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Conjugated Linoleic Acid Improves Endothelial Ca²⁺ Signaling by Blocking Growth Factor and Cytokine-Mediated Cx43 Phosphorylation

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Abstract

Sustained Ca²⁺ burst signaling is crucial for endothelial vasodilator production and is disrupted by growth factors and cytokines. Conjugated linoleic acid (CLA), a Src inhibitor in certain preparations, is generally regarded as safe during pregnancy by the FDA. Multiple CLA preparations; t10,c12 or c9,t11 CLA, or a 1:1 mixture of the two were administered before growth factor or cytokine treatment. Growth factors and cytokines caused a significant decrease in Ca²⁺ burst numbers in response to ATP stimulation. Both t10,c12 CLA and the 1:1 mixture rescued VEGF₁₆₅ or TNF α inhibited Ca²⁺ bursts and correlated with Src-specific phosphorylation of connexin 43. VEGF₁₆₅, TNF α , and IL-6 in combination at physiologic concentrations revealed IL-6 amplified the inhibitory effects of lower dose of VEGF₁₆₅ and TNF α . Again, the 1:1 CLA mixture was most effective at rescue of function. Therefore, CLA formulations may be a promising treatment for endothelial dysfunction in diseases such as preeclampsia.

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Keywords

Ca²⁺; conjugated linoleic acid; cytokine; growth factor; preeclampsia

1. INTRODUCTION

The vascular endothelium is responsible for sensing blood borne signals and maintenance of homeostasis in the cardiovascular system. Endothelial dysfunction is characterized in part by a reduction in vasodilator production and a shift towards a proinflammatory profile, and can be caused by local and systemic changes in the growth factor and cytokine environment (Visser, van Rijn, Rijkers et al., 2007, Saito, Shiozaki, Nakashima et al., 2007). There are a variety of diseases in which endothelial dysfunction plays a role, including hypertension, atherosclerosis, chronic kidney failure, and preeclampsia (PE) (Rajendran, Rengarajan, Thangavel et al., 2013, Boeldt and Bird, 2017). Despite the widespread acknowledgment of the important role endothelial dysfunction plays in such disorders, therapeutic targeting of the endothelium remains underutilized.

Endothelial intracellular free Ca²⁺ concentration ([Ca²⁺]_i) is an important determinant of vasodilator production (Tran, Leonard, Black et al., 2009). Such elevations in [Ca²⁺]_i often take the form of periodic Ca²⁺ bursts, which correlate directly with vasodilator production (Yi, Boeldt, Gifford et al., 2010, Yi, Magness and Bird, 2005, Krupp, Boeldt, Yi et al., 2013). Ca²⁺ bursting is dependent on functionally coupled Cx43 gap junctions (Yi et al., 2010, Boeldt, Grummer, Yi et al., 2015, Boeldt, Krupp, Yi et al., 2017). Cx43 gap junctions facilitate communication between adjacent cells through the passage of small molecules and ions, such as IP₃ and likely Ca²⁺ (Bird, Boeldt, Krupp et al., 2013, Solan and Lampe, 2009). The function of Cx43 gap junctions is highly influenced by growth factor and cytokine mediated kinase activation. Src activation results in direct phosphorylation of Cx43 at the inhibitory Y265 residue (Solan and Lampe, 2014). Phosphorylation of Y265 promotes downregulation of Cx43-mediated communication, most likely by destabilizing the Cx43 plaques and increasing the rate of protein turnover (Solan and Lampe, 2014). Because sustained endothelial Ca²⁺ signaling depends on functionally coupled Cx43 gap junctions, the net result of increased Y265 phosphorylation is reduced Ca²⁺ signaling capacity (Boeldt et al., 2015, Ampey, Boeldt, Clemente et al., 2019).

Previous studies have shown that exposure of intact endothelium to 10ng/ml VEGF₁₆₅ reduces ATP-stimulated Ca²⁺/nitric oxide (NO) production to a level that is not compensated for by the Ca²⁺/NO production attributable to VEGF₁₆₅ itself (Boeldt et al., 2017, Yi, Boeldt, Magness et al., 2011). Many of the growth factors and cytokines that may mediate endothelial dysfunction can be sorted into groups/classes based on how their receptors couple directly (VEGF₁₆₅, bFGF, EGF, TNF α) (D'Angelo, Struman, Martial et al., 1995, Deo, Axelrad, Robert et al., 2002, Eliceiri, Paul, Schwartzberg et al., 1999, Huang, Dudez, Scerri et al., 2003, Nwariaku, Liu, Zhu et al., 2002, Parenti, Morbidelli, Cui et al., 1998, Tang, Yang, Chen et al., 2007, Waltenberger, Claesson-Welsh, Siegbahn et al., 1994) or indirectly (IL-6, IL-8, IL-1 β) (Huang, Yang, Hsu et al., 2013, Liu, Kuo, Wang et al., 2016) to the same common post receptor kinase pathways. However, the literature has widely failed

to address the effects of these growth factors and cytokines when used in combination, as they exist in pathophysiologic context, instead of individually. Therefore, there is a need to construct “cocktails” utilizing multiple growth factors and cytokines to bridge the gap from single agonist experiments to experimental conditions that are more representative of the in-vivo environment of endothelial dysfunction. Co-stimulation with multiple growth factors and cytokines may be dependent on an unappreciated signaling hierarchy, and because these kinases may present a common pathway for endocrine-mediated endothelial dysfunction, they may be a prime target for therapeutic intervention (Bird et al., 2013).

In this study, we use human umbilical vein endothelial cells (HUVECs) to model the combined actions of these growth factors and cytokines on endothelial cell function. We use PE as a model system to construct our cocktails as it has been described as a condition of cytokine-mediated endothelial dysfunction (Redman, Sacks and Sargent, 1999). As appropriate levels of growth factors and cytokines are necessary to support normal pregnancy, we also use a cocktail representative of normal pregnancy as a control. For this study we include TNF α , IL-6 and VEGF in our cocktails, using doses attributed to ‘normal’ and PE pregnancies for TNF α and IL-6 (Lau, Guild, Barrett et al., 2013, Tosun, Celik, Avci et al., 2010, Xie, Yao, Liu et al., 2011), and varied doses of VEGF₁₆₅ to mimic systemic and local concentrations. While PE is used as the model for which we construct our dysfunctional phenotype, the resulting findings will likely apply to a broad range of diseases that have an endothelial dysfunction component.

The pharmacological Src inhibitor PP2 has already proven successful in rescuing VEGF₁₆₅ and TNF α pretreated endothelium from a dysfunctional Ca²⁺ signaling phenotype to one more congruent with normal function (Boeldt et al., 2015, Ampey et al., 2019). A pharmacological tool like PP2 is not used in humans due to significant toxicity and concerns of teratogenicity in pregnancy, and therefore offers no therapeutic value. A pair of isomers of conjugated linoleic acid (CLA) used at FDA recommended doses for treatment of inflammatory conditions including rheumatoid arthritis, Crohn’s disease and hypertension may hold promise as a safe method for Src inhibition, even in pregnancy. The t10,c12 CLA isomer has been shown to be a Src inhibitor in cancer cells (Shahzad, Felder, Ludwig et al., 2018). When used in ovine endothelial cells, VEGF₁₆₅-mediated inhibition of Ca²⁺ responses were rescued with as little as 5 μ m t10,c12 CLA, and 50 μ M t10,c12 CLA rescued TNF α -mediated Ca²⁺ response inhibition (Boeldt et al., 2015, Ampey et al., 2019). Herein, we examine the efficacy of individual c9,t11 and t10,c12 isomers, as well as a 1:1 mix to rescue Cx43-dependent Ca²⁺ signaling in HUVEC pretreated with individual growth factors or cytokines. We also examine the effect of combinations of a selection of growth factors and cytokines on Ca²⁺ signaling and the ability of CLA isomers alone or as a 1:1 mix, to restore function to the normal phenotype.

2. MATERIALS AND METHODS

2.1 Materials

ATP (disodium salt), Heparin sodium salt and all other chemicals were purchased from Sigma- Aldrich (St Louis, MO, USA) unless otherwise stated. Growth factors and cytokines (*bFGF*, *EGF*, VEGF₁₆₅, TNF α , IL-1 β , IL-6, IL-8) were purchased from R & D systems

(Minneapolis, MN, USA). Glass bottom microwell dishes for Ca^{2+} imaging studies were from MatTek Corporation (Ashland, MA, USA). Minimum Essential Medium (MEM) was purchased from Invitrogen (Life Technologies Inc., Grand Island, NY). Serum used in culture medium was fetal bovine serum (FBS) from Invitrogen and Endothelial Cell Growth Supplement ECGS was from Millipore (Temecula, CA). Experimental buffer was Krebs buffer (125mM NaCl, 5 mM KCl, 1 mM MgSO_4 , 1 Mm KH_2PO_4 , 6 mM glucose, 25 mM HEPES, 2 mM CaCl_2 , pH 7.4). The growth medium is referred in this article as HEH medium (Human, ECGS, Heparin) and is as described in (Krupp et al., 2013). PP2 is from Sigma- Aldrich (St. Louis, MO, USA), and t10,c12 CLA and c9,t11 CLA are from Matreya (Pleasant Gap, PA, USA).

2.2 Primary Cell Culture

Institutional review board (IRB) was approved from University of Wisconsin School of medicine and public health. HUVEC were isolated from fresh umbilical cords after informed consent from patients that delivered via elective Cesarean section with intact membranes and no evidence of labor or infection. Cords obtained were put into cold (4 C) PBS solution and were ready for cell harvest within 30 minutes from the delivery of the placenta. Complete methods of harvesting and storing of HUVECs has been described elsewhere in detail (Krupp et al., 2013). HUVECs were pooled from 5 subjects at passage 3 and then used for experimentation and data analysis. Patient information can be found in Krupp et al (Krupp et al., 2013).

2.3 Cell growth and Ca^{2+} imaging

HUVECs (passage 4) were grown to confluence in 35 mm, glass-bottom microwell dishes, and handled as described in (Krupp et al., 2013,Boeldt et al., 2015,Boeldt et al., 2017). HUVECs were incubated with 10uM Fura2-AM (Molecular Probes) with 0.05% Pluronic acid F127 (Life Technologies) in HEH for 1 hour at 37°C. HUVECs were washed with Krebs buffer following incubation and HUVECs were left in Krebs buffer for 30 minutes at room temperature, allowing for complete ester hydrolysis. The dish was then placed on a Nikon inverted microscope (Diaphot 150; Nikon, Melville, NY) for imaging. Up to 99 cells per field of view were identified and manually circled for analysis (Krupp et al., 2013). After 5 minutes baseline recording, HUVECs were stimulated by 100 uM ATP (ATP exposure 1) and recorded for a total of 30 minutes using alternate excitation at 340 nm and 380 nm at 1 second intervals. Number of cells that had 3 bursts were noted and calculated. Cells were then washed three times with Krebs' buffer for 20 minutes to bring cells to baseline for further imaging steps (A and B below). A standard curve was used to calculate concentration of intracellular Ca^{2+} ($[\text{Ca}^{2+}]_i$).

- a. For the effect of growth factors and cytokines on ATP-stimulated Ca^{2+} bursts, cells were exposed to 10ng/ml VEGF₁₆₅, bFGF, EGF, TNF α , IL-1 β , IL-6, IL-8, or combinations thereof for 30 minutes (except TNF α – 60 minutes) and 100uM ATP was subsequently added for an additional 30 minutes (ATP exposure 2). The number of Ca^{2+} bursts during the second ATP exposure were noted and compared to the number of Ca^{2+} bursts from the initial exposure.

- b. For recovery of ATP-stimulated Ca^{2+} bursts with CLA, 10 or 50 μM t10,c12 CLA or c9,t11 CLA (30 minutes) were added to the dish during the wash step. This was followed by treatment with growth factor or cytokine for the required time, and then ATP as in step A.

2.4 Western Blot Analysis

HUVEC were grown to 100% confluence on 60mm dishes before 4-hour serum withdrawal. Cells were then pretreated with t12,c12 CLA (50 μM), c9,t11 CLA (50 μM), or a 1:1 Mix of both isomers (10 μM) before 10ng/ml TNF α (1 hour) or 10ng/ml VEGF₁₆₅ (30minutes) application. Media was quickly aspirated and cells were snap frozen in liquid nitrogen, harvested, lysed, and loaded on 7.5% gels for SDS-PAGE at 20ug protein/well exactly as previously published (Boeldt et al., 2015, Grummer, Sullivan, Magness et al., 2009, Sullivan, Grummer, Yi et al., 2006). Phosphorylation of Cx43 was detected using an antibody to Tyr-265 Cx43 (#sc-17220-R; Santa Cruz Biotechnology) at 1:667. Secondary antibody was goat anti-rabbit HRP-conjugated (#7074, Cell Signaling Technology) at 1:3000. Loading control was Hsp90 (#4874, Cell Signaling Technology) at 1:2500.

2.5 Statistical analysis

For imaging, each agonist was administered in 4–6 dishes with each dish marking 90–95 cells. Burst numbers in each cell were counted, the difference from internal control was taken, and this difference is what was used for data analysis and creation of figures. Mann-Whitney rank sum test was used for statistical analysis. For Western Blot Analysis, a minimum of 6 independent experiments were run for each agonist, and statistical analysis was performed by post-hoc Tukey HSD one-way ANOVA with Holm interference. Data are presented as means \pm S.E.M. and $P < 0.05$ was considered statistically significant. Linear regression analyses were presented with 95% confidence intervals.

3. RESULTS

3.1 Effect of Individual Growth Factors and Cytokines on Ca^{2+} Bursts and CLA Rescue

Representative single cell tracings from Fura-2 loaded HUVECs are shown in Figure 1. A short baseline precedes 100 μM ATP stimulation at 0 seconds. Upon stimulation, an immediate initial Ca^{2+} peak is seen followed by multiple transient Ca^{2+} bursts over the duration of the 30-minute experiment. Figures 1A–F shows individual ATP-stimulated cells before and after a pretreatment with Krebs buffer control (top panels A, B), 10ng/ml VEGF₁₆₅ (middle panels C, D), or 10ng/ml TNF α (bottom panels E, F). Cells were selected as representative of the mean number of Ca^{2+} bursts observed for each condition. A reduction in Ca^{2+} burst numbers was observed for the cells treated with VEGF₁₆₅ and TNF α , but not for control cells.

Figure 2 shows the quantification of the loss of ATP-stimulated Ca^{2+} bursts in HUVECs with growth factor or cytokine treatments (ATP control - white bars; growth factor or cytokine treatment - black bars). These treatments were compared with control (defined as HUVEC cells stimulated with 100 μM ATP followed by Krebs wash and ATP stimulation again to rule out intrinsic Ca^{2+} burst loss). The physiologic functional range, as defined

previously in uterine artery endothelium and calibrated to the HUVEC model here by TPA treatment (Boeldt et al., 2017, Bird et al., 2013) is highlighted by the shaded region. In the absence of a non-pregnant control for HUVEC studies, we have previously established that the use of 1nM TPA produces a 60% reduction in Ca^{2+} bursting that parallels results from the non-pregnant ovine model, serving as a value that can represent pregnancy adapted function (Boeldt et al., 2017, Bird et al., 2013). This emphasizes that a relatively small difference in Ca^{2+} burst numbers of ~40% can span the entire range of pregnancy-adapted function, and can likely be applied equally to other vascular beds. While cells show significant ($p < 0.05$) loss of Ca^{2+} bursts with all growth factors or cytokines that were used, VEGF₁₆₅ (panel A) and TNF α (panel D) were the most potent Ca^{2+} burst inhibitors (55% and 70% of control bursting respectively, $p < 0.001$). Other treatments also showed a loss of Ca^{2+} bursts, with the smallest loss observed (e.g. EGF (panel C), IL-6 (panel F), IL-8 (panel G)) at 80–90% of control Ca^{2+} bursts, still achieving statistical significance.

HUVEC were also pretreated with two common CLA isomers prior to growth factor or cytokine treatment and subsequent 30-minute 100uM ATP stimulation (Figure 2). In initial studies, cells were exposed to 50uM of either c9,t11 CLA (non-Src inhibiting isomer (Shahzad et al., 2018)) or t10,c12 CLA (Src inhibiting isomer (Shahzad et al., 2018)) in order to assess the effect each may have individually on growth factor or cytokine-mediated inhibition of ATP-stimulated Ca^{2+} bursts. Pretreatment with c9,t11 CLA showed no capability in improving ATP-stimulated Ca^{2+} burst responses during treatment with any of the seven growth factors or cytokines used in this study. In the case of EGF (panel C), treatment with c9,t11 CLA exacerbated the growth factor or cytokine-mediated inhibition of Ca^{2+} bursts. However, when treated with the Src inhibiting t10,c12 CLA isomer, improvement in ATP-stimulated Ca^{2+} bursts was observed for VEGF₁₆₅ (panel A) and TNF α (panel D). Only in the case of IL-1 β (panel E) was any further reduction in ATP-stimulated Ca^{2+} bursts observed. Cells exposed to 50uM c9,t11 CLA with no growth factor or cytokine produced 80.5% ($\pm 4.4\%$) as many Ca^{2+} bursts compared to internal ATP control. Likewise, treatment with 50uM t10,c12 CLA alone produced 88.9% ($\pm 4.6\%$) of internal control Ca^{2+} bursts (also see Figure 7).

Because commercially available and FDA-approved formulations of CLA specify mixtures of CLA isomers, we included in our studies a 1:1 mix of c9,t11 and t10,c12 CLA as a Ca^{2+} burst recovery agent. Representative single-cell tracings for rescued Ca^{2+} bursting function for 10uM CLA Mix pretreatment with VEGF₁₆₅ and TNF α are shown in Figure 3, and quantification is shown in Figure 4. After a 30 minute pretreatment with a mixture of 50uM c9,t11 CLA and 50uM t10,c12 CLA, no positive effect on growth factor or cytokine inhibited Ca^{2+} bursts was observed. Significantly impaired Ca^{2+} burst responses were observed in response to EGF (panel 4C), IL-6 (panel 4F), and IL-8 (panel 4G). When the doses of c9,t11 and t10,c12 CLA were dropped to 10uM each, the 30-minute pretreatment was more effective in yielding a significant improvement for VEGF₁₆₅ (panel 4A), bFGF (panel 4B), and TNF α (panel 4C). The only exception was that further impairment was observed in the presence of IL-8 (panel 4G). Cells exposed to 50uM CLA mix with no growth factor or cytokine produced 68.9% ($\pm 5.2\%$) as many Ca^{2+} bursts compared to internal ATP control. Ca^{2+} Treatment with 10uM CLA mix alone produced 91.0% ($\pm 6.5\%$) of internal control Ca^{2+} bursts (also see Figure 7).

To better understand the relationship between degree of inhibition and degree of CLA effect or “rescue”, we applied two regression analyses (Figure 5) to each treatment group from Figures 2 and 4. When all growth factors and cytokines are pooled together (left panels A, C, E, G), there is a positive relationship between degree of inhibition and degree of rescue (i.e. those growth factors and cytokines which are the most inhibitory are the most rescued by CLA treatments) for each CLA treatment group. However, when we separated the interleukins that are not coupled directly to Src from the non-interleukin growth factors and cytokines that do couple directly to Src, another trend was apparent (panels B, D, F, H). For those CLA treatments which were mostly ineffective (50uM c9,t11 CLA (panel B) and 50uM 1:1 CLA mix (panel F), there was no clear trend in how inhibition correlated with rescue. But for the more effective treatments (50uM t10,c12 CLA (panel D) and 10uM 1:1 CLA Mix (panel H) the growth factors VEGF₁₆₅, bFGF as well as TNF α showed a positive relationship between inhibition and rescue, while interleukins showed a weaker negative relationship between inhibition and rescue.

In order to assess the effect of t10,c12 CLA on Src-mediated phosphorylation of Cx43, we performed phospho-specific western blots for HUVEC stimulated with VEGF₁₆₅ or TNF α . In Figure 6A and 6B, representative blots at the 43kD band are shown along with Hsp90 loading control. It should be noted that Y265 antibody binds rather weakly and as a result often gives blots that look quite variable before normalization and background correction. In Figure 6C, quantification of phosphorylation of the Src-sensitive Y265 site on Cx43 was increased by VEGF₁₆₅, TNF α , or 50uM c9,t11 CLA treatment. Phosphorylation of Y265 by TNF α was significantly reversed by 50uM t10,c12 CLA. Further validation of Src specificity of the Y265 site on Cx43 is shown by PP2 inhibition of phospho-Y265 under control and TNF α -stimulated conditions (Supplementary Figure 1). In Figure 7, we further analyzed the Y265 phosphorylation data by regression analysis of Ca²⁺ burst numbers versus Y265 phosphorylation. This plot showed a significant correlation between Y265 phosphorylation and ATP-stimulated Ca²⁺ burst responses when both VEGF₁₆₅ and TNF α data were pooled, indicating that Src-mediated Cx43 phosphorylation at Y265 is related to decreased Ca²⁺ burst ‘ numbers and is sensitive to CLA treatment.

3.2 Effect of Growth Factor and Cytokine Cocktails on Ca²⁺ Bursts and CLA Rescue

In order to model more physiologically relevant combinations of growth factors and cytokines in normal pregnancy and PE, we exposed HUVEC to cocktails (Normal or PE) of VEGF₁₆₅ and TNF α , with or without IL-6 for 30 minutes. Figure 8 illustrates representative single-cell tracings of the reduction in Ca²⁺ bursting due to VEGF₁₆₅, TNF α , and IL-6 cocktail exposure and the improvements achieved with 10uM CLA Mix. In Figure 9, we quantify the change in Ca²⁺ burst responses for the VEGF₁₆₅ and TNF α only cocktails. Doses corresponding to normal pregnancy were 0.5ng/ml VEGF₁₆₅ and 0.03ng/ml TNF α . For modeling PE pregnancies, we created three cocktails with varying concentrations of VEGF₁₆₅. Concentrations for (L)PE V+T were 0.1ng/ml VEGF₁₆₅ and 0.5ng/ml TNF α , for (M)PE V+T were 1 ng/ml VEGF₁₆₅ and 0.5ng/ml TNF α , and for (H)PE V+T were 10ng/ml VEGF₁₆₅ and 0.5ng/ml TNF α . The Normal V+T cocktail was moderately inhibitory and the 10uM CLA mix further inhibited the ATP-stimulated Ca²⁺ bursts. The PE V+T cocktails all inhibited ATP-stimulated Ca²⁺ bursts, in a VEGF₁₆₅ dose dependent manner. Pretreatment

with the 10uM CLA mix for 30 minutes exacerbated the (L)PE V+T mediated Ca^{2+} burst inhibition, but significantly improved ATP-stimulated Ca^{2+} bursts after (M)PE V+T and (H)PE V+T treatment.

In Figure 10, we added IL-6 to the V+T cocktails from Figure 9 and again pretreated for 30 minutes. The Normal cocktail thus contained 0.5ng/ml VEGF₁₆₅, 0.03ng/ml TNF α , and 0.1ng/ml IL-6. The (L)PE cocktail contained 0.1ng/ml VEGF₁₆₅, 0.5ng/ml TNF α , and 3ng/ml IL-6. The (M)PE cocktail contained 1ng/ml VEGF₁₆₅, 0.5ng/ml TNF α , and 3ng/ml IL-6. Finally, the (H)PE cocktail contained 10ng/ml VEGF₁₆₅, 0.5ng/ml TNF α , and 3ng/ml IL-6. The Normal cocktail had no significant effect on ATP-stimulated Ca^{2+} bursts, but the addition of the 10uM CLA Mix had a small inhibitory effect. All three PE cocktails were quite inhibitory to ATP-stimulated Ca^{2+} bursts, and only in the presence of the (H)PE cocktail was the 10uM CLA Mix more effective in improving Ca^{2+} bursts.

4. Discussion

The first goal of this study was to determine if individual growth factors or cytokines might induce endothelial dysfunction through inhibition of Ca^{2+} signaling mechanisms. Initially, we used a standard dose of 10ng/ml for all growth factors and cytokines, which was based on previous published and unpublished studies from ovine and HUVEC models in which effects were related to Cx43 inhibitory phosphorylations via Src kinase activity (Boeldt et al., 2015, Khurshid, 2015). In this study, we clearly observed a significant loss of Ca^{2+} bursts in response to growth factors and cytokines. Therefore, all agents investigated are capable of negatively influencing Ca^{2+} burst responses. The mechanism of signaling action for the most inhibitory factors alone (VEGF₁₆₅, TNF α , and bFGF) are all capable of promoting phosphorylation of Cx43 gap junction proteins because they all couple directly to Src (Boeldt et al., 2015, Bird et al., 2013, Vitale and Barry, 2015). This was demonstrated in detail with VEGF₁₆₅ and TNF α in the ovine uterine artery and human umbilical vein models (Boeldt et al., 2015, Boeldt et al., 2017, Ampey et al., 2019). Such a proposal is consistent with the findings of others who have suggested VEGF₁₆₅, EGF, bFGF, TNF α (directly) or IL-6, and IL-8 (indirectly) activate Src kinase to varying degrees (Deo et al., 2002, Eliceiri et al., 1999, Nwariaku et al., 2002, Tang et al., 2007, Huang et al., 2013, Liu et al., 2016), and is supported by Y265 phospho-Cx43 data herein. We were then able to assess the ability of potential nutraceutical therapeutic agent (t10,c12 CLA alone or mixed 1:1 with c9,t11 CLA) to inhibit Src kinase and rescue endothelial function. Because c9,t11 CLA is not known to block Src activation, it was included as a negative control. Indeed, c9,t11 CLA was unable to improve Ca^{2+} signaling after treatment with any of the growth factors or cytokines used. Since t10,c12 CLA has been shown to inhibit c-Src in cancer cells (Shahzad et al., 2018), we hypothesized that treatment with any CLA mixture that includes this isomer would rescue Ca^{2+} signaling from insult mediated by Src activation, whether directly or indirectly. Significant improvements were observed for VEGF₁₆₅, TNF α , and bFGF. Western blot data for Y265 phospho-Cx43 suggested that the t10,c12 CLA isomer does mediate Cx43 function through Src signaling, and as a pretreatment for TNF α , restores Y265 phosphorylation to control levels. The growth factors and cytokines which are known to couple directly to Src (i.e. the non-interleukins VEGF₁₆₅, EGF, bFGF, and TNF α) had a positive correlation between degree of inhibition and degree of rescue. Of note, EGF is a particularly weak

inhibitor and it is noteworthy that previous studies in HUVEC have shown EGF receptor numbers are very low in HUVEC (Al-Nedawi, Meehan, Kerbel et al., 2009). Those interleukin receptors that couple signaling to Src *indirectly* (i.e. receptors to IL-8, IL-6, and IL-1 β) all displayed a negative correlation between degree of inhibition and degree of rescue.

While the phospho-Cx43 blockade by t10,c12 CLA is clearer for TNF α than VEGF₁₆₅, it should be noted that only a portion of all Cx43 resides in functional gap junctional plaques, so it is a considerable challenge to detect significant changes in Y265 phosphorylation that mimic functional outcomes, such as Ca²⁺ burst responses. Indeed VEGF₁₆₅ failed to achieve the statistically significant Y265 Cx43 phosphorylation by western blot clearly observed for TNF α . However, when the VEGF₁₆₅ data for c9,t11 CLA, t10,c12 CLA, and the 1:1 mix of isomers are added to the equivalent data for TNF α and plotted against Ca²⁺ burst data, it is very clear that a significant negative correlation between Cx43 phosphorylation at Y265 and loss of Ca²⁺ bursts exists. This supports previous work in which we have shown that GAP peptide-mediated inhibition of Cx43 function results in a loss of Ca²⁺ bursting function as well (Yi et al., 2010). That the c9,t11 isomer results in significant phosphorylation of Y265 on its own, is an interesting result that warrants further study.

The Src specificity of Y265 on Cx43 has been demonstrated clearly in numerous studies, and has been clearly described in the context of cell function in seminal reviews by Lampe and Lau (Lampe and Lau, 2000, Lampe and Lau, 2004). Initial studies linked Src only to tyrosine phosphorylation on Cx43 (Loo, Berestecky, Kanemitsu et al., 1995, Loo, Kanemitsu and Lau, 1999), but other studies demonstrated that Y247 and Y265 were Src substrates (Kanemitsu, Loo, Simon et al., 1997, Giepmans, Hengeveld, Postma et al., 2001, Lin, Warn-Cramer, Kurata et al., 2001, Lin, Warn-Cramer, Kurata et al., 2001). However, Y247 phosphorylation appears to be dependent on Y265 phosphorylation (Lin et al., 2001) and therefore Y265 likely presents a better in vitro assay target for Src activity at Cx43. In the ovine uterine artery endothelial cell (UAEC) model, PP2 specifically inhibits Y265 pCx43 and not the MEK/ERK specific Ser279/282 pCx43 after VEGF₁₆₅ or TNF α stimulation, with TPA as a positive control (Boeldt et al., 2015, Ampey et al., 2019). Those studies also linked Cx43 phosphorylation with decreased sustained Ca²⁺ bursting in the context of pregnancy adapted endothelial function.

While such previous work in the ovine UAEC model demonstrated strongly the connection between Cx43 gap junction function and Ca²⁺ bursting capacity (Yi et al., 2010, Boeldt et al., 2015), the HUVEC model does not mirror those results exactly. The previous studies in UAEC and HUVEC also stopped short of directly implicating reversal of Cx43 phosphorylation as a mechanism by which CLA may be rescuing Ca²⁺ burst function after growth factor or cytokine pretreatment. In HUVEC, inhibition of gap junction function with GAP peptides revealed that while Cx43 was important for Ca²⁺ bursting function, there is likely also a role for Cx37 and Cx40 potentially via heteromeric or heterotypic gap junctions. Cx37 expression has been found to be increased in pregnancy in ovine uterine arteries (Morschauser, Ramadoss, Koch et al., 2014) though it does not appear to have a direct role in Ca²⁺ signaling. Thus, a potential role for Cx37 is more likely than Cx40, but still remains unclear at this time. These observed differences may be species specific or

could be due to the vasculature bed from which the endothelial cells are derived, and may contribute to the difficulty of clean Cx43 Y265 phosphorylation western blots, as the gap junctions may be comprised of a heterogeneous mixture of Cx isoforms (Boeldt et al., 2017).

In general, CLA may be most effective as a therapeutic rescue agent in those conditions that are most inhibitory to Ca^{2+} signaling, and particularly via Src and ERK. But in those conditions that are only modestly inhibitory, CLA may be a poor candidate for therapy. This fits with clinical studies where CLA has little effect on blood pressure in healthy subjects (Yang, Wang, Zhou et al., 2015) but can improve blood pressure in hypertensive subjects (Sato, Shinohara, Honma et al., 2011). Estimates of the dose of CLA the patients from these studies received would be around 10uM, also suggesting that our 10uM 1:1 mix dose could be therapeutically relevant. In a cohort at risk for complicated pregnancies (women age 18 and younger) treatment with calcium (addressing a nutritional deficit) and CLA resulted in a decreased incidence of PE. The decreased PE incidence was not observed in the group receiving only calcium supplementation, implicating a role for CLA in prevention of PE in human pregnancies (Alzate, Herrera-Medina and Pineda, 2015). In this study, the coupling of Cx43 phosphorylation with the Ca^{2+} data implicates a mechanistic role for CLA in protecting pregnancy-adapted Ca^{2+} signaling via blocking Src-mediated gap junction functional inhibition that may have been unappreciated in the Colombian trial. In further support of the potential role for CLA to support gap junction communication, Rakib et al have reported that CLA has been found to prevent TPA-mediated phosphorylation of Cx43 gap junctions (Rakib, Kim, Jang et al., 2010). Taken together, this work indicates that CLA treatment could potentially be beneficial for hypertensive diseases but warrants further mechanistic work.

Another goal of this study was to measure the effect of a cocktail of growth factors and cytokines on endothelial Ca^{2+} signaling. This was designed as the first step to more closely mimic the in vivo signaling environment of endothelial dysfunction, specifically that of PE. These combinations in turn may trigger combinations of signaling responses that may be additive or even synergistic on Cx43 inhibition. This is all the more relevant when one considers that interleukins such as IL-6 can possibly amplify Src activation by agents such as $\text{TNF}\alpha$ through GP130 signaling (Schaper and Rose-John, 2015). We used published data on levels of growth factors and cytokines in preeclampsia to guide our cocktail creation (Lau et al., 2013, Tosun et al., 2010, Xie et al., 2011). We used a dose response of VEGF_{165} in our cocktails as there is debate in the field of preeclampsia research on VEGF_{165} levels in the disease state. We rounded out our 'cocktails' by first adding in $\text{TNF}\alpha$ (0.5ng/ml) initially (Figure 9) and then further adding IL-6 (3ng/ml) in Figure 10. As a negative control, we also created cocktails which reflect the levels of VEGF_{165} , $\text{TNF}\alpha$, and IL-6 in a non-pathologic state.

When taken together, the V+T and V+T+IL-6 cocktails revealed some important points. For one, combination treatments seemed to magnify the degree of inhibition compared to any individual growth factor or cytokine, especially given that the doses for all but the high VEGF_{165} were well below the 10ng/ml doses from the individual factor experiments. Both high VEGF_{165} combinations dropped Ca^{2+} signaling well out of the normal physiological range. The 1:1 Mix CLA treatment was able to restore function to the physiological range,

however, considerable improvement capacity remains. In the absence of any IL-6, VEGF₁₆₅ mediated Ca²⁺ signaling in a dose dependent manner, with TNF α taking a secondary role in modulating the response amplitude. However, when IL-6 was also present, it had a profound impact. In the Normal cocktail, the small amount of IL-6 added improved the inhibitory Normal V+T response to one indistinguishable from control. When IL-6 was increased, it made the Low V+T cocktail profoundly more inhibitory than in the absence of IL-6. As VEGF₁₆₅ concentrations increased into the medium and high levels, it again reclaimed the dominant role. As with the individual growth factors and cytokines, it appears that the degree of rescue is related to the degree of Ca²⁺ burst inhibition.

It is also worth noting that in some cases cytokines and growth factors can stimulate Ca²⁺ and NO responses in their own right. Numerous studies report on the vasodilator properties of VEGF₁₆₅, and our own studies in UAEC and HUVEC confirm this (Boeldt et al., 2017, Yi et al., 2011). However, VEGF₁₆₅ and other growth factors and cytokines are relatively weak vasodilators and when compared to the concomitant inhibitory effects of better vasodilators such as ATP or Bradykinin, which result in a net deficit of vasodilator production (Reviewed in (Bird et al., 2013, Boeldt, Yi and Bird, 2011, Boeldt, Hankes, Alvarez et al., 2014)). It is therefore important to consider the role of such peptides on overall vasodilator production, and in inflammatory conditions such as PE where TNF α , IL-6, and in certain cases VEGF₁₆₅ are elevated, overall endothelial vasodilator production capacity is diminished. It is for this reason that selectively targeting the signaling pathways downstream of VEGF₁₆₅, TNF α , and IL-6 that negatively impact upon endothelial Ca²⁺ signaling may present a novel therapeutic approach. That CLA formulations may provide some benefit to endothelial Ca²⁺ signaling by blocking Y265 phosphorylation in Cx43 is an important consideration.

This study did not address the potential role of Ca²⁺ permeable channels contributing to sustained phase Ca²⁺ signaling. It is plausible that such channels may play a role in the remaining Ca²⁺ signaling function untouched by CLA treatment. TRP and ORAI channels are key regulators of extracellular Ca²⁺ influx (Silva and Ballejo, 2019) in endothelial cells. Traditionally, canonical transient receptor potential (TRP) channels have been thought to have an integral role in endothelial Ca²⁺ signaling. Specifically TRPC4 has been found to be expressed in endothelial cells and to contribute to Ca²⁺ homeostasis (Graziani, Poteser, Heupel et al., 2010), and TRPC1, TRPC3, and TRPC6 have been implicated in specific endothelial cell types (Gifford, Yi and Bird, 2006, Ge, Tai, Sun et al., 2009, Groschner, Hingel, Lintschinger et al., 1998, Cheng, James, Foster et al., 2006). ORAI channels are multimeric channels comprised of ORAI family member proteins, of which ORAI1 is best characterized, and are activated by Stim 1 and 2 (Bergmeier, Weidinger, Zee et al., 2013). Abdullaev et al have reported that in HUVECs downregulation of Stim1 or Orai1 suppressed store-operated Ca²⁺ entry (SOCE) and ectopic expression eYFP-Stim1 or CFP-Orai1 was able to rescue SOCE. They also reported that knockdown of TRPC1 or TRPC4 did not inhibit SOCE (Abdullaev, Bisailon, Potier et al., 2008). There are additional reports that TRPC channels may be involved in maintenance of ORAI channel components long term (Abdullaev et al., 2008), that there is cross-talk between ORAI and TRP channels (Graziani et al., 2010) and that both ORAI and TRP channels are involved in Ca²⁺ influx within the first five minutes after stimulant-mediated ER store release (Silva and Ballejo, 2019). Elucidating the role of these channels further in HUVECs may provide insight into the

strategy to achieve full rescue of growth factor and cytokine mediated inhibition of Ca^{2+} signaling.

In conclusion, our data suggest that several growth factors and cytokines alone or in combination may act additively or synergistically to cause Src-mediated loss of Ca^{2+} bursting via inhibition of Cx43 gap junctions. Prolonged exposure of such hormone combinations via Src signaling would certainly promote sustained phosphorylation of Cx43 gap junction protein at inhibitory sites, possibly resulting in disassembly of the membrane structural protein arrays in which Cx43 resides. The result of such Cx43 disassembly could be chronic loss of vasodilation and even edema, once cell-cell communication between cells is disrupted (Bird et al., 2013). Treatment with CLA, either as t10,c12 CLA or a 1:1 mix of c9,t11 and t10,c12 CLA at a suitable dose may be useful as a treatment for endothelial dysfunction based on their ability to protect against Src- mediated disruptions of Cx43 gap junction function.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Highlights

- GF/Cyt, alone or in combination, promote endothelial dysfunction and are rescued by the nutraceutical CLA.
- GF/Cyt coupled directly to Src mediate this damage, and rescue is achieved by t10,c12 CLA or a 1:1 mix of c9,t11 and t10,c12 CLA.
- Interleukins are less damaging than GFs, and are more resistant to CLA rescue.

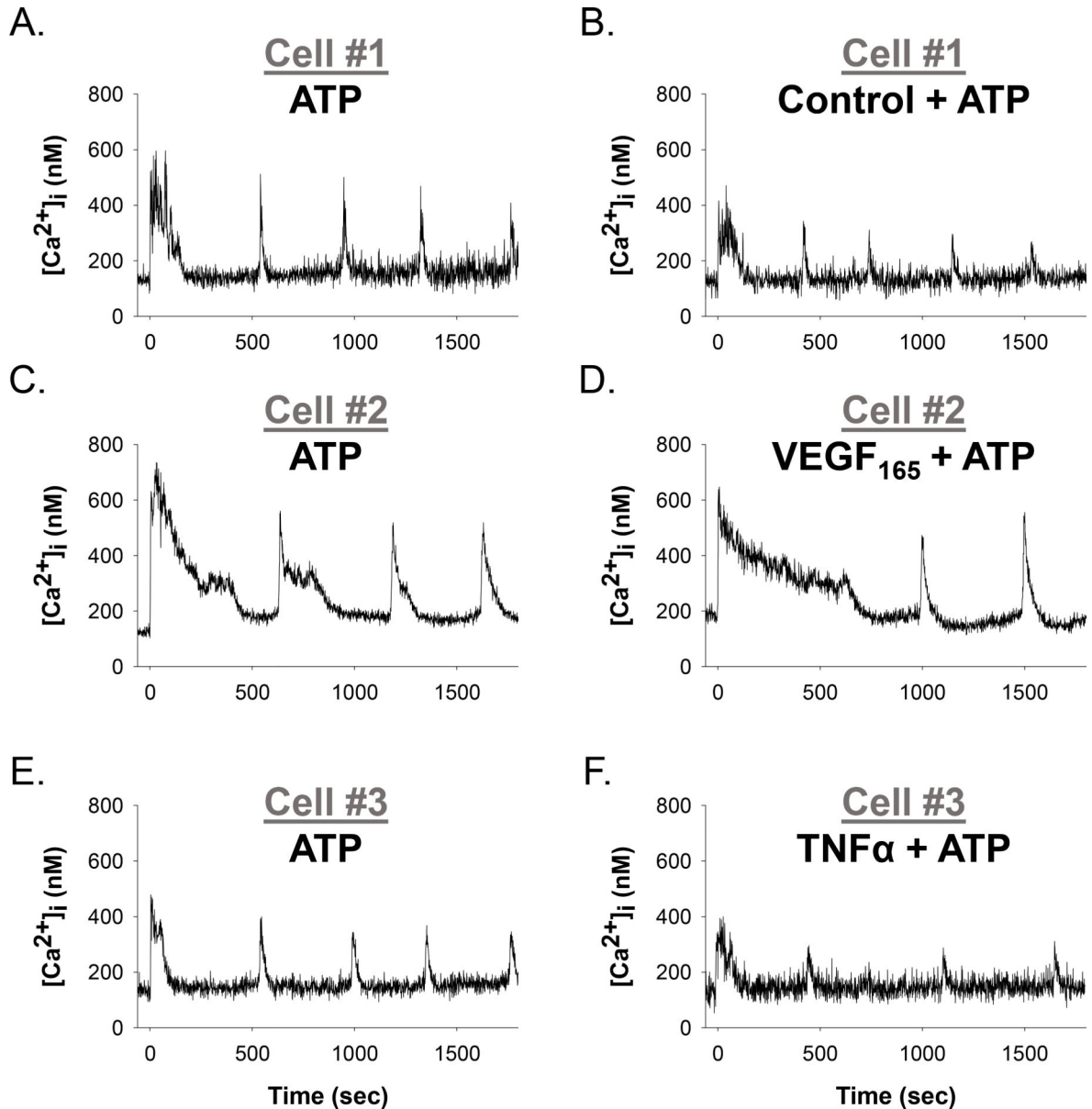


Figure 1: Representative individual HUVEC Ca^{2+} tracings before and after Control, VEGF_{165} , or $\text{TNF}\alpha$ treatment.

Panels A, C, and E show representative HUVEC ($[\text{Ca}^{2+}]_i$) tracings with a short baseline followed by 100 μM ATP stimulation (at time 0 sec), resulting in initial ($[\text{Ca}^{2+}]_i$) peak followed by repetitive Ca^{2+} bursts. Panel B shows the same cell as panel A, with addition of a vehicle control for 30 minutes and subsequently stimulated with ATP (at time 0 sec). Panel D shows the same cell as Panel C, with the addition of 10ng/ml VEGF_{165} for 30 minutes before subsequent stimulation with ATP (at time 0 sec). Likewise, Panel F shows the same cell as in Panel E, with the addition of 10ng/ml $\text{TNF}\alpha$ for 60 minutes before subsequent stimulation with ATP (at time 0 sec). Ca^{2+} The resulting Ca^{2+} burst loss provides an *in vitro* functional assay for growth factor and cytokine-mediated endothelial dysfunction as seen in

PE. For Panel A to Panel B, zero burst loss; for Panel C to Panel D, 1 burst loss – 25% reduction; for Panel E to Panel F, 1 burst loss – 20% reduction.

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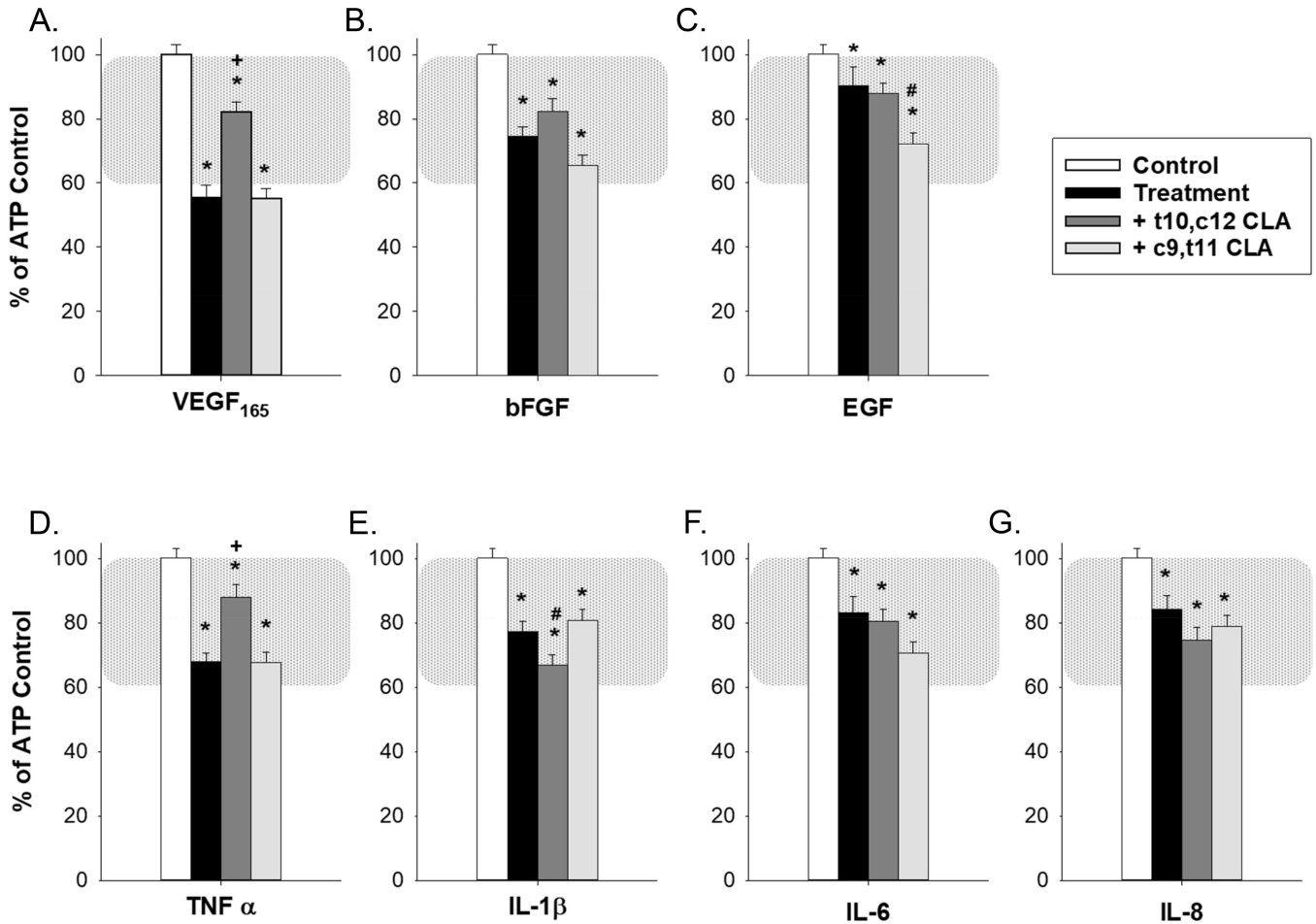


Figure 2: Growth factor and cytokine pretreatment inhibits ATP-stimulated Ca²⁺ bursts and is partially reversed by t10,c12 CLA for VEGF₁₆₅ and TNF α .

Pretreatment with each of seven growth factors and cytokines used at 10ng/ml for 30 minutes (VEGF₁₆₅ (panel A), bFGF (panel B), EGF (panel C), TNF α (panel D, 60 minutes), IL-1 β (panel E), IL-6 (panel F), and IL-8 (panel G)) inhibit ATP stimulated Ca²⁺ bursts in confluent HUVEC. Treatment with 50uM c9,t11 CLA for 30 minutes prior to growth factor or cytokine addition failed to reverse Ca²⁺ burst inhibition for any treatment and significantly exacerbated ATP-stimulated Ca²⁺ burst inhibition for EGF. Similar treatment with 50uM t10,c12 CLA partially reversed VEGF₁₆₅ and TNF α -mediated inhibition of ATP-stimulated Ca²⁺ responses and exacerbated ATP-stimulated Ca²⁺ burst inhibition for IL-1 β . Data is presented as mean Ca²⁺ burst numbers for those cells that showed 3+ bursts on initial ATP treatment \pm SEM on a minimum of 75 cells from at least 4 separate dishes. Significance by Rank-sum test is indicated as p<0.05 for * Control vs Treatment (w/ or w/o CLA); + Significant improvement after CLA pretreatment vs Treatment; # Significant exacerbation of Ca²⁺ burst inhibition by CLA pretreatment vs Treatment. Typical physiological range is shown with shaded region as defined in (Boeldt et al., 2017, Bird et al., 2013), adjusted to HUVEC.

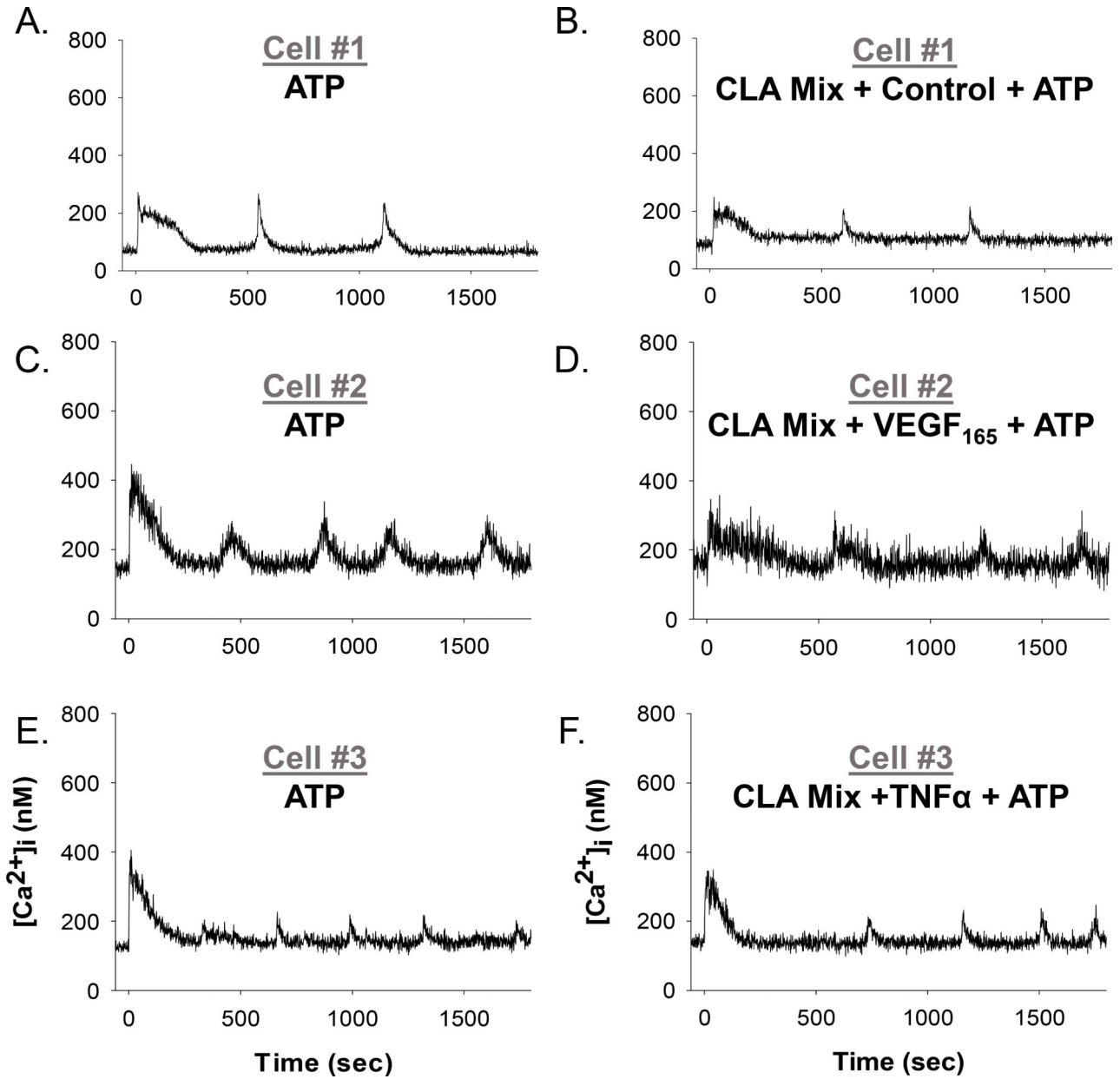


Figure 3: Representative individual HUVEC Ca^{2+} tracings before and after CLA Mix pretreatment and Control, $VEGF_{165}$, or $TNF\alpha$ treatment.
 Panels A, C, and E show representative HUVEC ($[Ca^{2+}]_i$) tracings with a short baseline followed by 100uM ATP stimulation (at time 0 sec), resulting in initial ($[Ca^{2+}]_i$) peak followed by repetitive Ca^{2+} bursts. Panel B shows the same cell as panel A, with addition of a 10um CLA mix pretreatment for 30 minutes before vehicle control treatment for 30 minutes and subsequently stimulated with ATP (at time 0 sec). Panel D shows the same cell as Panel C, with the addition of 10um CLA mix pretreatment for 30 minutes before 10ng/ml $VEGF_{165}$ for 30 minutes and subsequent stimulation with ATP (at time 0 sec). Likewise, Panel F shows the same cell as in Panel E, with the addition of 10um CLA mix pretreatment for 30 minutes before 10ng/ml $TNF\alpha$ for 60 minutes and subsequent stimulation with ATP

(at time 0 sec). An improvement in the degree of lost bursting as seen in Figure 1 established CLA-mediated rescue of Ca^{2+} bursting function.

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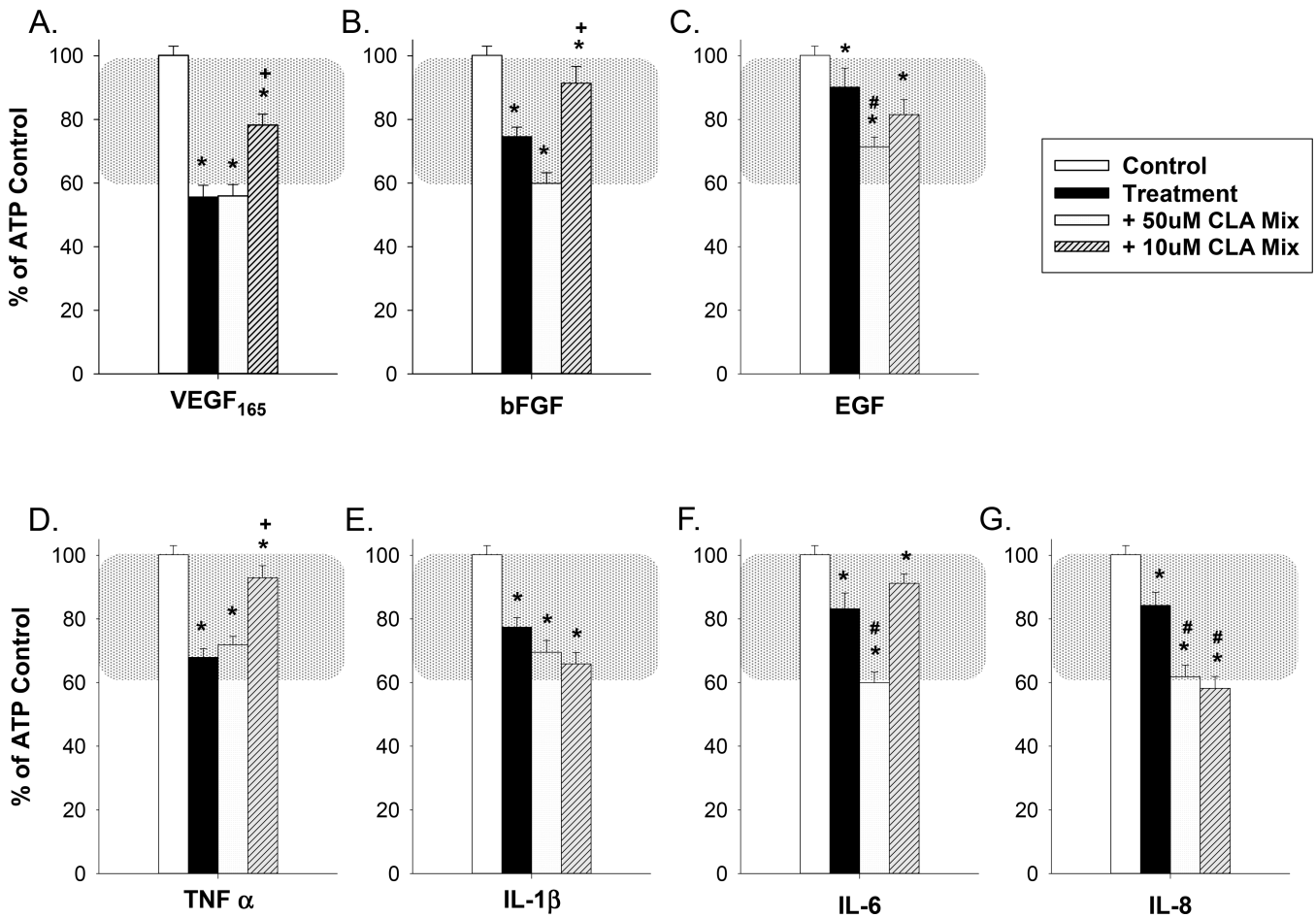


Figure 4: Inhibition of ATP-stimulated Ca^{2+} bursts is partially rescued by a 10uM 1:1 mix of c9,t11 and t10,c12 CLA isomers for VEGF₁₆₅, TNF α , and bFGF.
 Confluent HUVEC were pre-exposed to a 1:1 mix of either 50uM c9,t11 CLA and 50uM t10,c12 CLA or 10uM c9,t11 CLA and 10uM t10,c12 CLA for 30 minutes before growth factor or cytokine treatment and subsequent 100uM ATP treatment. ATP-stimulated Ca^{2+} bursts remained significantly reduced compared to control for all CLA-exposed treatments. The 50uM CLA mix failed to improve ATP-stimulated Ca^{2+} bursts for any growth factor or cytokine and exacerbated the Ca^{2+} burst inhibition for EGF (panel C), IL-6 (panel F), and IL-8 (panel G). The 10uM CLA mix significantly improved ATP-stimulated Ca^{2+} bursts for VEGF₁₆₅ (panel A), bFGF (panel B), and TNF α (panel D). The 10uM CLA mix exacerbated Ca^{2+} burst loss for IL-1 β (panel E) and IL-8 (panel G). Data is presented as mean Ca^{2+} burst numbers for those cells that showed 3+ bursts on initial ATP treatment \pm SEM on a minimum of 75 cells from at least 4 separate dishes. Significance by Rank-sum test is indicated as $p < 0.05$ for * Control vs Treatment (w/ or w/o CLA); + Significant improvement after CLA pretreatment vs Treatment; # Significant exacerbation of Ca^{2+} burst inhibition by CLA pretreatment vs Treatment. Typical physiological range is shown with shaded region as defined in (Boeldt et al., 2017, Bird et al., 2013), adjusted to HUVEC.

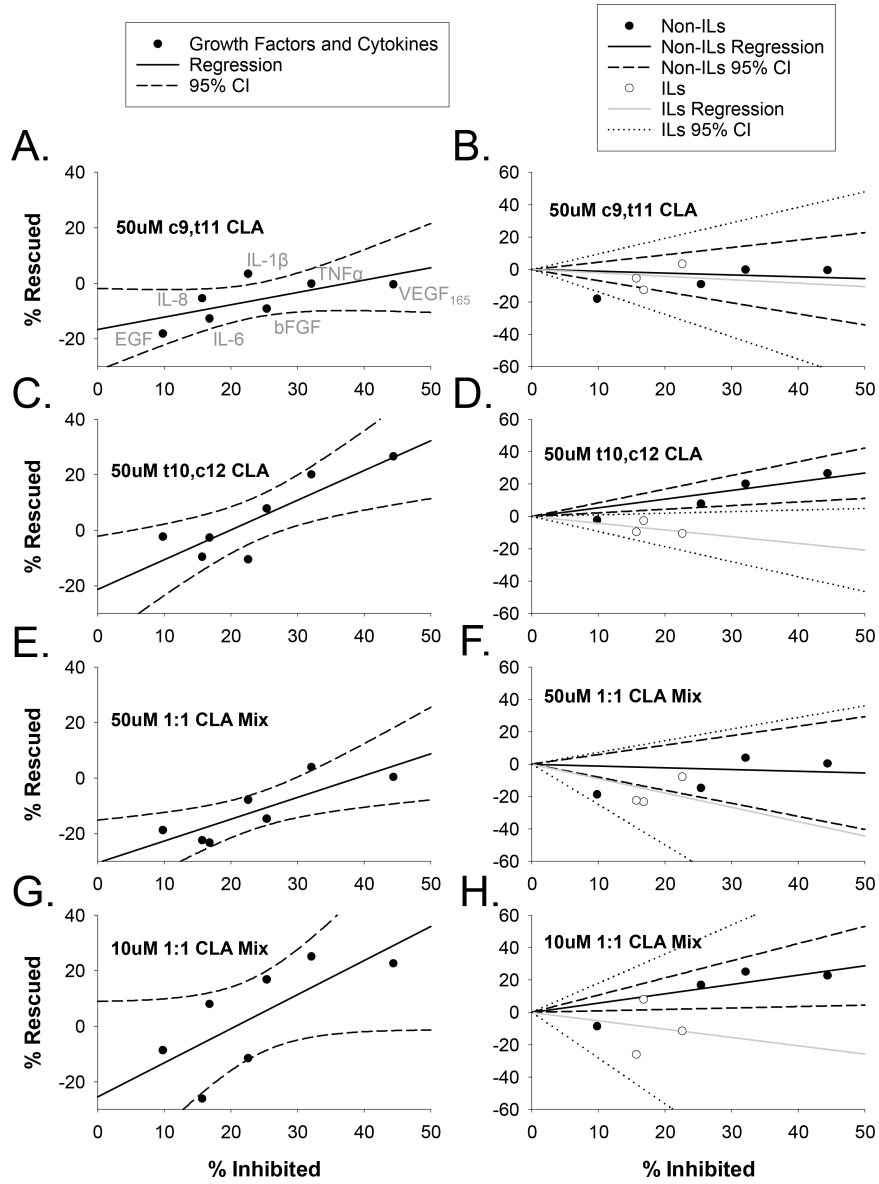


Figure 5: Regression analysis for the effect of CLA treatments on individual growth factors and cytokines.

Linear regressions were executed with percentage of bursts inhibited plotted on the x-axis and percentage “rescued” indicating change in burst numbers for all CLA and growth factor treatments from Figures 2 and 4 plotted on the y-axis. Experimental conditions are matched left (all data together; panels A,C,E,G) and right (separated into non-interleukins (Non-ILs) and Interleukins (ILs); panels B,D,F,H). All data points correspond to the same growth factors and cytokines from L to R: EGF, IL-8, IL-6, IL-1 β , bFGF, TNF α , and VEGF₁₆₅; as indicated in the upper left panel. For the left panels (A,C,E,G), regression lines are indicated by a solid black line and 95% confidence intervals (CI) are indicated by dashed black lines. For panels to the right (B,D,F,H), Non-ILs are indicated by black circles, black regression line, and black dashed CIs; for left panels, ILs are indicated by open circles, gray regression line, and dotted CIs.

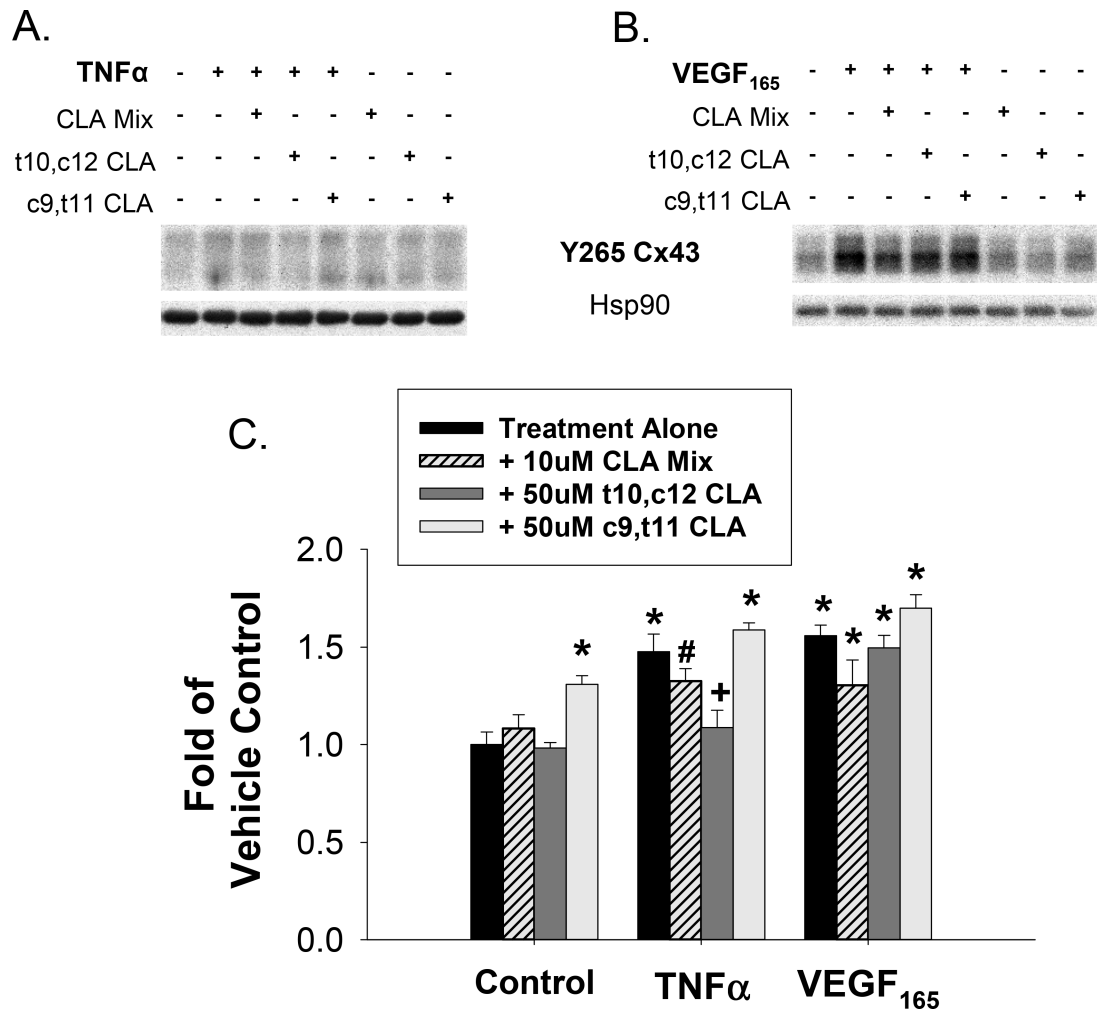


Figure 6: TNF α and VEGF₁₆₅ promote phosphorylation of Cx43 at residue Y265 and TNF α stimulated phosphorylation is reversed by t10,c12 CLA.
 Phospho- specific western blot analysis for Y265 on Cx43 were done and representative blots are shown. HUVEC were pretreated with CLA isomers or a 1:1 mix for 30 minutes prior to TNF α (1 hr, panel A) or VEGF₁₆₅ (30 min, panel B) stimulation. Each band was normalized to Hsp90 loading control and expressed as fold of unstimulated control. In panel C, values are depicted as mean \pm S.E.M. with a minimum of 6 independent experiments. Statistical significance is indicated by * $p < 0.01$ vs unstimulated control, # $p < 0.05$ vs unstimulated control, and + $p < 0.01$ vs treatment alone by post-hoc Turkey HSD one-way ANOVA with Holm interference.

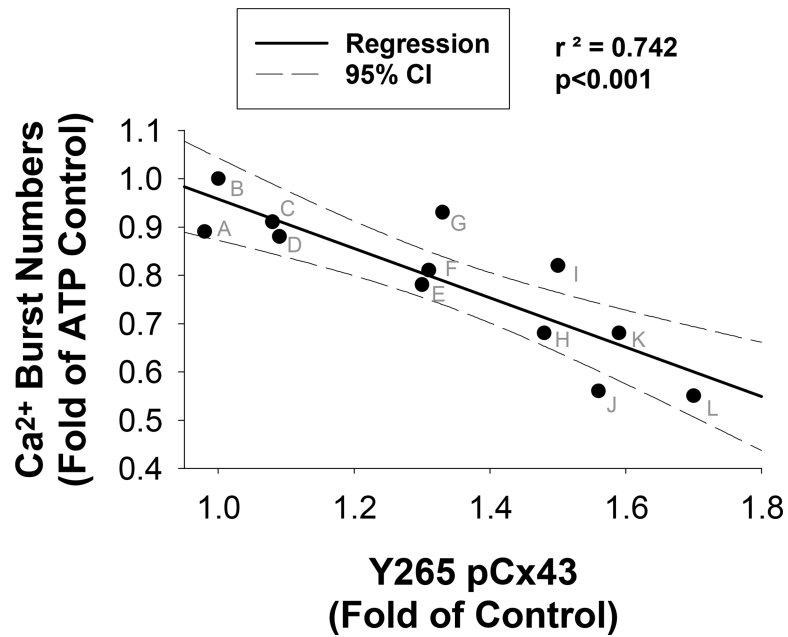


Figure 7: Regression Analysis of Ca²⁺ Burst and phospho-Cx43 for VEGF₁₆₅, TNF α , and CLA isomers.

Mean Ca²⁺ Burst numbers as fold of ATP control (from figures 2 and 4) on the X-axis and mean Y265 phosphor-Cx43 western blot data as fold of control (from figure 6) are plotted. A linear regression with 95% confidence intervals reveal a significant negative correlation between ATP-stimulated Ca²⁺ bursts and Y265 phosphorylation ($p < 0.001$). Treatments consisted of the following (indicated by letters above according to mean Y265 phosphorylation); A: 50uM t10,c12 CLA; B: Control; C: 10uM CLA mix; D: TNF α + 50uM t10,c12 CLA; E: VEGF₁₆₅ + 10uM CLA Mix; F: 50uM c9,t11 CLA; G: TNF α + 10uM CLA Mix; H: TNF α ; I: VEGF₁₆₅ + 50uM t10,c12 CLA; J: VEGF₁₆₅; K: TNF α + 50uM c9,t11 CLA; L: VEGF₁₆₅ + 50uM c9,t11 CLA.

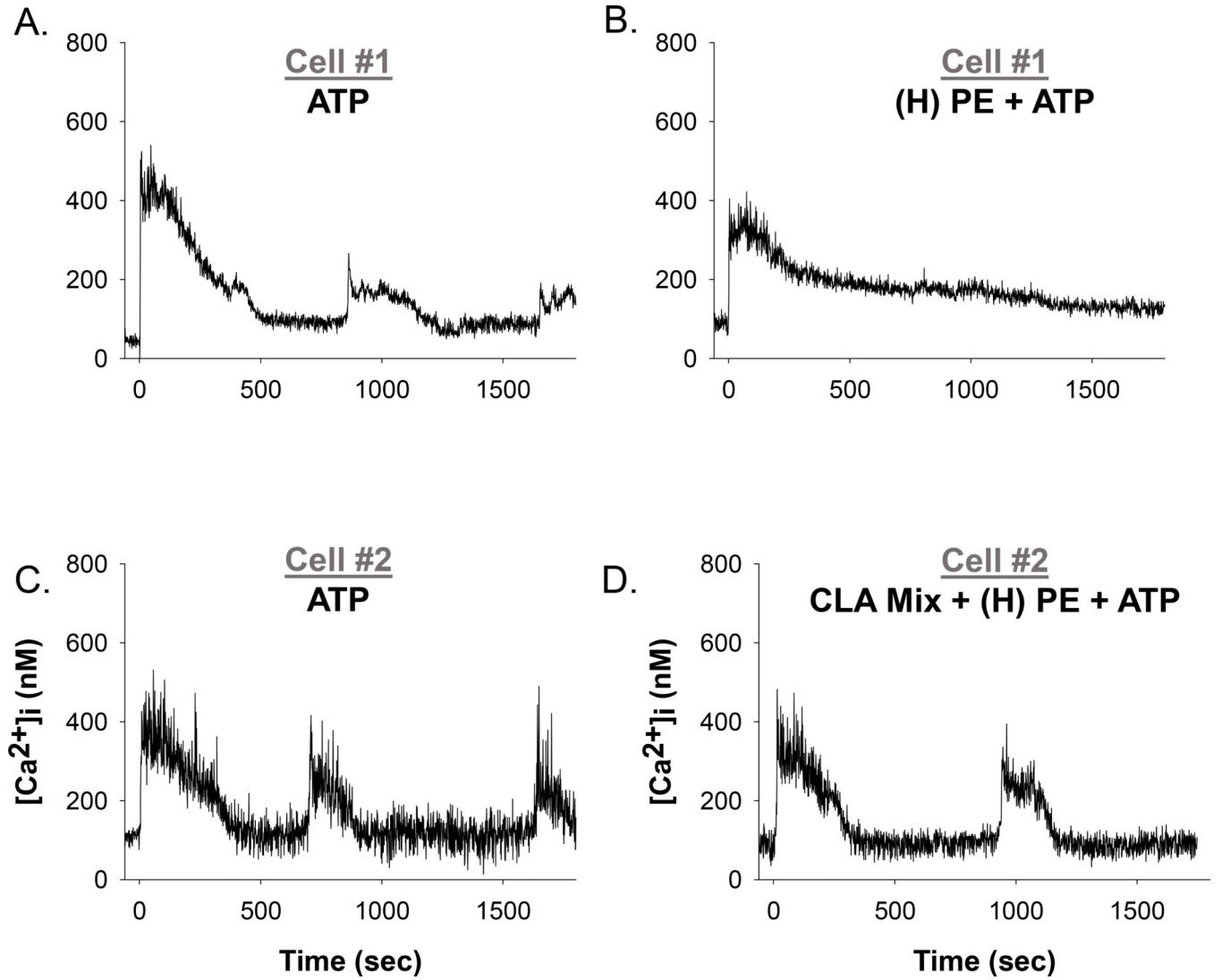


Figure 8: Representative individual HUVEC Ca^{2+} tracings before and after (H) PE cocktail treatment and in combinations with CLA Mix pretreatment.

Panels A and C show representative HUVEC ($[Ca^{2+}]_i$) tracings with a short baseline followed by 100 μ M ATP stimulation (at time 0 sec), resulting in initial ($[Ca^{2+}]_i$) peak followed by repetitive Ca^{2+} bursts. Panel B shows the same cell as panel A, with addition of (H)PE cocktail treatment for 30 minutes and subsequently stimulated with ATP (at time 0 sec). Panel D shows the same cell as Panel C, with the addition of 10 μ M CLA mix pretreatment for 30 minutes before (H)PE cocktail treatment for 30 minutes and subsequent stimulation with ATP (at time 0 sec).

VEGF₁₆₅ + TNF α Cocktails

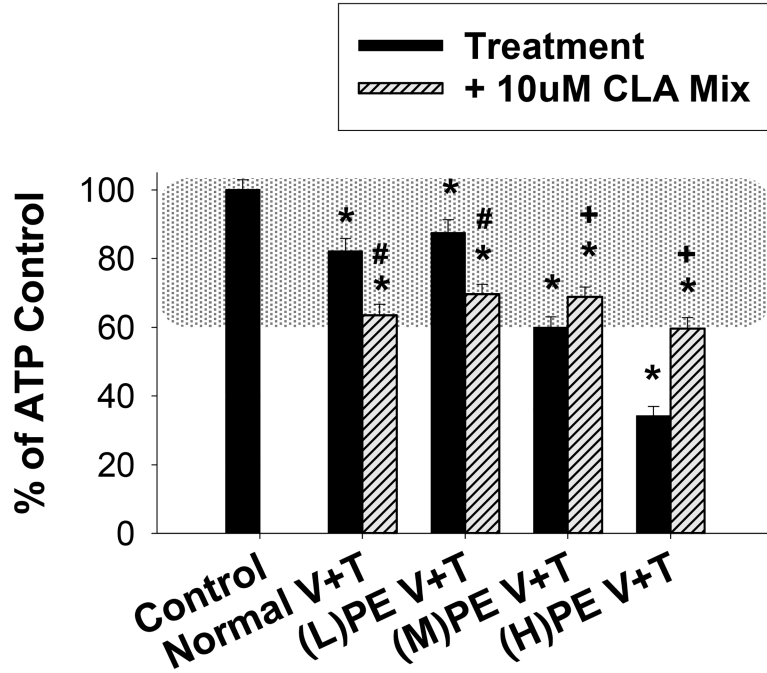


Figure 9: The 10uM CLA mix rescues ATP-stimulated Ca²⁺ bursts only under Medium and High levels of VEGF₁₆₅ and TNF α .

Normal pregnancy (30 minutes 0.5ng/ml VEGF₁₆₅ and 0.03ng/ml TNF α) and PE-mimicking combination pretreatments (30 minutes) of varying doses of VEGF₁₆₅ and a single dose of TNF α ((L)PE V+T: 0.1ng/ml VEGF₁₆₅ and 0.5ng/ml TNF α ; (M)PE V+T: 1ng/ml VEGF₁₆₅ and 0.5ng/ml TNF α ; (H)PE V+T: 10ng/ml VEGF₁₆₅ and 0.5ng/ml TNF α) inhibited ATP-stimulated Ca²⁺ bursts in a dose-dependent manner. A 30 minute pretreatment with the 10uM 1:1 mix of CLA isomers exacerbated the inhibition due to the Normal V+T, (L)PE V+T combination, but rescued ATP-stimulated Ca²⁺ bursts for the (M)PE V+T and (H)PE V+T combinations. Data is presented as mean Ca²⁺ burst numbers for those cells that showed 3+ bursts on initial ATP treatment \pm SEM on a minimum of 75 cells from at least 4 separate dishes. Significance by Rank-sum test is indicated as p<0.05 for * Control vs Treatment (w/ or w/o CLA); + Significant improvement after CLA pretreatment vs Treatment; # Significant exacerbation of Ca²⁺ burst inhibition by CLA pretreatment vs Treatment. Typical physiological range is shown with shaded region as defined in (Boeldt et al., 2017, Bird et al., 2013), adjusted to HUVEC.

VEGF₁₆₅ + TNF α + IL-6 Cocktails

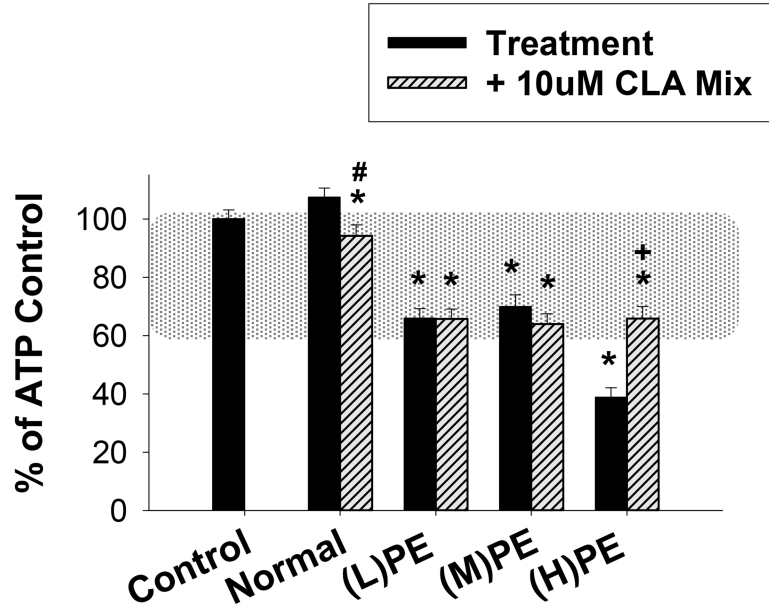


Figure 10: Preeclamptic-like cytokine cocktails strongly inhibit ATP-stimulated Ca²⁺ bursts and are partially rescued by CLA in the high VEGF₁₆₅ condition.

Cocktails corresponding to Normal (0.5ng/ml VEGF₁₆₅, 0.03ng/ml TNF α , 0.1ng/ml IL-6) or PE ((L)PE: 0.1ng/ml VEGF₁₆₅, 0.5ng/ml TNF α , 3ng/ml IL-6; (M)PE: 1ng/ml VEGF₁₆₅, 0.5ng/ml TNF α , 3ng/ml IL-6; (H)PE: 10ng/ml VEGF₁₆₅, 0.5ng/ml TNF α , 3ng/ml IL-6) circulating concentrations were applied to HUVEC. Cells exposed to the Normal cocktail were unchanged in number of ATP-stimulated Ca²⁺ bursts, and 30 minutes pretreatment with the 10uM CLA mix showed a slight reduction in Ca²⁺ bursts. Cells exposed to (L)PE and (M)PE cocktails showed modest levels of ATP-stimulated Ca²⁺ burst inhibition which were unchanged by CLA pretreatment. Exposure to the (H)PE cocktail gave severe Ca²⁺ burst inhibition, which was partially rescued using the 10uM 1:1 CLA mix. Data are presented as mean Ca²⁺ burst numbers for those cells that showed 3+ bursts on initial ATP treatment \pm SEM on a minimum of 75 cells from at least 4 separate dishes. Significance by Rank-sum test is indicated as p<0.05 for * Control vs Treatment (w/ or w/o CLA); + Significant improvement after CLA pretreatment vs Treatment; # Significant exacerbation of Ca²⁺ burst inhibition by CLA pretreatment vs Treatment. Typical physiological range is shown with shaded region as defined in (Boeldt et al., 2017, Bird et al., 2013), adjusted to HUVEC.