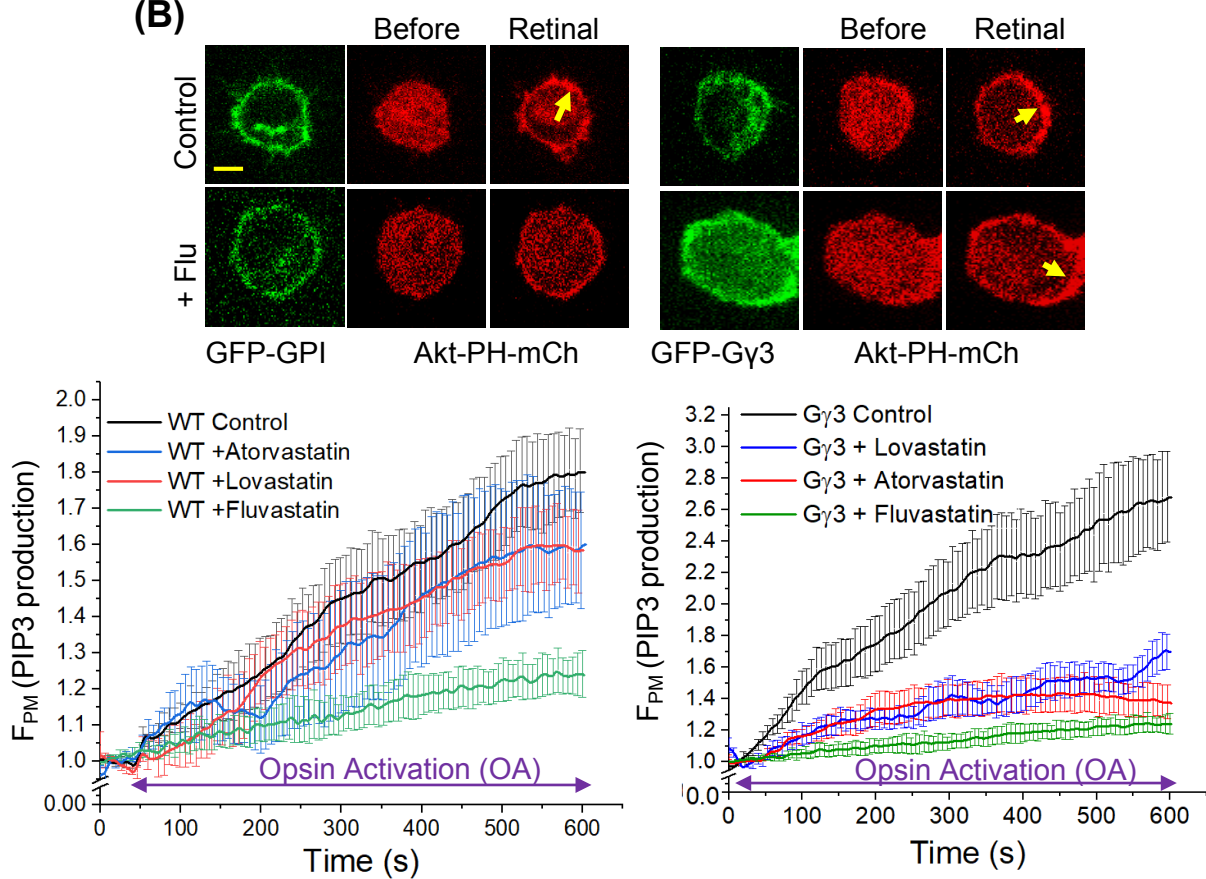
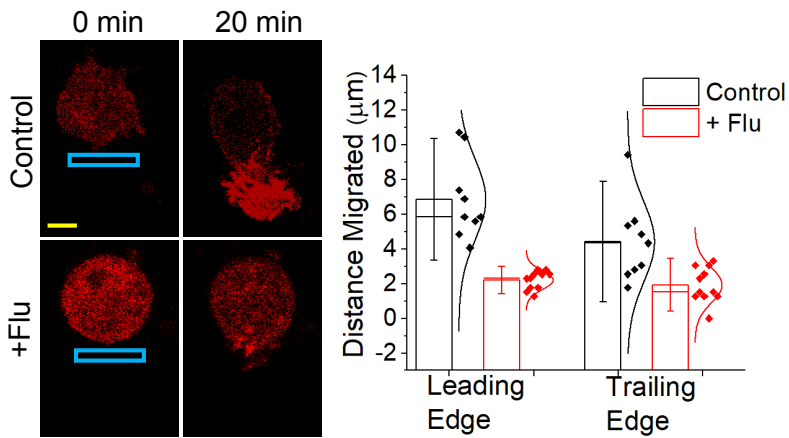
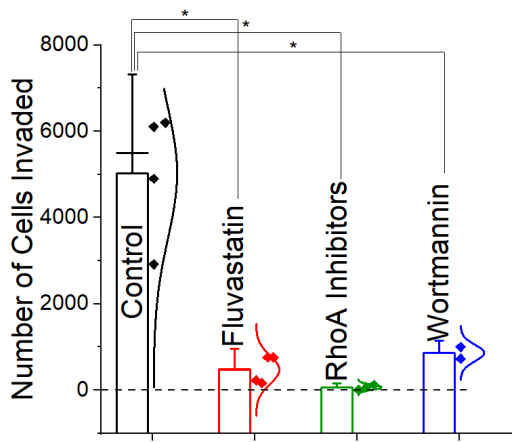


Figure 6

(A)



(B)



(C)

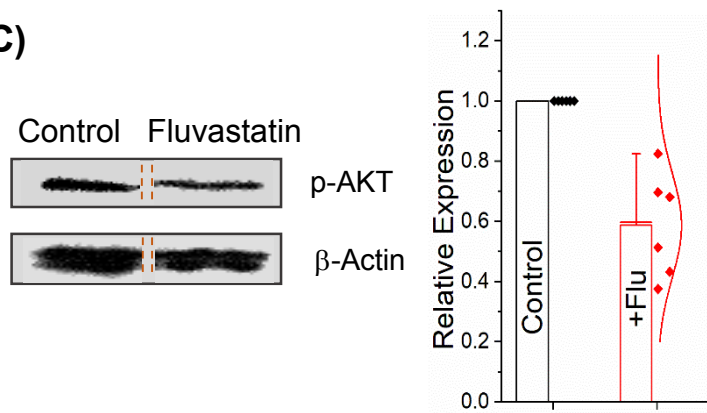
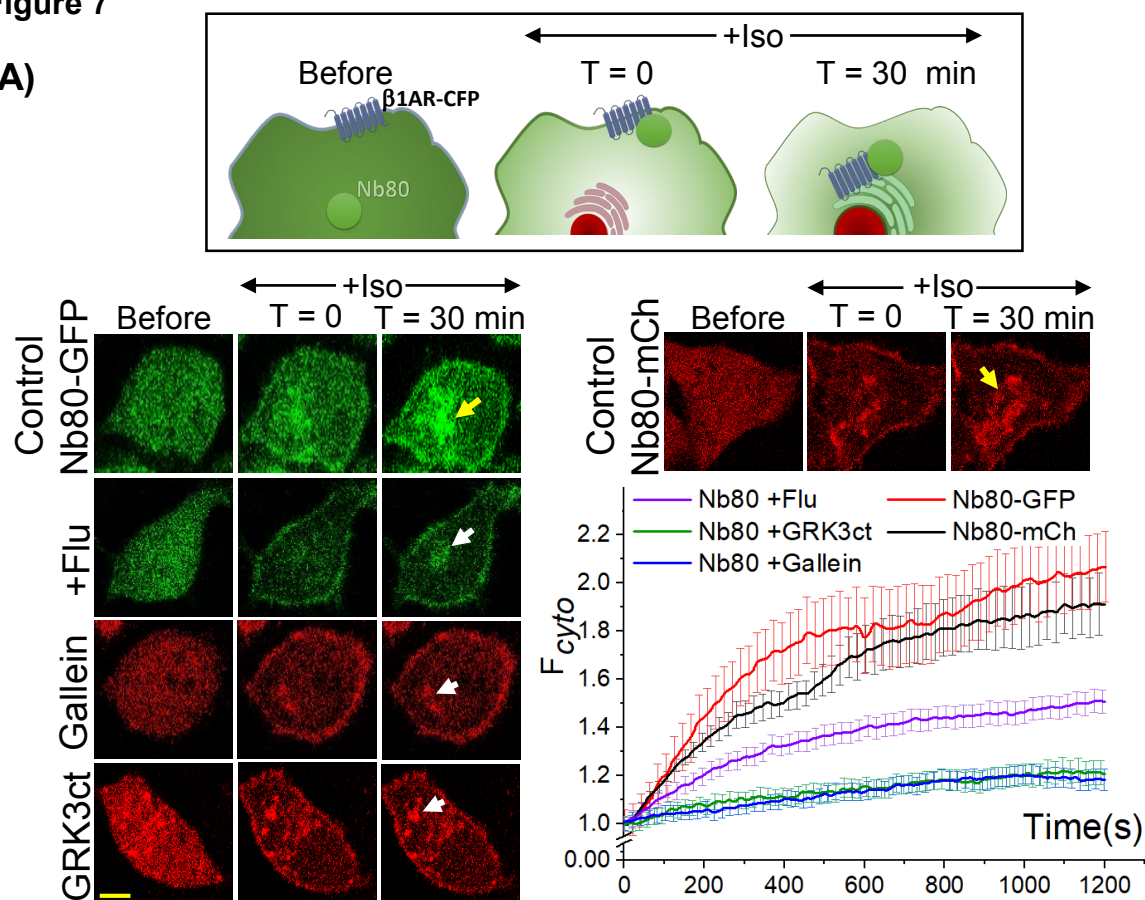
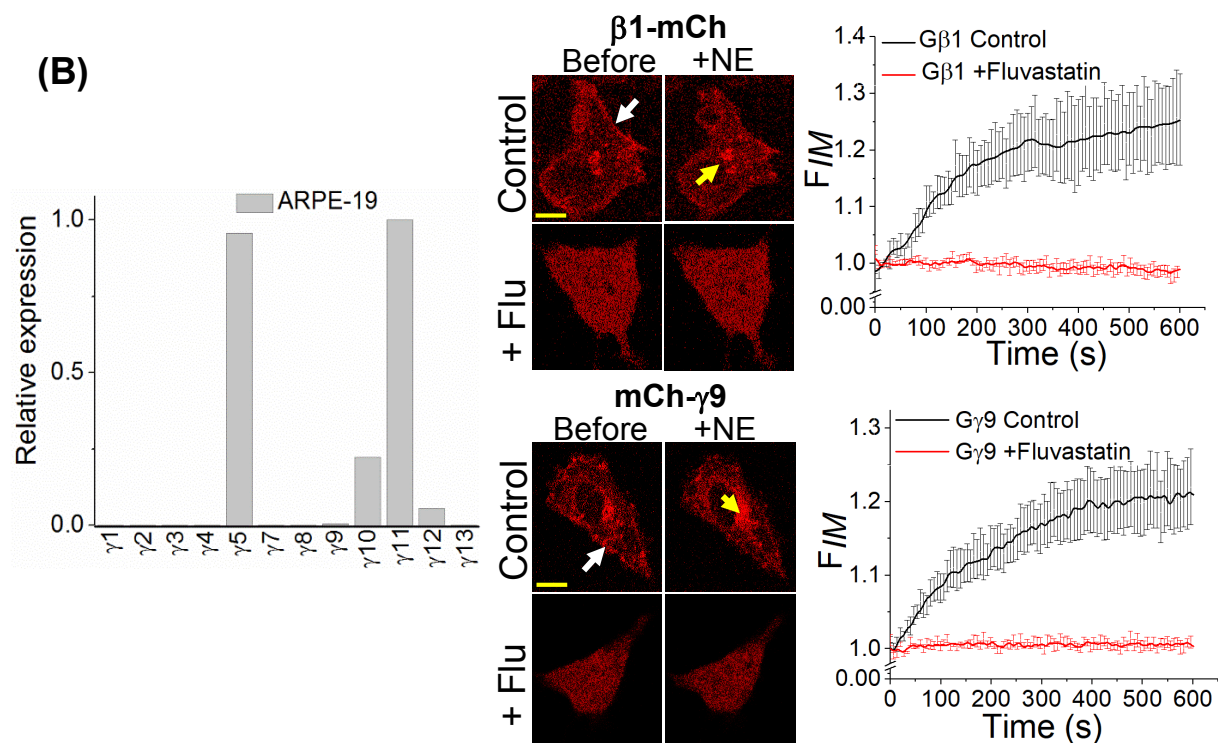


Figure 7

(A)



(B)



Supplemental Data

Statins perturb G $\beta\gamma$ signaling and cell behavior in a G γ subtype dependent manner

Mithila Tennakoon, Dinesh Kankanamge, Kanishka Senarath, Zehra Fasih and Ajith Karunarathne

Molecular Pharmacology

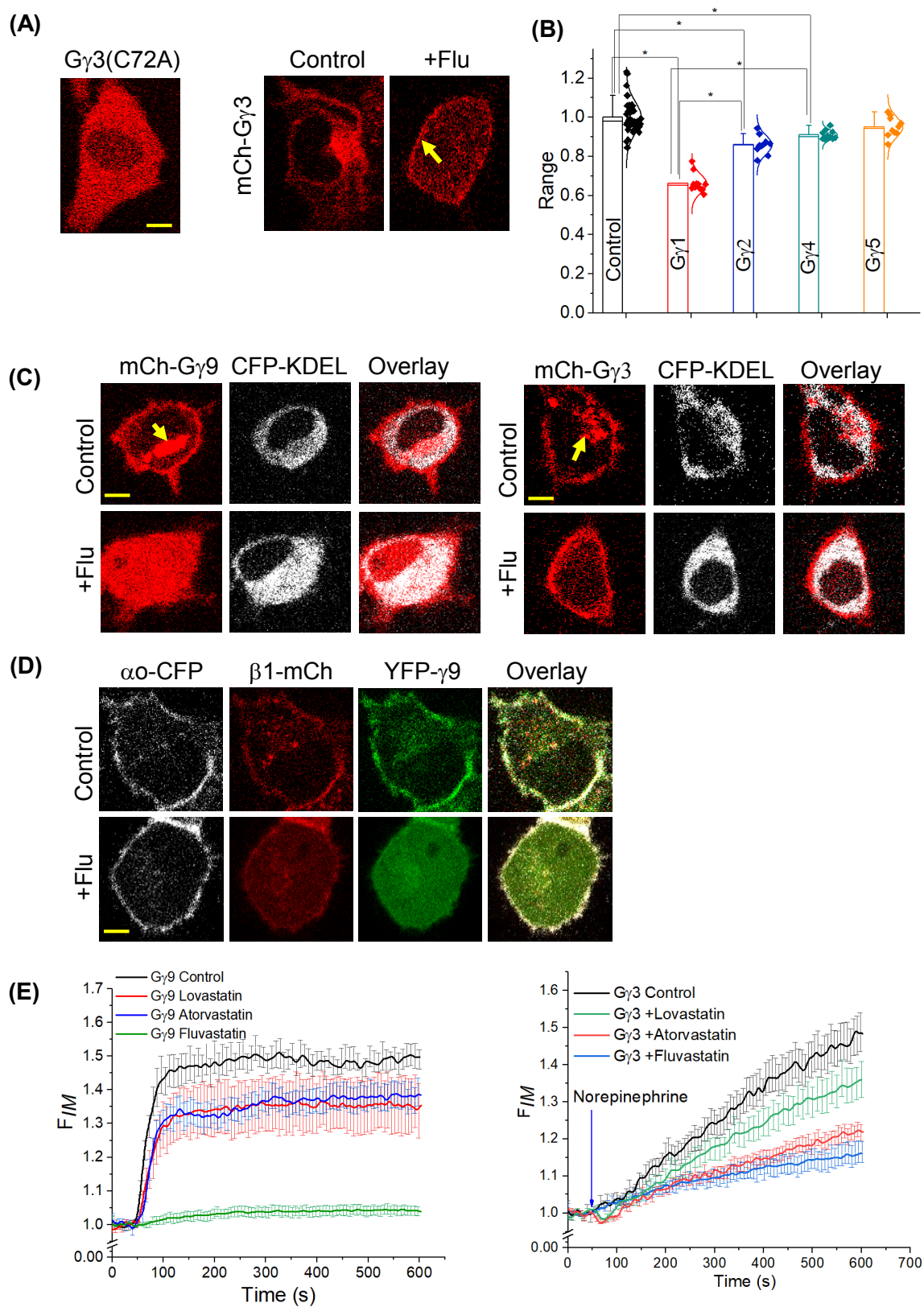
Supplemental Figure Legends

Supplemental Figure 1

- A) **Comparison of cellular distribution of WT mCh-G γ 3 with and without +Flu and mCh-G γ 3 (C72A) mutant.** While WT mCh-G γ 3 expressing control cells showed a strong distribution of G γ 3 on both PM and IMs, +Flu cells having lost their ability to bind to IM exhibited G γ 3 only on the PM (Yellow arrow). Comparatively mCh-G γ 3(C72A) mutant expressing cells showed G γ 3 distribution all over the cytosol, but not bound to either PM or IMs (scale bar: 5 μ m).
- B) **Comparison of the extents of G γ membrane binding inhibition by fluvastatin.** As the numerical value indicates, while farnesylated G γ 1 exhibited the highest inhibition, the other geranylgeranylated G γ s showed comparatively low inhibition. One-way ANOVA was performed, and the data were statistically different $F(4, 77) = 79.19$, $P < 0.05$ (error bars: SD, $n > 10$, $p < 0.05$).
- C) **G γ -type independent inhibition of IM binding by fluvastatin.** HeLa cells expressing CFP-KDEL and either mCh-G γ 9 or G γ 3. Compared to control cells of both G γ 9 and G γ 3 with clear distribution of G γ bound to IMs (yellow arrow), regardless of the G γ subtype, +Flu cells exhibited absence of G γ on the IMs. CFP-KDEL shows the intact IMs even after +Flu treatment to cells (scale bar: 5 μ m).
- D) **Effects of fluvastatin treatment on subcellular distribution of heterotrimeric G protein subunits.** HeLa cells expressing G α -CFP, G β 1-mCh, and G γ 9-YFP. Compared to control cells, PM localization of G α subunit of many +Flu cells remained intact. However, both G β 1 and G γ 9 showed primarily a cytosolic distribution (scale bar: 5 μ m).
- E) **The varying effect of G γ 3 and G γ 9 translocation from PM to IM by different drug.** Plots shows the extent of translocation of G γ 9 (left) and G γ 3 (right) with different drug treatment. Compared to the effect on +Flu cells, in the G γ 9 and G γ 3 expressing cells the effect by +Ator and +Lov is moderate. (error bars: SEM, $n > 12$, $p < 0.05$)

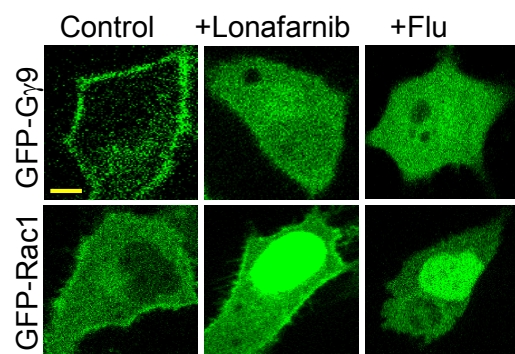
Supplemental Figure 2

- A) **Comparison of prenylation-inhibition by fluvastatin and farnesyl transferase inhibitor, Lonafarnib.** HeLa cells expressing GFP-G γ 9 and GFP-Rac1 exhibited identical and complete inhibitions of PM-binding when exposed to fluvastatin and lonafarnib, indicating that both inhibit farnesylation (scale bar: 5 μ m).
- B) The original images of Western blot analysis showing the expression of p-Akt and β -Actin with and without Fluvastatin. The yellow dashed line indicates the spliced-border where the band splicing was done in the main figure, due to merging of bands.

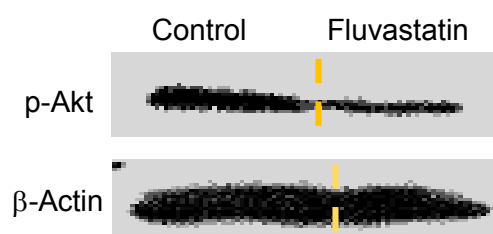


Supplemental Figure 1

(A)



(B)



Supplemental Figure 2