



**UNIVERSIDADE FEDERAL DE UBERLÂNDIA  
INSTITUTO DE GENÉTICA E BIOQUÍMICA  
PÓS-GRADUAÇÃO EM GENÉTICA E BIOQUÍMICA**

**Análise do papel da metformina na via insulínica, não-insulínica e inflamatória**

**Aluno:** Leonardo Gomes Peixoto

**Orientador:** Foued Salmen Espindola

**UBERLÂNDIA – MG  
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(Foued Salmen Espindola)

## Dedicatória

Dedico a todas as pessoas que conseguiram compartilhar comigo além daquilo que tem, aquilo que é.

Despedidas e saudades

Encontros e desencontros

Começo e recomeço

Tristezas e alegrias

Sonhos e agora realidade

Tudo isso, misturado com um sincero sorriso demistifica a apessoa que sou!

E tenho que admitir: - Sou grato. Por cada palavra amiga, cada abraço inesperado, cada sonho partilhado e por cada realidade vivida.

Palavras estas vindas de abraços que se tornaram sonhos nesta realidade que sou.

Obrigado por tudo...

Pais: pela força, expectativa, confiança e amor.

Irmãos: Fabrícios e Luciana pelos momentos partilhados.

Parentes: pelo incentivo e determinação.

Namorada: pelos sorrisos e abraços constantes.

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## **Apresentação**

Os estudos apresentados nesta tese foram delineados com o intuito de (1) avaliar o papel da metformina sobre os marcadores inflamatórios séricos de indivíduos diabéticos; de (2) descrever o papel da metformina e da insulina na via de sinalização relacionada à diminuição da sensibilidade insulínica no músculo esquelético de ratos diabéticos e de (3) avaliar o papel da metformina e da insulina na via de sinalização inflamatória e apoptótica em músculo esquelético de ratos diabéticos.

A metformina é uma droga amplamente usada no tratamento do diabetes tipo 2. Esta droga é considerada um sensibilizador de insulina, uma vez que diminui os níveis glicêmicos sem aumentar a secreção de insulina. Além disso, há estudos que comprovam que a metformina diminui a produção hepática de glicose, além de aumentar a disponibilidade de glicose periférica e reduzir a absorção intestinal deste carboidrato.

O mecanismo de ação molecular da metformina ainda não é claro. Até o momento, evidências sugerem quatro vias: 1ª inibir o complexo 1 mitocondrial (NADH-coenzima Q oxidoredutase, também conhecida como NADH desidrogenase), 2ª ativar a proteína quinase dependente de AMP (AMPK), 3ª inibir a adenosina 3',5'-monofosfato cíclico induzida por glucagon e 4ª inibir a enzima mitocondrial glicerol-3-fostato desidrogenase (GPDH).

Dentre estas 4 vias de sinalização, a melhor estudada é a inibição do complexo 1 mitocondrial, que leva a um desbalanço na relação da adenosita tri-fosfato/adenosina di-fosfato (ATP/ADP). Este desbalanço desencadeia uma complexa rede de reações de ativação e inibição das vias de sinalização inflamatória e apoptótica. Assim, no capítulo 1 são resumidos os estudos sobre a captação da metformina e seu mecanismo de ação, bem como seu mecanismo de ação no músculo esquelético, destacando a via de sinalização inflamatória desencadeada por “toll like receptor” (TLRs).

Além disso, e devido a seu complexo mecanismo de ação, a metformina tem sido avaliada em diferentes patologias. Vários estudos demonstraram que a metformina além de atenuar a resistência insulínica e melhorar a homeostase da glicose, tem sido relacionada ao tratamento de várias doenças, como o câncer e a síndrome do ovário policístico. Sabe-se que o diabetes associado ou não à outras patologias leva a um aumento do quadro de inflamação do paciente. Assim, o capítulo 2 é um manuscrito de meta-análise sobre o papel da metformina sobre os marcadores inflamatórios séricos e parâmetros metabólicos em pacientes diabéticos. Aqui, excluímos todos os manuscritos que relacionavam o diabetes a outras doenças que aumentariam os marcadores inflamatórios. Assim, neste estudo, mostramos que a inflamação causada pelo diabetes aumenta marcadores inflamatórios como a proteína C reativa (CRP), o fator de necrose tumoral (TNF $\alpha$ ) e o hormônio adiponectina. Como esperado, a metformina contribui diminuindo os valores dos marcadores inflamatórios séricos regulando a glicemia e hemoglobina glicada (HbA1c).

Embora existam vários estudos mostrando as diferentes vias de sinalização intracelular desencadeadas pela metformina, ainda há poucas evidências sobre o papel desta droga sobre a via de sinalização inflamatória/apoptótica no músculo esquelético de ratos diabéticos induzidos por estreptozotocina. Por isso, neste estudo avaliamos a via de sinalização relacionada à resistência insulínica no músculo esquelético de ratos diabéticos tratados com metformina. Assim demonstramos no capítulo 3 que a metformina melhora a sensibilidade insulínica a partir da via de ativação da AMPK, Proteína Kinase dependente de Cálcio (CAMKK $\beta$ ) e Miosina-V, além de diminuir a via inflamatória intracelular relacionada à resistência insulínica. No capítulo 4, avaliamos a via de sinalização inflamatória e apoptótica desencadeada pelo sinal TLRs em ratos hiperglicêmicos. A metformina, além de diminuir os marcadores inflamatórios desencadeados por TLRs, inibe a via de sinalização apoptótica.

Em resumo, demonstramos que a metformina desempenha um importante papel tanto no controle dos marcadores inflamatórios séricos quanto nos intracelulares, melhorando a sensibilidade à insulina e aumentando a captação de glicose.

### ***1.1 Metformin uptake and its action mechanism: systematic review***

Metformin is an oral antidiabetic drug of the biguanide class. According of American Diabetes Association 2015 [1] metformin is one of the first-line drug of choice for the treatment of type 2 diabetes. This drug decreases hyperglycemia primarily by suppressing hepatic gluconeogenesis [2], and increases the peripheral glucose uptake as well as reduces hepatic glucose output [3].

It is well documented that the leg muscles have different morphological characteristics. Muscle fibers can be classified by their contractile property and its color: slow fibers – oxidative, red fibers, or Type I, and fast twitch fibers – anaerobic, white fibers, or Type II. The gastrocnemius muscle contains approximately 50% of anaerobic fibers while the soleus muscle has 75% to 90% of oxidative fibers. In this study we evaluated the effect of metformin action on gastrocnemius and soleus muscles of diabetic rats.

Metformin is actively transported into cells by organic cation transporters (OCT) 1, 2, and 3 (encoded by *SLC22A1*, *SLC22A2* and, *SLC22A3*), which are tissue specifically expressed at significant levels in various organs such as liver, muscle, and kidney [4, 5]. Dujic et al., showed that reduced OCT1 transport leads to metformin intolerance [6]. The liver is a major site of drug action, as it is exposed to high levels of oral metformin and expresses the metformin transporter OCT1, facilitating uptake and allowing for drug accumulation in this tissue. However, another tissues has been identify as sites to metformin action [7].

Evidence suggest that mechanisms of metformin action are: inhibition of the mitochondrial respiratory chain (complex I), activation of AMP-activated protein kinase (AMPK), inhibition of glucagon-induced elevation of cyclic adenosine monophosphate (cAMP) [8] as well as inhibition of mitochondrial glycerophosphate dehydrogenase [9].

To date, the best-known role for metformin is on the inhibition of mitochondrial complex. Complex I inhibition by metformin interrupts mitochondrial respiration and decreases proton-driven synthesis of adenosine triphosphate (ATP) [10], causing cellular energetic stress and elevation of the AMP/ATP ratio [11]. However, Hawley

et al., showed that effects of metformin on mitochondrial respiration vary among cells [12] indicating that it has not yet been possible to confirm whether or not complex I is the only mitochondrial target of metformin. Thus, more work is required to understand the underlying reason(s) for these variations.

The imbalance of ATP/ADP ratio from metformin action leads to activation of 5' AMP-activated protein kinase (AMPK). However, there are evidences that metformin activates AMPK independent of imbalance of APT/ADP [13]. Other studies showed that metformin activates AMPK through inhibition of AMP deaminase [14] or by upstream kinases serine/threonine kinase 1 (LKB1) in liver [15]. Recent finding show that metformin promote the formation of the AMPK  $\alpha\beta\gamma$  heterotrimeric complex, resulting in an increase of AMPK $\alpha$  phosphorylation by LKB1 [16]. Taken together, it is suggested that AMPK activation by metformin may occur by different pathway. In addition, it is know that this activation of the AMPK by metformin can inhibit gluconeogenesis and activates glycolysis in liver [17] and increase glucose consumption in muscle [18].

In condition as fasting or starvation the hormone glucagon is released and acts via cyclic AMP (cAMP) and cAMP-dependent protein kinase, promoting hepatic glucose production by inhibiting glycolysis and activating gluconeogenesis, both acutely and by long term effects on gene expression. Study shows that metformin rapidly reduces cAMP levels induced by glucagon in primary hepatocytes, and reduces phosphorylation of protein kinase A (PKA) substrates, including the 6-phosphofructo-2-kinase isoform (PFKFB1) [19]. Lowering of cAMP could inhibit the switch from glycolysis to gluconeogenesis triggered by glucagon [20]. These finding also suggest that metformin action is involved in inhibition of glucagon action by decrease of cAMP, and represents one of the AMPK-independent mechanisms by which biguanides acutely inhibit glucose production.

Finally, metformin treatment inhibits mitochondrial glycerophosphate dehydrogenase non-competitively and modulates cytosolic and mitochondrial redox state, inducing an effective reduction in endogenous glucose production [9]. These results identify mitochondrial glycerophosphate dehydrogenase as one of the primary molecular targets by which guanides/biguanides inhibit hepatic gluconeogenesis

Thus, metformin suppresses gluconeogenesis independently of AMPK, instead altering hepatic energy charge and inducing allosteric inhibition of glycolytic enzymes or adenylate cyclase and glucagon-activated gluconeogenic transcription.

### ***1.2 Intracellular mechanism of metformin in muscle cells***

Insulin is a glucose regulating hormone. When blood glucose is high, insulin facilitates the uptake of glucose into cells via translocation of glucose transporter 4 (GLUT4) independently AMPK activation. Insulin binds to its receptor (IR) which phosphorylates the substrate (IRS1) following an activation cascade: phosphatidylinositol 4,5 biphosphate - inositol 1,4,5 triphosphate - serine threonine kinase (AKT) - TBC1 domain family members; Akt substrate 160 (AS160) triggering the translocation of GLUT4.

On the other hand, metformin activates GLUT4 translocation by alternative pathway to uptake glucose from AMPK activation [14-16]. Several evidence indicated that in peripheral tissue the activation of AMPK increase GLUT4 translocation [21]. These facts indicate the existence of a metformin–AMPK–GLUT4 axis.

Lee et al., show that metformin induces GLUT4 translocation via increase of Rab4 expression by AMPK pathway [22]; and that the activities of Akt substrate 160 (AS160) and PKC-zeta are involved in metformin-induced Rab4 regulation [22]. The Rab protein is a member of the Ras G proteins. Rab GTPases regulate many steps in membrane traffic, including vesicle formation, movement, and membrane fusion [23]. Furthermore, study showed that myosin Va and RAB8a were co-localized in exocytose of GLUT4 vesicle of muscle cells [24].

Another mechanism for action of metformin is related to the decreased of insulin resistance by the inactivation of pro-inflammatory mediators produced by diabetes [25-27]. Increase of tumor necrosis factor- $\alpha$  (TNF $\alpha$ ) inhibits AMPK [28] which also may activates the insulin-mediated glucose uptake pathway [29]. Thus, metformin inhibits TNF $\alpha$  – induced IKK $\alpha$ / $\beta$  phosphorylation [28]. Thereby, there is a decrease of the intracellular signal for dissociation of nuclear factor (NF- $\kappa$ B) from its inhibitor (I $\kappa$ B) [30] allowing the action of NF- $\kappa$ B. Inflammatory mediator stimulated by NF- $\kappa$ B leads to phosphorylation of c-Jun-N-terminal kinase (JNK) [29]. p-JNK, in

turn, induces serine phosphorylation in insulin receptor substrate (IRS1), which inhibits insulin signal transduction [31] and decreased GLUT4 translocation. In addition, the metformin increases the tyrosine kinase activity of the IR in liver [32]. See below the scheme of molecular mechanism of insulin and metformin action (Figure 1).

The scheme – (figure 1. studied in chapter 3) shows different signaling pathways of the metformin action: 1<sup>th</sup> inhibition of complex I mitochondrial, 2<sup>th</sup> activation AMPK, 3<sup>th</sup> maintenance of NF- $\kappa$ B/I $\kappa$ B complex and 4<sup>th</sup> inhibition of TNF $\alpha$ . Once AMPK activates, this mechanism can regulate the insulin and non-insulin pathway by phosphorylation both IR and PI3K (insulin pathway) as AS160 (non-insulin pathway). On the other hand, metformin may improve insulin sensitivity by decrease of NF- $\kappa$ B and increase of I $\kappa$ B concentration and consequently inhibition of TNF $\alpha$ . This inhibition decrease the p-JNK and the phosphorylation of IRS1ser. Thus, metformin enhance of IR activity and GLUT4 translocation.

### ***1.2 Inflammatory pathway on skeletal muscle triggered by toll like receptor and the metformin role.***

Inflammation is pivotal in the development of insulin resistance in part via the activation of the toll-like receptors (TLRs), as TLR4 and TLR2 [33, 34]. TLRs are a class of proteins that play a key role in the innate immune system. TLR4 expression have been shown to be increased in conventional insulin resistance target tissues like skeletal muscle tissue of type 2 diabetic subjects [35]. TLR4 and TLR2 expression are increased in diabetes [36] and the activation of these receptors contributes to insulin resistance in muscle [35]. The signaling pathways triggered by TLRs involve the activation of the transcription factors NF- $\kappa$ B and JNK, which initiates the transcription of proinflammatory cytokine genes [37-39].

NF- $\kappa$ B plays a key role in regulating of genes involved in inflammatory pathways such as TNF $\alpha$ , adhesion molecules [40, 41] and chemokines CXCL1/KC [42]. Boyd et al show that TLR4 activates 20-fold NF- $\kappa$ B pathway than TLR2 in skeletal muscle [43]. NF- $\kappa$ B is inhibited by I $\kappa$ B under basal condition, and remain into cytoplasm. Upon proinflammatory pathway activation, as present in diabetes,

the kinase complex I $\kappa$ B is activated and catalyzes the phosphorylation of I $\kappa$ B leading to its degradation, which liberates NF- $\kappa$ B to translocate into the cell nucleus and stimulate the transcription of inflammatory mediators. JNK is activated by both mitogen-activated protein kinases (MKK4 and MKK7), which phosphorylate Thr-Pro-Tyr residues induced by several stress stimuli and is inhibited by JNK interacting protein (JIP) [44].

Evidence indicates that both JNK and NF- $\kappa$ B also participate of pro-apoptotic and anti-apoptotic pathways. JNK activation stimulated apoptosis by bcl-2-like protein 4 (BAX) translocation to mitochondria [45] and the phosphorylation of B-cell lymphoma 2 (Bcl2), suppressing its function [46]. NF- $\kappa$ B in turn, stimulates anti-apoptotic signal activates Bcl-2 expression [47]. BAX and Bcl2 are members of Bcl-2 family of proteins, which can be divided into pro-apoptosis (BAX, Bid, Bad, Bak, Bic, Bok, Bcl-XS, and Hrk) or anti-apoptosis (Bcl2, Bcl-XL, Bcl-W, A1 and Mcl-1) proteins [48].

There are few studies about metformin role in inflammatory pathway triggered by TLRs. Two studies show that metformin treatment is able the decrease TLR4 mRNA [49, 50]. On the other hand, several studies show that metformin acts on inflammatory markers triggered by TLRs as describe above.

The scheme – (figure 2. studied in chapter 4) shows the molecular mechanism of metformin action on inflammatory markers triggered by TLRs in different signaling pathways: 1<sup>th</sup> maintenance of NF- $\kappa$ B/I $\kappa$ B complex and 2<sup>th</sup> inhibition of both TNF $\alpha$  and CXCL1/KC and consequently inhibition of p-JNK, 3<sup>th</sup> inhibition of BAX and activation of Bcl2. Taken together, our data point out that metformin may attenuate the activation of the inflammatory pathway TLRs/NF- $\kappa$ B/TNF $\alpha$ /CXCL1/KC and the apoptotic signaling BAX:Bcl2 ratio/p-JNK independent of p-AMPK, which could be accompanied by a reduction of the inflammatory damage caused by hyperglycemia in skeletal muscle of diabetic rats.



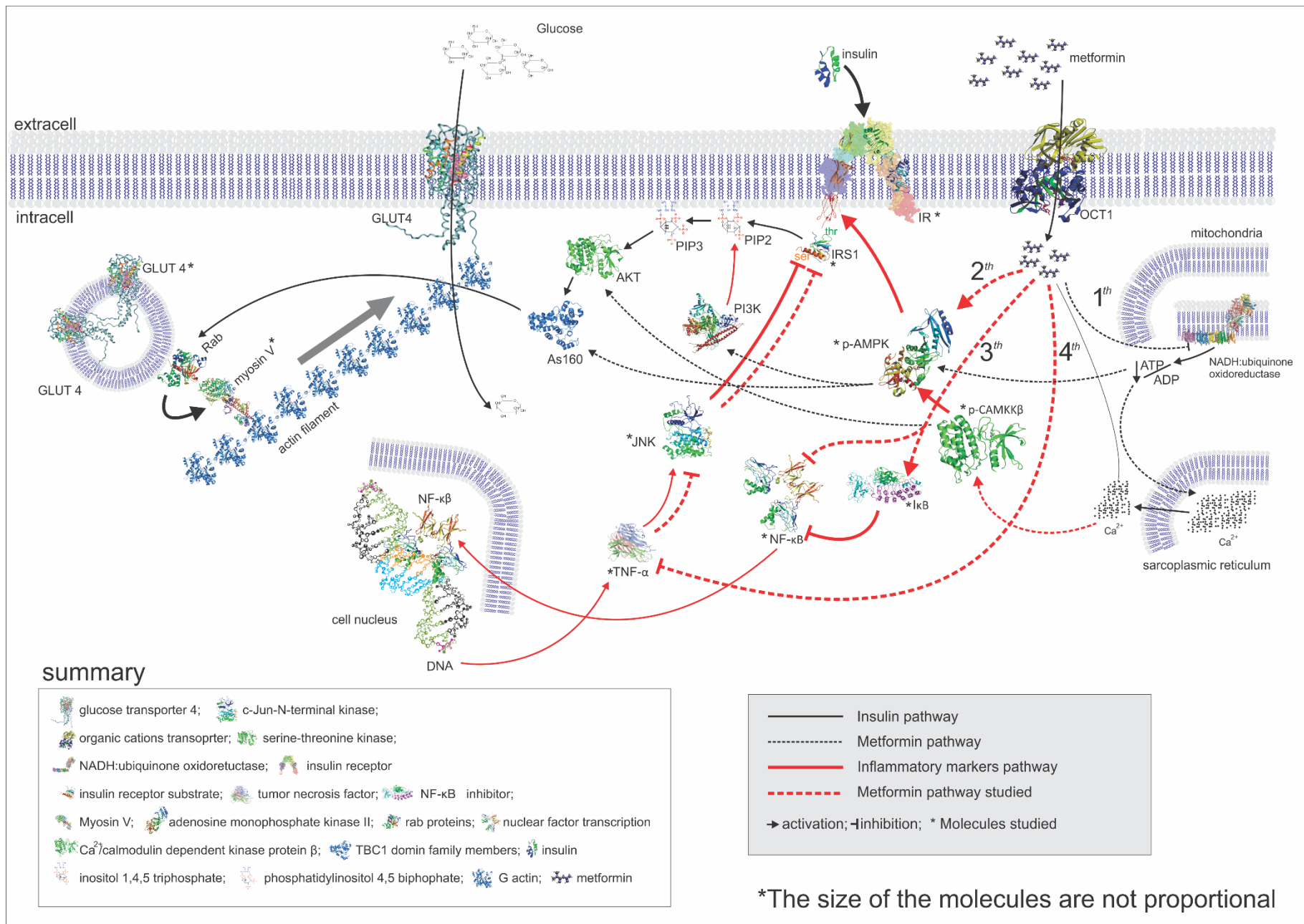
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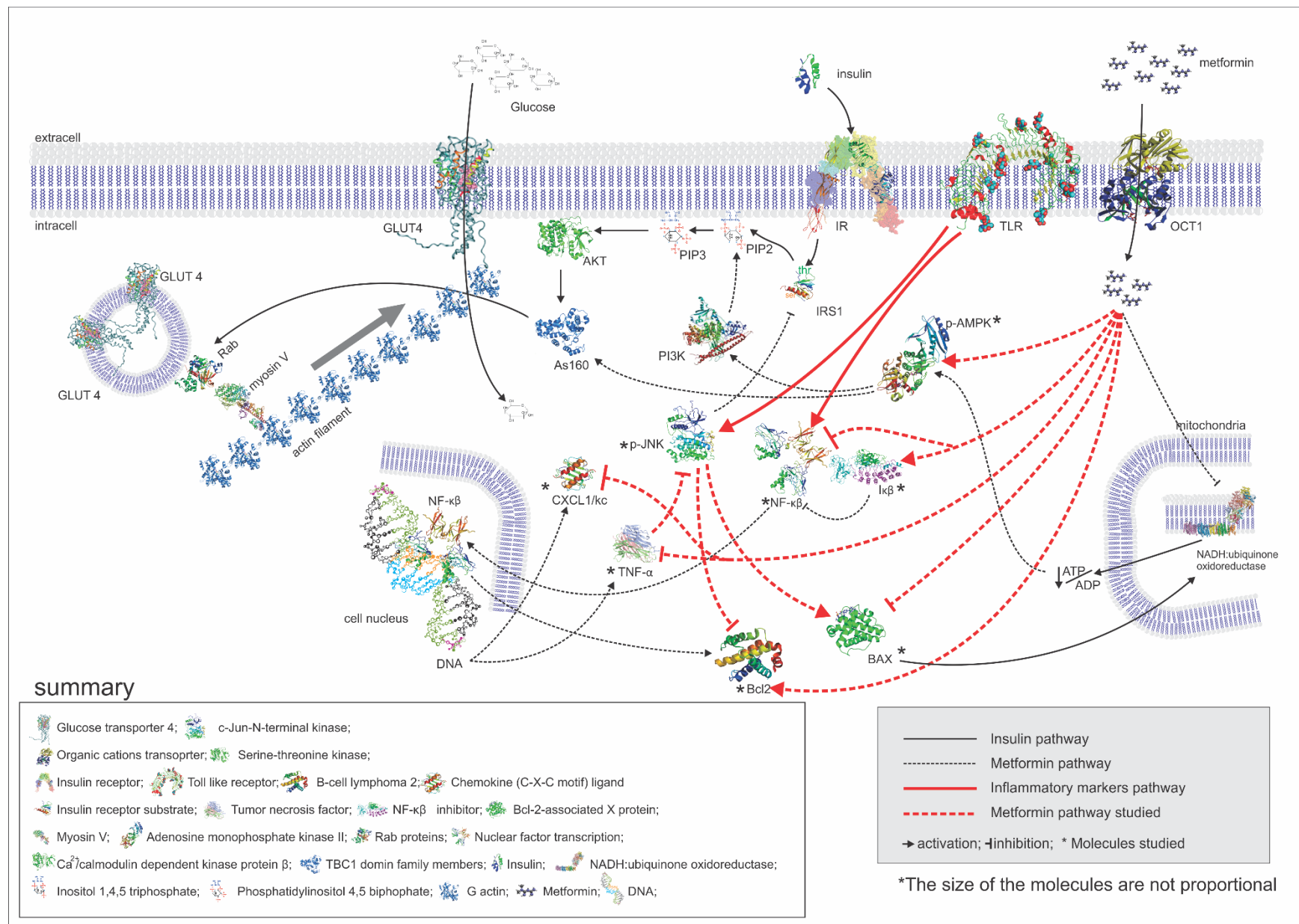
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**Figure 1. Scheme of insulin and metformin role on skeletal muscle of diabetic rat**



**Figure 2. Scheme of metformin role on inflammatory and apoptotic pathway triggered by TLRs in skeletal muscle of diabetic rats.**

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**Metformin on inflammatory markers: Meta-Analysis.**

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## Abstract

**Purpose:** We performed a meta-analysis of randomized trials to assess the effect of metformin on inflammatory markers and metabolic parameters in subjects with diabetes.

**Methods:** We performed comprehensive searches on NCBI, Cochrane, Science Direct databases from 1966 to Jun of 2015. We included randomized trials of at least 4 weeks duration that compared groups with diabetes before and after the treatment with metformin or metformin plus other drugs, and evaluated body mass index, blood glucose, HbA1c and inflammatory parameters such as C-reactive protein, tumor necrosis factor and adiponectin.

**Results:** Pooled results of the 26 trials, with 1760 participants at the end of treatment reduce BMI in 0.9%  $p=0,0043$ , as well as, decrease of blood glucose level [SMD -0,411 mg/dL, 95%CI -0,463 to -0,369,  $I^2= 56.62\%$ ], HbA1c [SMD -0.479%, 95%CI -0,568 to -0,390,  $I^2= 55.02\%$ ], CRP levels [SMD -0,274mg/dL, 95%CI -0,419 to -0,129,  $I^2= 72.78\%$ ], TNF $\alpha$  concentration [SMD -0,103pg/ml, 95%CI -0,514 to 0,309,  $I^2= 87.67\%$ ] and increase of adiponectin [SMD 0,171 $\mu$ g/ml, 95%CI 0,098 to 0,440,  $I^2= 81.09\%$ ] compared with pretreatment.

**Conclusion:** The long-treatment with metformin monotherapy or metformin plus other drugs improves metabolic parameters and induced changes in inflammatory markers in diabetic subject.

## Key words:

C-reactive proteins, tumor necrosis factor (TNF $\alpha$ ), adiponectin, blood glucose level, glycated hemoglobin (HbA1c)



## Introduction

Metformin is one of the first-line drug of choice for the treatment of diabetes [1]. This drug controls hyperglycemia by suppressing hepatic gluconeogenesis [2], and improve the peripheral glucose uptake [3]. Evidence suggests that metformin has an important role on inflammatory markers linked to diabetes and obesity. Risk factors such as family history of diabetes, physical inactivity, obesity, insulin resistance, hypertension among others can lead to the development of diabetes [1]. Diabetes is a group of metabolic diseases characterized by the high blood sugar levels over a prolonged period. Diabetes leads to the increase of the inflammatory markers such as C-reactive protein (CRP), proinflammatory cytokines such as tumor necrosis factor- $\alpha$  (TNF $\alpha$ ) and increasing insulin resistance [4].

Several studies have shown that the use of metformin decrease body mass index (BMI) attenuating obesity [5]. The increase of BMI is associated with the inflammation caused by diabetes [6]. Recent finding suggest that metformin decreases both TNF $\alpha$  and C-reactive protein [7]. Higher TNF $\alpha$  and CRP levels have been linked to the increase of insulin resistance in obese subjects [8, 9] and to the increase of the risk on later development of diabetes [10]. On the other hand, when the inflammatory signal increases there are an overexpression of the adiponectin by adipocytes. This hormone modulates a number of metabolic processes, including glucose regulation and fatty acid oxidation [11]. Adiponectin decreases insulin resistance and has a potent anti-inflammatory and anti-atherogenic effects [12]. Recently, an study showed that metformin is capable to up-regulate adiponectin gene expression [13]. These finding together probably indicate that metformin decreases the systemic inflammation.

This meta-analyses study on data of inflammatory markers and metabolic parameters from randomized trials with diabetics subjects treated with metformin or metformin plus other drugs was conducted to add new evidence to the effect of metformin on these parameters.

## Material and Methods

### Selection of Studies and statistical analysis

We performed comprehensive searches on NCBI, Cochrane, Science Direct databases from 1966 to Jun of 2015 using the terms metformin, diabetes, inflammatory cytokines, glucose uptake, TNF and trial and scanned selected journals and references of identified articles. We included randomized trials of at least 4 weeks duration that compared groups with diabetes before and after the treatment with metformin or metformin plus other drugs, and evaluated body mass index, blood glucose, HbA1c and inflammatory parameters such as C-reactive protein, tumor necrosis factor and adiponectin. We excluded trials evaluating inflammation associated with other diseases, because the metabolic derangements may be different from other forms of inflammatory. Interventions included metformin alone or in combination with other drugs.

Inclusion and exclusion criteria are shown in Table I.

The included studies used different outcome measures to quantify CRP. We used the mean values to this analyses and the difference was verified with the test sensitivity. Some studies reported the p-value in the group before and after treatment and, in cases that this value was not available the p-value was calculated by the value of the mean, standard deviation and number of sample by the statistical method of multiple t-test in the GraphPad Prism 6.

We used the Comprehensive Meta-Analysis software, version 3 free trial ([www.meta-analysis.com](http://www.meta-analysis.com)) for statistical calculations. We analyzed the *effect size* data *continuous* for one group – pre and post treatment with metformin or metformin and other drugs – by standardized mean difference (SMD) to *normalize the results*, the *size* and *expression interference effect* in each study regarding the observed variability.

The random effect was decided at  $I^2$  statistic value which represent heterogeneity (by the Chi-squared test). A value of 0% to 40% may not be important, from 30% to 60% may represent moderate, 50% to 90% may represent substantial and 75% to 100% considerable heterogeneity. Results were displayed in tables and graphically by Forest plots.

## Results and Discussion

### *Search results*

The search identified approximately 3,015 articles, of which 182 were potentially relevant (Figure 1). Of these, 82 were according to the inclusion criteria. Trials were excluded for the following reasons: 32 trials did not compare metformin before and after treatment, 11 trials conducted the study for less than 4 weeks duration, 13 trials evaluated another disease beyond diabetes, and 2 trials provided data of participants included in another trial.

### *Trial characteristics and data synthesis*

The analysis included 26 trials, with a total of 2006 participants at the beginning of the treatment with a 13% dropout. The mean duration of the trials was 22 weeks (range, 4-52 weeks), with a mean study size of 95 participants (range 20-473). The mean dose of metformin was 1.2 g/day (range, 500-2250 mg/day). In some studies the treatment with metformin was combined with the following drugs: Arcabose, Rosiglitazone, Pioglitazone, Garlic, Glargine, Exenatide and Atovarstatin (Tables 2).

BMI is expressed as weight (kilograms) divided by height (meters) squared, this parameter is often used to classify overweight (BMI>25) and obesity (BMI>30) [5]. This meta-analysis showed that metformin treatment reduced BMI (0.9%) compared with pretreatment (table 2). In the studies evaluated was not reported physical exercise for the participants, indicating that this weight loss was associated with the metformin action. Study shows that 1% reduction in weight is associated with a reduction in insulin resistance of 3% to 7% and a decrement in new cases of diabetes of 5% to 15% [37, 38]. Considering the 1760 participants, we observed a reduction of approximately 0.9% in BMI, representing a reduction in insulin resistance by approximately 4% and new cases of diabetes by approximately 7%. Thus, these findings suggest that metformin treatment can attenuate weight gain and decrease insulin resistance.

### *Glycaemia and HbA1c levels*

We performed the meta-analysis to investigate the metformin role on serum inflammatory and metabolic markers levels in diabetic patients. It is well documented that metformin is an anti-glycemic drug. Corroborating with this

postulation, we show that metformin treatment decrease both blood glucose level [SMD -0,411 mg/dL, 95%CI -0,463 to -0,369,  $I^2= 56.62\%$ ] and HbA1c levels [SMD -0.479%, 95%CI -0,568 to -0,390,  $I^2= 55.02\%$ ]. Another study showed that metformin alone and combined with vildagliptin decrease glycaemia and HbA1c levels [14]. Metformin decreases hyperglycemia by suppressing hepatic glucose production [2] and as well as reducing hepatic glucose output [3]. In addition, metformin increases insulin sensitivity, enhances peripheral glucose uptake and decreases insulin-induced suppression of fatty acid oxidation [15]. In animal models metformin decreases the absorption of glucose from the gastrointestinal tract [16].

In addition, HbA1c is formed in a non-enzymatic/enzymatic glycation pathway by the hemoglobin's exposure to plasma glucose. It is a form of hemoglobin that is measured primarily to identify the average plasma glucose concentration over prolonged periods of time. Our finding is in accordance with other studies that showed a decrease of HbA1c in metformin monotherapy treatment [17, 18].

#### *Serum inflammatory markers*

Our findings show that metformin treatment with or without combination with other drug reduced CRP levels [SMD -0,274mg/dL, 95%CI -0,419 to -0,129,  $I^2= 72.78\%$ ] (Figure 3). CRP is generated by the liver during acute infection or inflammation, and its concentration in serum may increase as much as 1,000-fold under the condition of injury and infection [29]. Elevated CRP levels have been linked to an increase risk of later development of diabetes [10]. Metformin also reduces serum CRP levels in women with polycystic ovary syndrome [39]. The mechanism through which metformin decreases CRP is still unclear. Evidence indicates that the decrease in CRP is associated with the decrease of waist circumference [40] and serum levels of endothelin-1 [41]. In summary, the main results of our meta-analysis were in accordance to previous findings in which metformin may reduce serum CRP levels in diabetic patients, possibly indicating a decrease of the degree of low-grade inflammation.

TNF $\alpha$  is an inflammatory marker also find in serum of diabetic patients. Overproduction of TNF $\alpha$  by adipose tissue is involved in insulin resistance and obesity, which leads to inflammatory responses triggered by macrophages [8, 9].

Thus, TNF $\alpha$  promote unnecessary systemic inflammation and trigger insulin resistance [42]. In this study, we verified that the treatment with the combination of metformin and another drug decrease TNF $\alpha$  concentration compared to pretreatment [SMD -0,261 pg/ml, 95%CI -0,397 to -126, I<sup>2</sup>= 87.67%] (Figure 4). Interestingly, our results show that metformin treatment without pioglitazone do not reduced the TNF $\alpha$  levels, this can be related to the dose and duration of the treatment. Kim et al. showed that 12 weeks of metformin treatment without rosiglitazone combination do not decrease CRP, TNF $\alpha$  and IL6 levels [28]. Instead, studies in animal models show that metformin modulates the expression of oxidative stress genes, restoring the mRNA of glutathione S-transferase- $\alpha$  (GST $\alpha$ ), NAD(P)H quinone oxidoreductase (NQO1), and catalase genes and inhibiting TNF $\alpha$  and IL-6 pro-inflammatory genes [43]. In addition, metformin is associated with the decrease of transcriptional factor –  $\kappa$ B (NF- $\kappa$ B) [44], which decreases TNF $\alpha$  expression. Thus, it is necessary more studies to understand how metformin acts in the decreased of TNF $\alpha$  in diabetic patients.

On the other hand, adiponectin decreases insulin resistance and has a potent anti-inflammatory and anti-atherogenic effect [12]. Adiponectin is a protein hormone exclusively secreted from adipose tissue. This hormone modulates various metabolic processes, including glucose regulation [11]. Several clinical studies indicate that circulating total adiponectin strongly correlates with markers of insulin resistance. In animal models, adiponectin has an insulin-like function [45, 46]. Here, we show that metformin increase the adiponectin level compared to pretreatment [SMD 0,171 $\mu$ g/ml, 95%CI -0,098 to 0,440, I<sup>2</sup>= 81.09%] (Figure 5). Arita et al. showed that the mean plasma adiponectin levels were 3.7 mg/ml in a group of obese patients, whereas in non-obese subjects these values reached a mean of 8.9 mg/ml [47]. Furthermore, adiponectin levels in patients with type 2 diabetes mellitus were lower than in non-diabetic patients, and were particularly low in subjects with coronary artery disease [48]. The increase of adiponectin may be associated with anti-inflammatory action and decrease of insulin resistance. However, the molecular mechanism to the increase of adiponectin by metformin is unclear. There is an evidence that point out that metformin is able to up-regulate adiponectin gene expression, both in vivo and in vitro, and to stimulate adiponectin protein secretion from human subcutaneous adipose tissue in vitro [13]. Therefore, the main results

of our study were in concordance to previous findings that metformin may increase serum adiponectin levels of diabetic patients.

In summary, our meta-analysis study provides evidence of metformin-induced changes in inflammatory markers and metabolic parameters in diabetic subjects. Taking into account that in the articles analyzed for this meta-analysis the subjects had no disease to alter inflammatory markers level, our findings indicate that metformin monotherapy for more than 4 weeks is able to reduce BMI and inflammatory markers like CRP and TNF $\alpha$  besides increasing adiponectin levels in diabetics. These indicate that this drug has an important clinical effect on inflammation, controlling glucose homeostasis and HbA1c level.

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### **Author contributions**

Conceived and designed the experiments: L.G.P., L.N.B., F.S.E. Collection and tabulation of data: L.G.P., L.N.B., Analyzed the data: L.G.P., L.N.B., Wrote the paper: L.G.P., L.N.B., Supervised the entire project: F.S.E.

### **Declaration of interest**

The authors report no conflicts of interest. The authors alone are responsible for the content and writing of the paper.

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Figures and legends

**Table 1:** Selection criteria for inclusion of studies in the meta-analysis

Inclusion criteria	Exclusion criteria
<i>Study design:</i> Retrospective or prospective study	<i>Study design:</i> Review, meta-analysis, letters to the editor and animal studies
<i>Population:</i> both gender of any age, ethnicity, BMI, with diabetes	<i>Population:</i> Diabetes associated with others disease
<i>data assessment:</i> pre and post metformin treatment	<i>data assessment:</i> No pre and post outcomes
Exposure to monotherapy metformin or metformin and other oral anti-diabetic agents	Healthy control group  Incomplete information data
<i>Outcomes:</i> Blood glucose/HbA1c Inflammatory markers CRP/hsCRP, TNF, Adiponectin, Language: English	Glucose in values normal in pretreatment < 130mg/dL

**Table 1. Data synthesis and Meta-analysis**

**Table 2: Studies characteristic with dose and time of treatment show difference between pre and post BMI.**

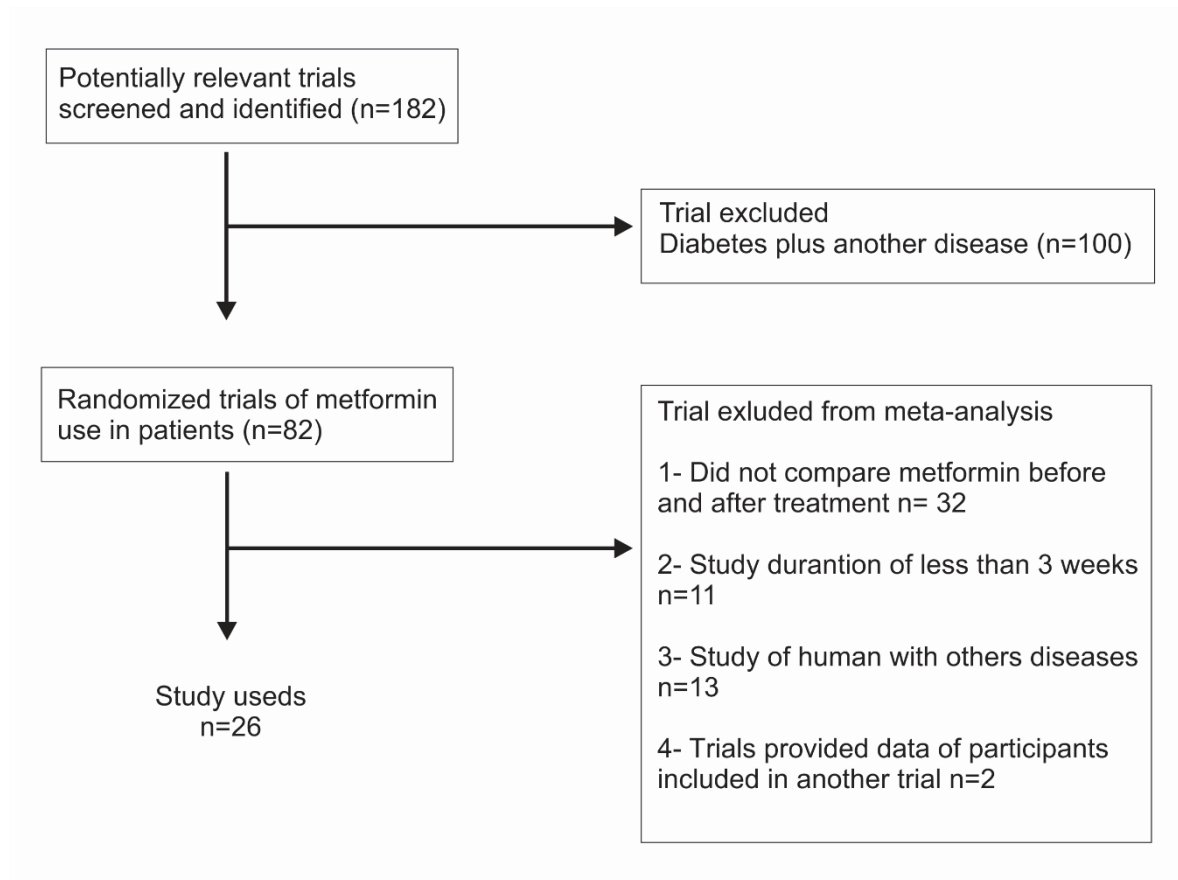
Study	Treatment		Subgroup	dose/day	Time	age		pre % BMI		post % BMI	
	Pre	Post									
Hsieh et al., 2007 <sup>[19]</sup>	24	24	MetforminSR	SR Metformin 1000-2000mg 1/d	14 weeks	58	±1,4	25,30	±0,8	25,00	±0,8
Hsieh et al., 2007 <sup>[19]</sup>	23	23	MetforminRR	RR Metformin 500-1000mg 2/d	14 weeks	56,6	±1,7	25,60	±0,7	25,50	±0,7
Bulcao et al., 2007 <sup>[20]</sup>	21	21	Metformin	Metformin 850mg 2/d	16 weeks	48,6	±9,1	36,10	±0,8	37,30	1,1
Chakraborty et al., 2011 <sup>[21]</sup>	110	110	Metformin	Metformin 850-2000mg/d 1 week and 2/d 2 weeks	4 weeks	42,5	±17,68	27,00	±2,4	23,00	±3,9
de Jager et al., 2005 <sup>[22]</sup>	196	182	Metformin	Metformin 850mg 3/d	16 weeks	63	±9	29,60	±13,9	29,50	±14,1
de Jager et al., 2014 <sup>[23]</sup>	196	131	Metformin	Metformin 850mg 1-3/d	16 weeks	59,1	±11	30	±5	ND	ND
Derosa et al., 2013 <sup>[24]</sup>	85	82	Metformin	Metformin	52 weeks	56,7	±7,3	31,7	±1,5	30,9	±1
Dolasik et al., 2013 <sup>[24]</sup>	60	60	Metformin	Metformin	25 weeks	53	±9	27,8	±3,6	ND	ND
Fidan et al., 2011 <sup>[25]</sup>	20	20	Metformin	Metformin 850 mg 1-3/d	12 weeks	52,6	±7,2	30,1	±5,3	29,3	±5,1
Genovese et al., 2013 <sup>[26]</sup>	29	26	Metformin	Metformin	16 weeks	56,4	±7,9	31,70	±3,6	ND	ND
Glintborg et al., 2014 <sup>[27]</sup>	19	19	Metformin	Metformin	ND	31	±6,36	25,90	±3,9	24,90	±6,6
Kim et al., 2007 <sup>[28]</sup>	56	56	Metformin	Metformin 1000mg 1/d	12 weeks	57,6	±9,4	25,80	±3,3	25,63	±2,3
Kumar et al., 2013 <sup>[29]</sup>	30	30	Metformin	Metformin 500mg 2/3/d	12 weeks	ND	ND	27,23	±2,77	26,86	±12,69
Mao et al., 2009 <sup>[30]</sup>	43	30	Metformin	Metformin 1750mg/d	8 weeks	54,12	±8,86	27	±2,01	26,67	±2,02
Pradhan et al., 2009 <sup>[31]</sup>	126	124	Metformin	Metformin	ND	53,8	±11,5	36,20	±8,1	ND	ND
Stocker et al., 2007 <sup>[32]</sup>	47	38	Metformin	Metformin 850mg 1/d for 2 weeks, 1700mg 1/d for more 2 weeks (4 weeks)	4 weeks	65	±10	29,72	±0,74	ND	ND
Tousoulis et al., 2011 <sup>[33]</sup>	17	17	Metformin	Metformin 850mg 1/d	12 weeks	53,88	±11,06	ND	ND	ND	ND
Derosa et al., 2010 <sup>[34]</sup>	473	175	Metf+Other2	Metformin 2000±500mg + Acarbose	12 weeks	57	±6	26,86	±0,7	26,97	±0,8
Derosa et al., 2010 <sup>[34]</sup>	473	175	Metf+Other1	Metformin 2000±500mg + Pioglitazone	12 weeks	56	±7	26,86	±0,7	26,18	±0,6
Derosa et al., 2007 <sup>[35]</sup>	48	48	Metf+Other1	Metformin + Rosiglitazone 2.250±750mg/d	52 weeks	56	±4	28,10	±1,6	26,00	±1,2
Derosa et al., 2007 <sup>[35]</sup>	48	48	Metf+Other	Metformin + Pioglitazone 2.250±750mg/d	52 weeks	55	±5	28,20	±1,7	26,60	±1,1
Derosa et al., 2010 <sup>[36]</sup>	76	68	Metf+Other	Metformin 850mg + Pioglitazone 15mg 2/d	52 weeks	58	±6	27,7	±1,3	26,7	±0,7
Derosa et al., 2013 <sup>[24]</sup>	85	81	Metf+Other	Metformin + Exenatide	52 weeks	57,3	±7,7	31,9	±1,7	29,6	±0,5

Kumar et al., 2013 <sup>[29]</sup>	30	30	Metf+Other	Metformin + garlic 500mg 2/3/d	12 weeks	ND	ND	27,72	±1,74	25,97	±10,7
Pradhan et al., 2009 <sup>[31]</sup>	126	124	Metf+Other	Metformin + insulin + glargine	ND	54	±11,7	35,60	±7,9	ND	ND
Tousoulis et al., 2011 <sup>[33]</sup>	18	18	Metf+Other	Metformin 850mg/d + Atorvastatin 10mg/d	12 weeks	52,53	±9,57	ND	ND	ND	ND

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ND, non-describe.

**Figure 1.**

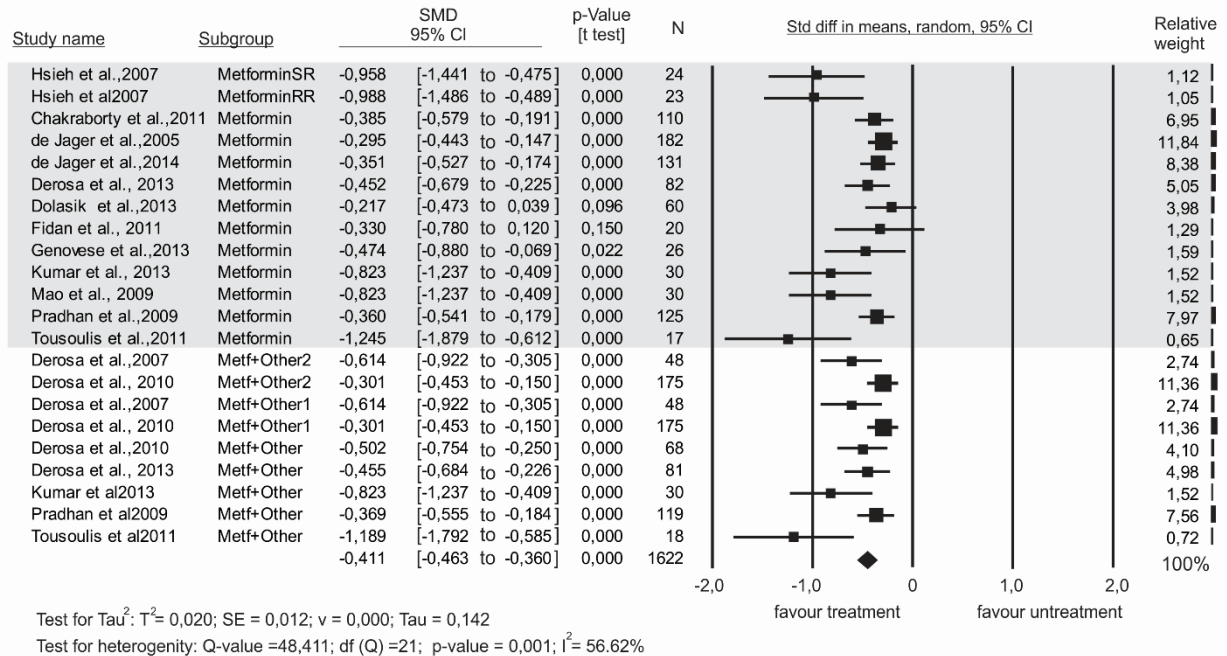


**Figure 1. Flow card of trial research.**

**Figure 2**

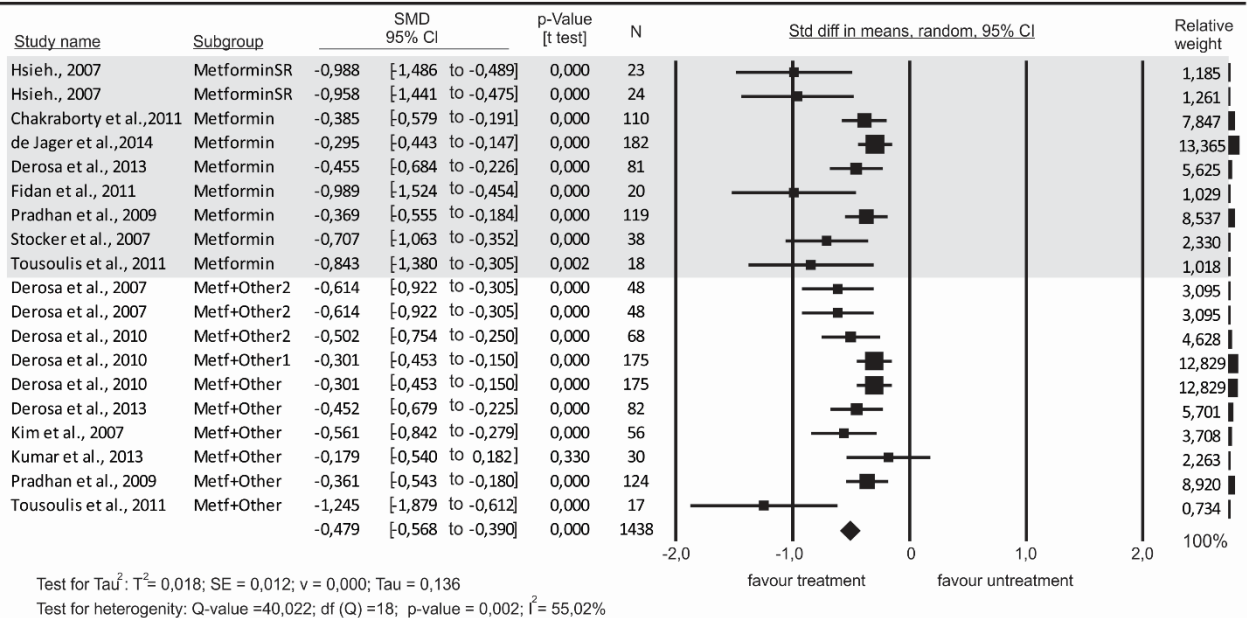
**A**

Outcome: Blood glucose



**B**

outcome: HbA1c



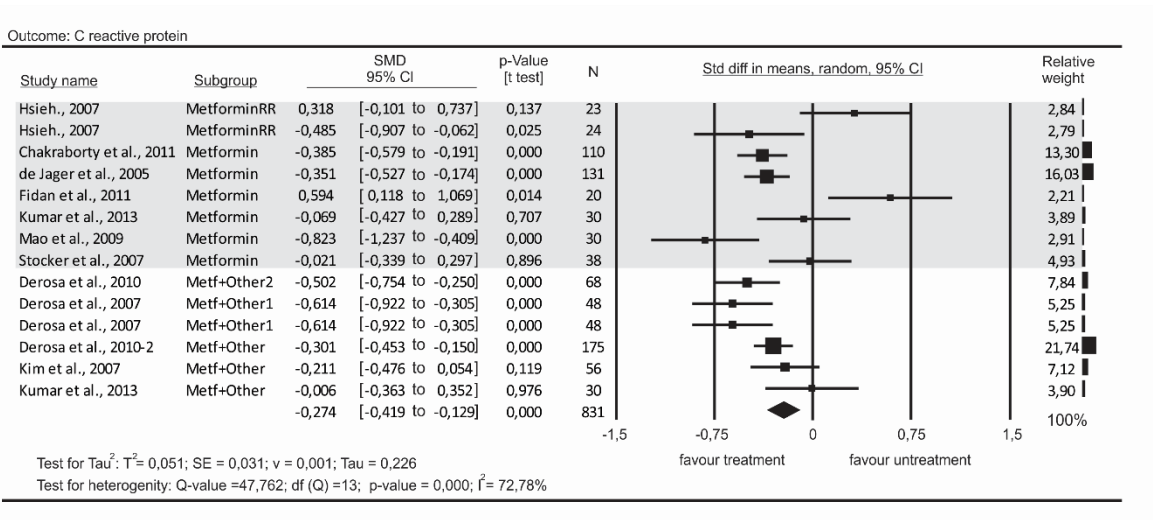
**Figure 2. Effect of post treatment with metformin on blood glucose level and glycated hemoglobin (HbA1c) in diabetic subjects.**

Forest plot from the meta-analysis of metformin on hematological parameters.

Gray frame corresponds to metformin monotherapy and, the white frame corresponds metformin associated with another drug.

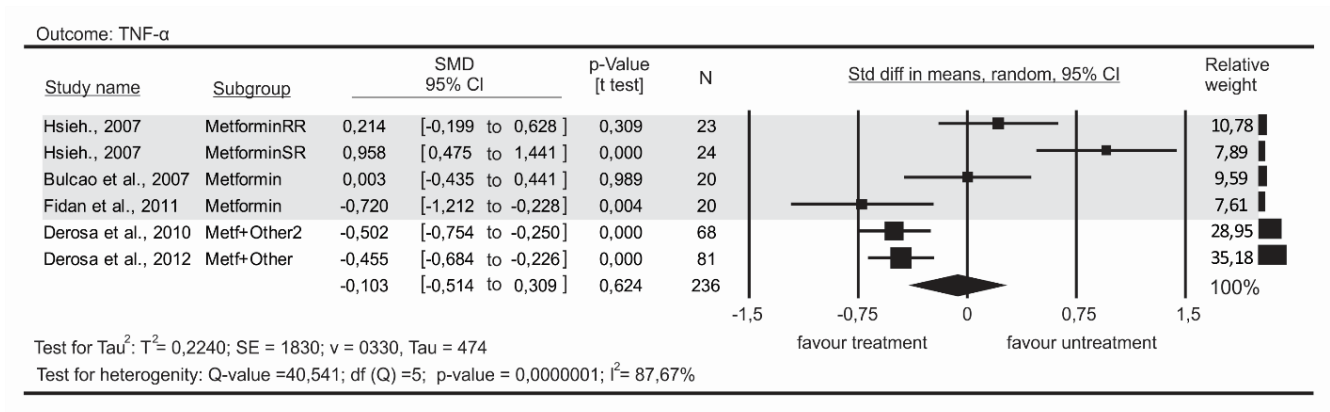


Figure 3.



**Figure 3. Forest plot from the meta-analysis of metformin evaluating C-reactive protein (CRP) in diabetic subjects post metformin treatment.**  
Gray frame corresponds to metformin monotherapy and the white frame corresponds metformin associated with another drug.

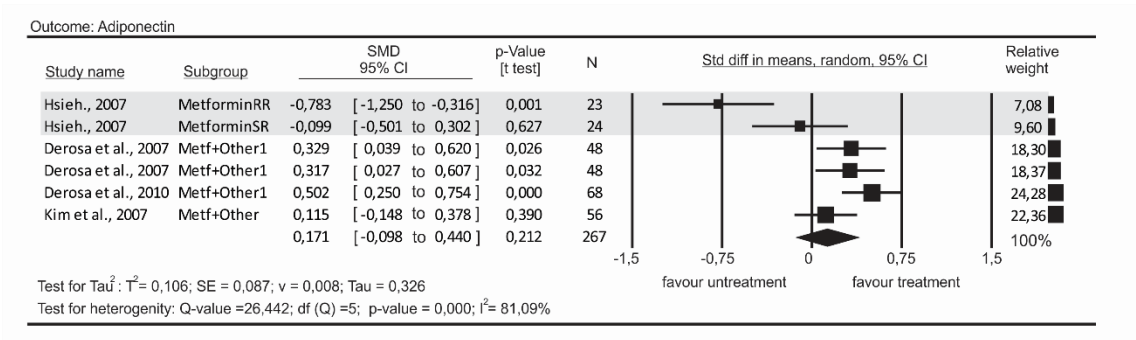
**Figure 4.**



**Figure 4. Forest plot from the meta-analysis of metformin on tumor necrosis factor (TNF $\alpha$ ) in diabetic subjects.**

Gray frame corresponds to metformin monotherapy and the white frame corresponds metformin associated with another drug.

Figure 5.



**Figure 5. Effect of the treatment with metformin on adiponectin in diabetic subjects.**

Gray frame corresponds to metformin monotherapy and the white frame corresponds metformin associated with another drug.

Submits to Journal of Biochemistry & Pharmacology

**Anti-inflammatory and CaMKK $\beta$ /AMPK/Myosin V pathway effects of metformin in skeletal muscle ameliorates insulin resistance in diabetic rats**

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## **Abstract**

**Background:** Metformin increases insulin sensitivity by decreasing hepatic glucose production and increasing glucose disposal in skeletal muscle. However, modulation of inflammatory response and CaMKK $\beta$ /AMPK/Myosin V activation in gastrocnemius muscle by metformin treatment has not been demonstrated in hypoinsulinemic diabetic rats.

**Objective:** The present study investigated how the metformin improve insulin sensitivity in skeletal muscle of hypoinsulinemic diabetic rats.

**Methods:** Diabetes was induced by streptozotocin (45 mg/kg, intraperitoneally) 10 days prior treatments. On 11<sup>th</sup> day, diabetic rats were treated with metformin (500 mg/kg, oral gavage), insulin (2U at 08:00 h and 4U at 17:00 h, subcutaneously) or untreated. After 20 days, glycemia was measured and insulin sensitivity was determined by KITT. Serum Insulin, GLUT4, *IRS $\theta$* , inflammatory markers (NF- $\kappa$ B, I $\kappa$ B, TNF- $\alpha$  and p-JNK) and CAMKK, AMPK and Myosin V in gastrocnemius muscle were determined by ELISA.

**Results:** As expected, insulin and metformin improved the insulin sensitivity. Besides, metformin treatment promoted reduction in inflammatory response mediated by NF- $\kappa$ B, I $\kappa$ B, TNF- $\alpha$  and p-JNK, and that was accompanied by increased CaMKK $\beta$ /AMPK/Myosin V/GLUT4 pathway activity in gastrocnemius muscle of diabetic rats.

**Conclusion:** Our findings suggest that metformin induces significant reductions in several inflammatory markers in skeletal muscle of diabetic rats. Metformin-induced increase in CaMKK $\beta$ /AMPK/Myosin V/GLUT4 pathway activity was associated with higher insulin sensitivity.

## **Key words:**

Inflammatory pathway, insulin sensitivity, insulin pathway, hypoinsulinemic rat.

## Introduction

Skeletal muscle constitutes the largest insulin-sensitive tissue in the body and is the primary site for insulin-stimulated glucose uptake. Skeletal muscle resistance to insulin is fundamental to the development of the metabolic syndrome [1] and diabetes mellitus [2].

It is well documented that higher plasma glucose induces the expression of tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) in skeletal muscle. TNF- $\alpha$  is a potent activator of nuclear factor- $\kappa$ B (NF- $\kappa$ B), which may increase inflammatory cytokines [3-5]. NF- $\kappa$ B is found in all cell types [6-8] and its activity is controlled by regulatory proteins, called inhibitors of NF- $\kappa$ B (I $\kappa$ B) [9]. I $\kappa$ B exists in two phosphorylation states: a hyperphosphorylated state in unstimulated cells, and a hypophosphorylated newly synthesized state in LPS stimulated cells [10]. The phosphorylation of I $\kappa$ B and its proteolytic degradation, allowing the action of NF- $\kappa$ B [9]. Inflammatory mediators lead to phosphorylation of c-Jun-N-terminal kinase (JNK) [5, 11] *which may decrease* insulin sensitivity. However, there are few data in the literature about NF- $\kappa$ B/JNK effects on insulin resistance in skeletal muscle of hypoinsulinemic diabetic rats.

Metformin is an effective hypoglycemic drug [12] that decreases glycemia by decreasing hepatic glucose production and increasing glucose disposal in skeletal muscle [13]. This drug activates AMP-activated protein kinase (AMPK) and decreases NF- $\kappa$ B and TNF- $\alpha$  signaling in endothelial cells [14]. The incubation of isolated epitrochlearis muscles with metformin promotes an increase in the activity of catalytic subunits of AMPK [15]. AMPK activity in vastus lateralis biopsies of subjects with Type 2 Diabetes increased after 4 weeks and 10 weeks of treatment with metformin [13]. In addition, metformin decreases JNK activation and increases insulin receptor substrate (IRS1) concentration in skeletal muscle *in vitro* [16], which can increase translocation of glucose transporter 4 (GLUT4) in cardiac myocytes [17].

Reduced GLUT4 protein expression plays a significant role to promote a decrease in insulin sensitivity of skeletal muscle, which has been described in murine models of insulinopenic diabetes [18]. Furthermore, the improvement in insulin action of skeletal muscle contributes to the therapeutic effects of metformin, mainly resulting in increased non-oxidative glucose disposal in these cells [19]. Myosin V, an unconventional motor protein family, participates in GLUT4 translocation in L6 muscle cells [20]. However, at this time, the myosin V expression induced by metformin treatment has not been evaluated. Although there is a lot of evidence of the role of metformin in intracellular signaling pathway, it is unclear how this drug improves insulin sensitivity in skeletal muscle of hypoinsulinemic diabetic rats. In this report we tested the hypothesis that activation of CaMKK $\beta$ /AMPK/Myosin V pathway in skeletal muscle cells mediates the beneficial metabolic effects of metformin. The present study was also designed to determine the role of metformin in the regulation of inflammatory response in skeletal muscle of hypoinsulinemic diabetic rats.

## Material and Methods

### *Animals*

Male Wistar rats (12 weeks, 181–223g) were maintained under standard conditions ( $22\pm 1^{\circ}\text{C}$ , humidity  $60\pm 5\%$ , 12 h light/dark cycle) with food and water *ad libitum*. All procedures for handling, use and euthanasia of the animals followed the guidelines proposed by the Brazilian Society of Laboratory Animal Science and by the Ethics Committee for Animal Research of the Federal University of Uberlandia, Brazil (CEUA/UFU 053/13).

### *Induction of experimental diabetes*

Animals were fasted 24h and rendered diabetic by a single injection of streptozotocin (STZ) ( $60\text{ mg kg}^{-1}$ , intraperitoneally). STZ was dissolved in 0.01 M citrate buffer (pH 4.5). Non-diabetic rats received a placebo dose (0.01 M citrate buffer). After 10 days, the fasting plasma glucose level was monitored by tail vein puncture using reactive strips (Biocheck Glucose Test Strip; Bioeasy, Belo Horizonte, MG, Brazil). Diabetes was defined as non-fasting glycaemia  $>250\text{ mg/dL}$ .

### *Groups and treatments*

The animals were randomly divided into four groups ( $n=12$  rats/group). Non-Diabetic (ND), Diabetic (D), Diabetic rats treated with NPH insulin - 6U/day divided in two subcutaneous injections: 2U at 08:00 h and 4U at 17:00h (D+I), and Diabetic rats treated with oral metformin -  $500\text{ mg kg}^{-1}\text{day}$  by gavage (D+M). Treatments were performed for 20 days. After that period, rats were anesthetized with xylazine ( $10\text{ mg kg}^{-1}$ ) and ketamine ( $100\text{ mg kg}^{-1}$ ) for blood collection and euthanized for the removal of gastrocnemius skeletal muscle.

### *Intravenous insulin tolerance test*

Tail blood samples were collected from animals fed *ad libitum* before (0 min) and 4, 8, 12, and 16 min after i.v. injection of regular insulin ( $0.75\text{ U kg}^{-1}\text{ BW}$ ). The constant rate of plasma glucose disappearance during insulin tolerance test (KITT) was calculated based on a linear regression of the Neperian logarithm of the plasma glucose concentrations (test strips, Advantage, Roche). The tests were performed from 09:00 to 11:00 h [21].

### *Muscle homogenization*

Gastrocnemius muscle (0.1 g, removed before intravenous insulin tolerance test) was homogenized in buffer (25 mM Tris-HCl pH 7.4, cocktail protease inhibitor, 10 mM EDTA and 10 mM EGTA) for 3 min on ice. The homogenate was centrifuged at  $15.000\times g$  for 40 min. at  $4^{\circ}\text{C}$ . The pellet was resuspended in the same homogenization buffer. The supernatant was used for following analysis (GLUT4, CAMKK $\beta$ , p-CAMKK $\beta$ , IRS 1 thr, AMPK, p-MAPK, NF- $\kappa$ B, I $\kappa$ B, p-JNK) and pellet to (Myosin V) by *Western blotting* (WB) and ELISA methods.

## ELISA

Gastrocnemius muscles from all groups were diluted to a final concentration of  $0.1 \mu\text{g mL}^{-1}$  in coating buffer (100 mM bicarbonate/carbonate, pH 9.6). The microplate wells were coated with these samples ( $100 \mu\text{L} - 1 \mu\text{g}$  total protein well<sup>-1</sup>) overnight at  $4^{\circ}\text{C}$  and washed 3 times with PBS-T. Then, the wells were blocked with  $200 \mu\text{L}$  of blocking buffer (2% BSA in PBS-T) for 4h at RT.  $100 \mu\text{L}$  well<sup>-1</sup> of the antibodies (anti-GLUT4, anti- CAMKK $\beta$ , anti- CAMKK $\beta$  phosphorylate, anti-IRS1 tyrosine, anti-AMPK, anti-AMPK phosphorylate, anti-myosin V, anti-NF- $\kappa\text{B}$ , anti-I $\kappa\text{B}$ , anti-JNK phosphorylate) diluted 1:1000 were added for 2h at RT and washed 3 times with PBS-T.  $100 \mu\text{L}$  well<sup>-1</sup> of RP-conjugated antibodies diluted 1:5000 were added for 2h at RT and washed 3 times with PBS-T. The secondary antibody (rabbit and mouse – conjugated HRP) was detected by adding  $100 \mu\text{L}$  of the OPD-substrate at 20 min and read at 490 nm.

## SDS–PAGE and Western blotting

The *Western blotting* was used to confirm the protein expression. Protein samples ( $10 \mu\text{g}$ ) were applied to 5–22% polyacrylamide gradient gels under denaturing conditions [22] and stained with Coomassie brilliant blue. For immunoblotting, proteins were transferred to nitrocellulose membranes in Tris–glycine buffer[22]. The membranes were incubated with 5% dried milk in Tris-buffered saline (TBS-T) (50 mM Tris–HCl, pH 8.0, 150 mM NaCl, 0.05% Tween 20) and probed with antibodies (anti- CAMKK $\beta$ , anti- CAMKK $\beta$  phosphorylate, anti-IRS1 tyrosine, anti-AMPK, anti-AMPK phosphorylate, anti-myosin V, anti-NF- $\kappa\text{B}$ , anti-I $\kappa\text{B}$ , anti-JNK phosphorylate) diluted to  $0.2 \mu\text{g/mL}$  in TBS-T. The reaction was detected using an ECL kit (Amersham Biosciences, Buckinghamshire, UK) according to the manufacturer's instructions, and Amersham Imager 600 was used for exposure. The intensities of the protein bands were analyzed and compared using the ImageQuant TL software, version 8.1 (GE Healthcare Life Science).

## Insulin measurement

Plasma insulin was analyzed by an enzyme immunoassay (EIA) using an insulin (mouse/rat) EIA kit (Bertin Pharma, Montigny-le-Bretonneux, France).

## Statistical analysis

Prior to the statistical analysis, the data were tested for normality using the Shapiro-Wilk test. For the statistical analysis, data were normalized to the non-diabetic group, and the mean of each group was compared with the mean of the non-diabetic rats (positive control) and diabetic rats (negative control), using ANOVA followed by a Bonferroni test for multiple comparison. A Student's *t* test was used to compare the values of the intravenous insulin tolerance test of diabetic rats and diabetic rats treated with metformin. The significance level for all analyses was  $p < 0.05$ . The results are shown as the means  $\pm$  SEM.



## Results

Glycemic characteristics of the animals are shown in Figure 1. Compared with non-diabetic rats (ND), diabetic (D) and metformin-treated diabetic (D+M) rats presented hyperglycemia ( $p<0.001$ ). As expected, insulin treatment reduced ( $p<0.05$ ) plasma glucose in diabetic rats. The metformin treatment did not change ( $p>0.05$ ) plasma glucose in diabetic rats during removal of samples (Figure 1.A). For analysis of acute metformin treatment in diabetic rats (Figure 1.B), plasma glucose was evaluated 2h, 4h, 6h and 12h after a single administration by oral gavage. Metformin treatment of diabetic (D+M) rats induced a similar plasma glucose reduction ( $p<0.05$ ) from 2h to 6h of treatment (Figure 1B).

As expected, the insulin sensitivity in diabetic (D) rats was lower ( $p<0.05$ ) than non-diabetic (ND) rats. In addition, the insulin sensitivity was higher in both insulin-treated diabetic (D+I) and metformin-treated diabetic (D+M) rats, compared to diabetic (D) rats (Figure 2.A). Figure 2.B shows that diabetes decreased ( $p<0.05$ ) the GLUT4 protein expression in gastrocnemius muscle. Insulin- (D+I) and metformin-treated (D+M) rats were able to increase ( $p<0.01$ ) the GLUT4 protein expression as compared to diabetic condition level.

Low plasma insulin levels ( $p<0.05$ ) were observed in diabetic (D) and metformin-treated diabetic (D+M) rats compared to non-diabetic (ND) rats. Increased insulin levels ( $p<0.05$ ) were found in insulin-treated diabetic (D+I) rats compared with non-diabetic (ND) rats (Figure 3.A). The diabetes did not change *IRS1<sup>thr</sup>* expression in gastrocnemius muscle. As expected, chronic (20 days) treatment of insulin resulted in the upregulation of *IRS1<sup>thr</sup>* in gastrocnemius muscle of diabetic rats as compared to non-diabetic (ND) and to diabetic (D) rats. Furthermore, the chronic (20 days) metformin treatment also increases ( $p<0.05$ ) *IRS1<sup>thr</sup>* expression in this tissue compared to diabetic (D) rats (Figure 3.B).

To assess the role of insulin and metformin treatment on the activation of CAMKK $\beta$ /AMPK/Myosin V pathway under diabetic conditions, these proteins were analyzed in gastrocnemius muscle. In diabetic (D) rats CAMKK $\beta$  phosphorylation was decreased ( $p<0.05$ ) compared to the non-diabetic (ND) rats. Compared with non-diabetic rats (ND), 20-day diabetic rats (D) showed no significant changes in AMPK phosphorylation and myosin V content. After 20 days of insulin treatment, the effect of diabetes on CAMKK $\beta$  phosphorylation and myosin V content were reversed in gastrocnemius muscles. Compared with non-diabetic (ND) and diabetic (D) rats, insulin-treated diabetic (D+I) rats showed no significant changes in AMPK phosphorylation (Figure 4.A-4.C). As shown in Fig. 4., metformin treatment increased CAMKK $\beta$  phosphorylation ( $p<0.05$ ), AMPK phosphorylation ( $p<0.05$ ) and myosin V protein content ( $p<0.05$ ) than non-diabetic (ND) and diabetic (D) rats.

To determine the kinetics of inflammatory response, we next tested if the insulin and metformin treatment would impact NF- $\kappa$ B and other related factors like I $\kappa$ B, TNF- $\alpha$  and p-JNK in skeletal muscle of diabetic rats (Figure 5.A-5.D). We found that diabetic (D) animals had higher ( $p>0.05$ ) NF- $\kappa$ B and TNF- $\alpha$  expression compared to non-diabetic (ND) animals. Besides, the JNK phosphorylation was also increased ( $p<0.05$ ) in diabetic (D) rats compared to non-diabetic (ND) rats. In gastrocnemius muscle of insulin-treated diabetic (D+I) rats, decreased ( $p<0.05$ ) NF- $\kappa$ B expression, TNF- $\alpha$  expression and JNK phosphorylation were observed as

compared to diabetic (D) rats. I $\kappa$ B expression was increased ( $p < 0.05$ ) in insulin-treated diabetic (D+I) rats as compared to diabetic (D) rats. Metformin-treated diabetic (D+M) rats overexpressing ( $p < 0.05$ ) the NF- $\kappa$ B inhibitory protein, I $\kappa$ B, associated with reduction ( $p < 0.05$ ) in NF- $\kappa$ B protein expression than non-diabetic (ND) rats. TNF- $\alpha$  expression and JNK phosphorylation significantly decreased ( $p < 0.05$ ) in metformin-treated diabetic rats (D+M) than diabetic (D) rats.

## Discussion

The present study investigated the effects of metformin or insulin and their association with CaMKK $\beta$ /AMPK/Myosin V/GLUT4 pathway and inflammatory response in gastrocnemius muscle of hypoinsulinemic diabetic rats. To the best of our knowledge, the CaMKK $\beta$ /AMPK/Myosin V/GLUT4 pathway associated with inflammatory markers (NF- $\kappa$ B, I $\kappa$ B, TNF- $\alpha$  and p-JNK) has never been directly measured in hypoinsulinemic diabetic rats despite its critical role in regulating insulin resistance, which is linked to the development of the diabetes. The present results show evidence of increased CaMKK $\beta$ /AMPK/Myosin V/GLUT4 pathway activity in metformin-treated diabetic rats compared with diabetic rats. Furthermore, metformin reduced the inflammatory response mediated by NF- $\kappa$ B, I $\kappa$ B, TNF- $\alpha$  and p-JNK in skeletal muscle of diabetic rats. These effects were associated with reduction in insulin resistance.

The terminal half-life of metformin after oral administration occurs in 6h - 7h at doses of 100 and 200 mg/kg, respectively [23]. These data are consistent with our findings which indicate that metformin decreases glycaemia between 2 to 6hs after the oral administration, returning to baseline blood glucose after 12h. In the present study, the diabetic state of the animals was similar to that described in previous studies, and effect of insulin upon glycaemic control was also similar [24-26]. In addition, we showed that metformin treatment increases the plasma glucose disappearance rate (KITT) compared to non-treated diabetic rats indicating an improving the insulin sensitivity.

Metformin has been used in different pathologies [27-30], but, its major application is to decrease insulin resistance in the liver [31]. On the other hand, skeletal muscle is the predominant site of insulin-mediated glucose uptake. Therefore, the reduction of damage caused by proinflammatory cytokines in this tissue is essential for glucose homeostasis. Diabetes is associated with systemic inflammation and metabolic disorders [32, 33], increasing the level of TNF- $\alpha$  [5, 32, 33] and decreasing the GLUT4 translocation [34, 35]. It is well documented that TNF- $\alpha$  induce insulin resistance in skeletal muscle [5] by increase of phosphorylation of JNK and IRS1 ser. Studies show that metformin decrease TNF- $\alpha$  via AMPK activation and inhibit NF- $\kappa$ B activation in vascular endothelial cells [36, 37]. Here, we have found that metformin have a similar mechanism of action in skeletal muscle of hypoinsulinemic diabetic rats. Our findings suggest that treatment with metformin improve insulin sensibility by maintenance of NF- $\kappa$ B/I $\kappa$ B complex, decrease of TNF- $\alpha$  expression, JNK phosphorylation associated with the activation of CaMKK $\beta$ /AMPK pathway in skeletal muscle. This could indicate that metformin regulates proinflammatory pathway in skeletal muscle by activation of

CAMKK $\beta$ /AMPK pathway. However, metformin is not specific AMPK activators. This drug activate AMPK by the upstream kinases serine/threonine kinase 1 (LKB1) in liver. The deletion of LKB1 in the liver did not impair AMPK activation in culture of muscle tissue [38], but deficiency of LKB1 in skeletal muscle prevents AMPK activation even in the presence of the AMP-mimetic agent, 5-aminoimidazole-4-carboxamide riboside (AICAR), the antidiabetic drug phenformin, or by muscle contraction [39]. Higher metformin dose induce AMPK phosphorylation independent of CAMKK $\beta$  in arterial smooth muscle [40]. On the other hand, other studies show that CAMKK $\alpha$ , regulates glucose uptake independent of AMPK and Akt activation, and that the CAMKK $\beta$  has a critical role in the regulation of contraction-induced glucose uptake in mouse skeletal muscle [41]. CAMKK $\beta$  activate AMPK in response to increase of intracellular calcium [42]. So we suggest that metformin can activate AMPK both as CaMKK independently.

Furthermore, intracellular calcium promotes GLUT4 translocation by CAMKK $\beta$  and AMPK activation [43]. Thus, CAMKK activity depends of the calcium intracellular level and can activate AMPK independently of changes in AMP [44]. Although we did not evaluate the release of calcium from the sarcoplasmic reticulum, we show for the first time an increase of CAMKK $\beta$  in skeletal muscle of rats treated with metformin. More studies are needed to understand how metformin activates the CAMKK $\beta$  in skeletal muscle. Additionally, we show that metformin decrease p-JNK and increase CaMKK $\beta$ /AMPK/Myosin V pathway IRS1 $thr$ . Our results is consistent with a previous study which suggested that metformin decrease JNK activation and increase IRS1 concentration in skeletal muscle *in vitro* [16]. Furthermore, to our knowledge, we also demonstrate for the first time that metformin treatment increase myosin V expression in skeletal muscle of hypoinsulinemic diabetic rats. The molecular mechanism involved in this process is due to activation of AS160 by p-AMPK that can occurs independently of the insulin signal [45]. In addition, study shows the co-localization of Myosin V and RAB8a in the GLUT4 vesicle exocytosis of muscle cells [46]. These finding indicated that metformin treatment is able to increase expression of myosin V involved in GLUT4 translocation independent of the insulin signal. Thus, we propose a metformin-induced pathway leading to GLUT4 translocation into the plasma membrane of muscle cells in gastrocnemius muscle (Fig. 6).

In summary, we show that metformin induces significant reductions in several inflammatory markers as NF- $\kappa$ B, I $\kappa$ B, TNF- $\alpha$  and p-JNK of skeletal muscle in hypoinsulinemic diabetic rats. Additionally, in this tissue, metformin treatment was followed by increased CaMKK $\beta$ /AMPK/Myosin V/GLUT4 pathway. The degree of pro-inflammatory markers and CaMKK $\beta$ /AMPK/Myosin V/GLUT4 pathway activation in skeletal muscle cells are related with the insulin sensitivity in metformin-treated diabetic rats compared with diabetic rats. Collectively, these novel findings may provide a significant step toward the elucidation of the metabolic and inflammatory signaling pathways regulating metformin-dependent skeletal muscle glucose uptake.

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## **Author contributions**

Conceived and designed the experiments: L.G.P., R.R.T., F.S.E. Performed the experiments: L.G.P., R.R.T., N.B.B., H.L.M., D.C.C., Analyzed the data: L.G.P., R.R.T., D.D.V., R.S.S. Wrote the paper: L.G.P., R.R.T., Participated in interpretation and revised the manuscript for publication: RS-S. Supervised the entire project: F.S.E.

## **Declaration of interest**

The authors report no conflicts of interest. The authors alone are responsible for the content and writing of the paper.

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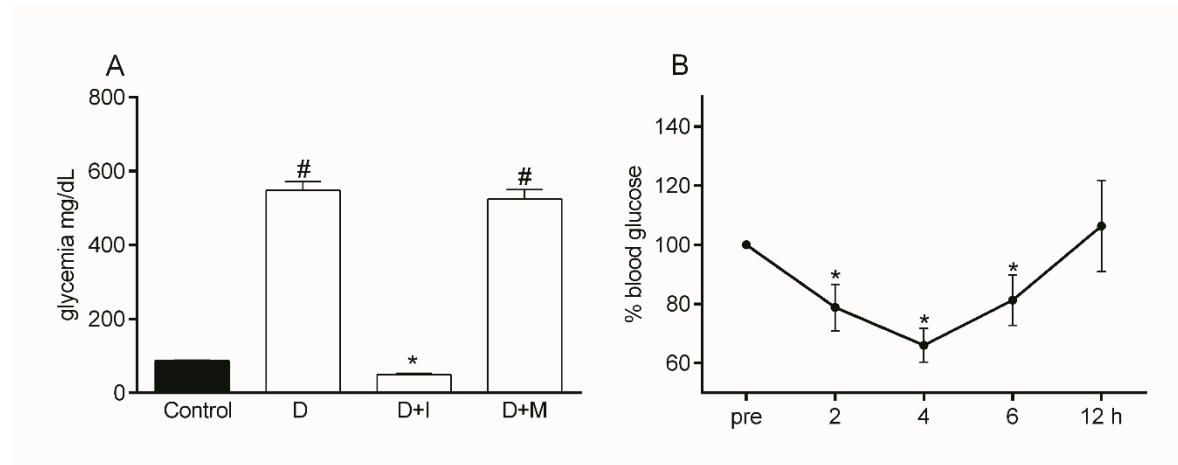
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## Figures and Legends

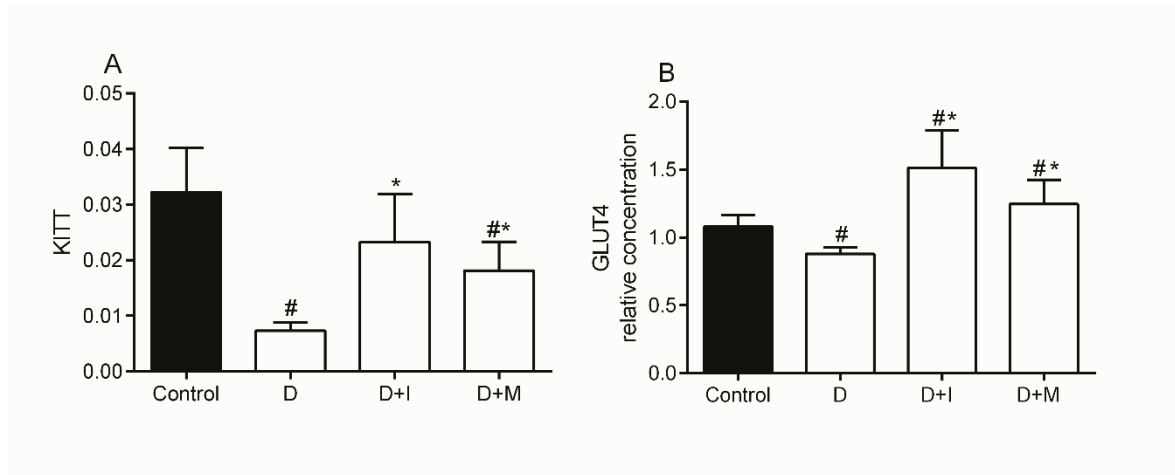
**Figure 1.**



**Figure 1. Fasting plasma glucose in non-diabetic (ND), untreated diabetic (D), and insulin (D+I)- or metformin (D+M)- treated diabetic rats.**

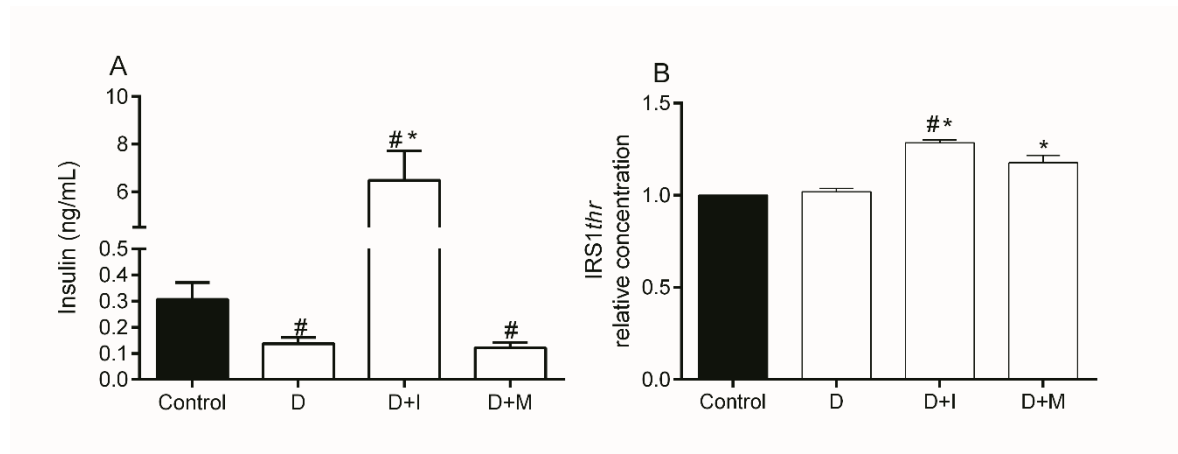
(A) Blood glucose levels. (B) Daily blood glucose variation in rats treated with metformin. Data expressed as mean  $\pm$  SEM; # vs. non-diabetic; \* vs. diabetic;  $p < 0.05$ ;  $n = 8-10$  per group.

**Figure 2.**



**Figure 2. Insulin sensitivity (A) and GLUT4 expression in gastrocnemius (B) muscles of on-diabetic (ND), untreated diabetic (D), and insulin (D+I)- or metformin (D+M)- treated diabetic rats.**  
Data expressed as mean  $\pm$  SEM; # vs. non-diabetic; \* vs. diabetic;  $p < 0.05$ ;  $n = 8-10$  per group.

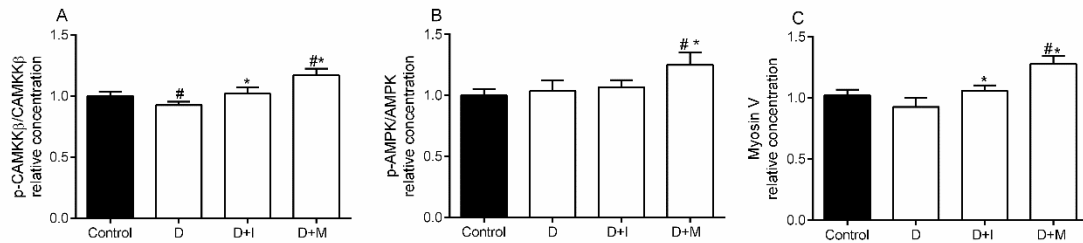
**Figure 3.**



**Figure 3. Basal plasma insulin and IRS1thr expression in gastrocnemius (B) muscles of on-diabetic (ND), untreated diabetic (D), and insulin (D+I)- or metformin (D+M)- treated diabetic rats.**

Data expressed as mean  $\pm$  SEM; # vs. non-diabetic; \* vs. diabetic;  $p < 0.05$ ;  $n = 8-10$  per group.

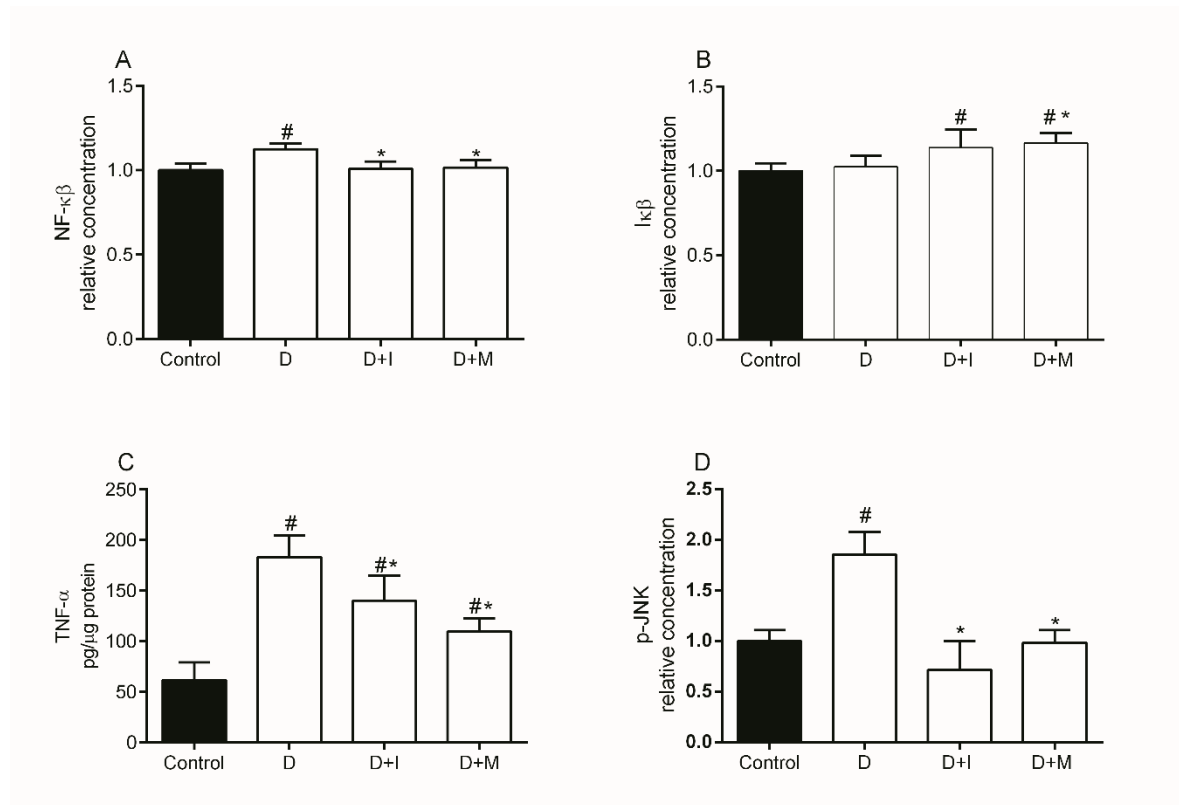
**Figure 4.**



**Figure 4. Intracellular signal triggered by metformin action.**

(A) p-CAMKK $\beta$ /CAMKK $\beta$  concentration; (B) p-AMPK/AMPK concentration; (C) Myosin V concentration. Non-diabetic (ND), untreated diabetic (D), and insulin (D+I)- or metformin (D+M)-treated diabetic rats. Data expressed as mean  $\pm$  SEM; # vs. non-diabetic; \* vs. diabetic;  $p < 0.05$ ;  $n = 8-10$  per group.

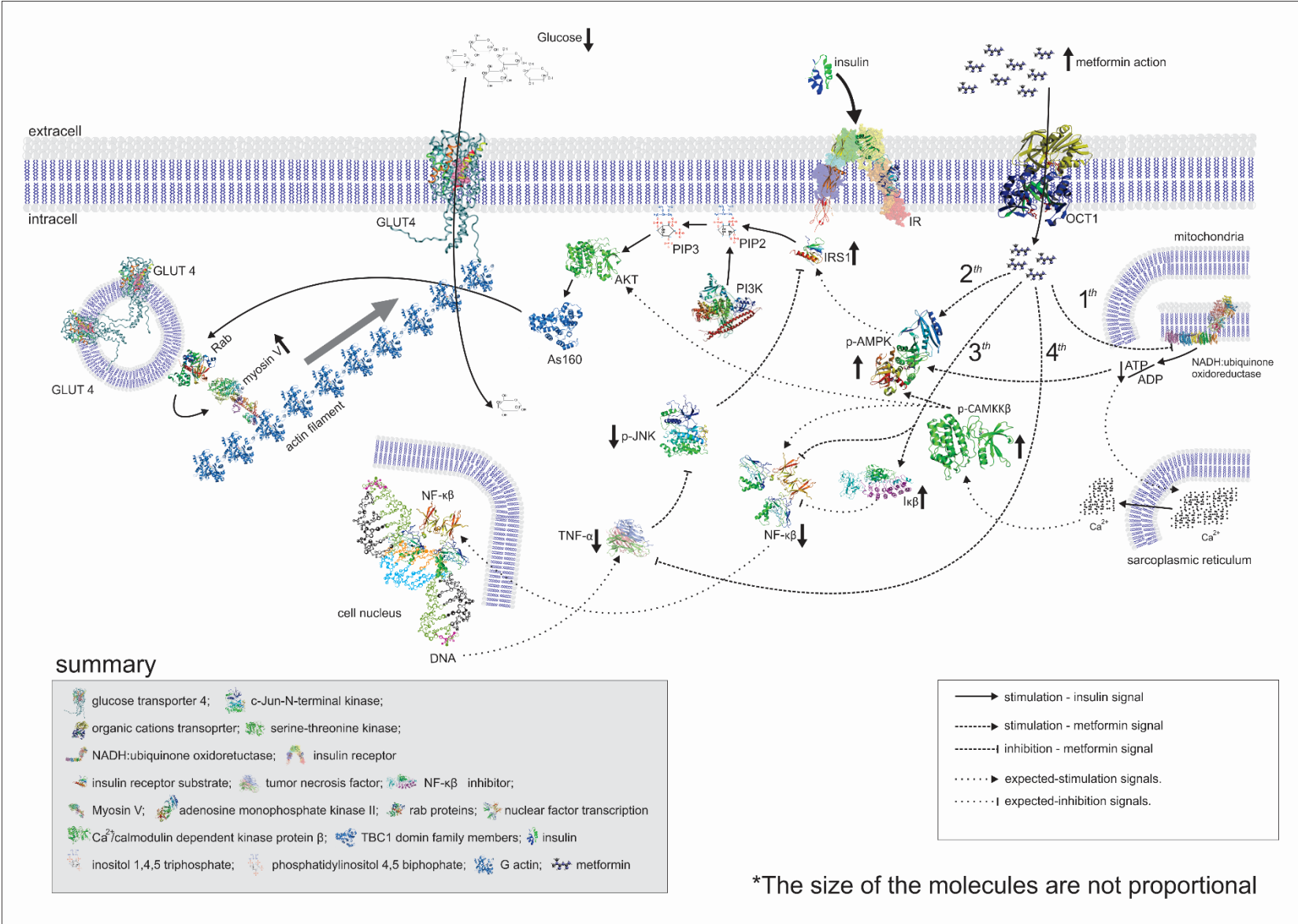
**Figure 5.**



**Figure 5. Metformin role on inflammatory pathway.**

(A) NF- $\kappa$ B concentration; (B) I $\kappa$ B concentration; (C) TNF- $\alpha$  concentration; (D) p-JNK concentration. Non-diabetic (ND), untreated diabetic (D), and insulin (D+I)- or metformin (D+M)-treated diabetic rats. Non-diabetic (ND), untreated diabetic (D), and insulin (D+I)- or metformin (D+M)-treated diabetic rats. Data expressed as mean  $\pm$  SEM; # vs. non-diabetic; \* vs. diabetic;  $p < 0.05$ ;  $n = 8-10$  per group.

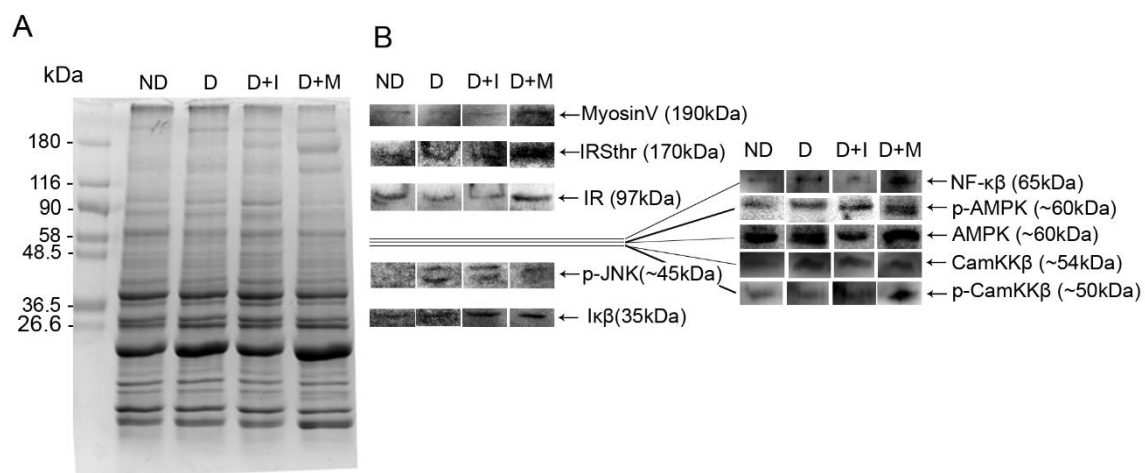
Figure 6.



**Figure 6: Metformin role on skeletal muscle.**

Schematic draw of a proposed pathway for the molecular mechanism of metformin on skeletal muscle. Arrows indicate the modulations (↑ increase, ↓ decrease) of each parameter in insulin (D+I)- or metformin (D+M)-treated diabetic rats. Scheme shows the molecular mechanism of metformin on skeletal muscle by different signaling pathways: 1<sup>th</sup> inhibition of complex I mitochondrial, 2<sup>th</sup> activation AMPK, 3<sup>th</sup> maintenance of NF-κB/IκB complex and 4<sup>th</sup> inhibition of TNFα. Indirectly, the metformin active AMPK and CaMKKβ, which are regulatory points. In addition, metformin increases IκB and decrease NF-κB concentration and consequently inhibits TNFα expression and p-JNK concentration. Furthermore, metformin increases *IRS<sup>thr</sup>*, Myosin V and GLUT4 concentrations. Taken together, metformin improve glucose uptake decrease inflammatory pathway and increase GLUT4 translocate.

## Supplementary figure



### Supplementary figure.

**Representative figure of inflammatory signaling pathways induced by hyperglycemia of gastrocnemius muscle of hypoinsulinemic diabetic rats.** A- SDS-PAGE 5-22% stained with coomassie blue (b- before; a- after). B- WB of inflammatory pathway. Antibody uses: anti-myosin V, anti-IRSthr, anti-IR, anti-NF- $\kappa$ B, anti-p-AMPK, anti-AMPK, anti, p-JNK, anti-CAMKKII, anti-p-CAMKKII and, anti-I $\kappa$ B.



Submits to Journal of Diabetic Research

**The effects of metformin on inflammatory and apoptotic pathways in soleus muscle of diabetic rats.**

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Running title: Anti-inflammatory mechanism of metformin in soleus muscle

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## **Abstract**

Diabetes is characterized by a proinflammatory state which can activate TLR2 and TLR4, and these receptors could induce NF- $\kappa$ B and JNK activation in skeletal muscle. In this study, we investigated the inflammatory and apoptotic signaling pathways triggered by TLRs/NF- $\kappa$ B and JNK activation in skeletal muscle of diabetic rats treated with metformin before and after an insulin tolerance test. Metformin treatment decreased p-JNK and NF- $\kappa$ B, and increased I $\kappa$ B concentrations. This attenuation leads to a decrease of TNF $\alpha$  and CXCL1/KC, and an increase of p-AMPK, BAX and Bcl2 concentration. Furthermore, KITT revealed an improvement of the insulin sensitivity in the diabetic rats treated with metformin. In addition, metformin was not capable of attenuating the changes in the inflammatory pathway triggered by insulin injection as the increase of TNF $\alpha$  and TLR4 in metformin treated rats, and I $\kappa$ B, CXCL1/KC, TNF $\alpha$  and p-AMPK increase in the untreated group. Taken together, these results point out that metformin may attenuate the activation of the inflammatory pathway TLRs/NF- $\kappa$ B/TNF $\alpha$ /CXCL1/KC and the apoptotic signaling BAX/Bcl2/p-JNK, which could be accompanied by a reduction of the inflammatory damage caused by hyperglycemia and an improvement of insulin sensitivity in diabetic rats.

## **Key words:**

Insulin resistance, toll like receptor, NF- $\kappa$ B/I $\kappa$ B pathway, skeletal muscle

## Introduction

Diabetes is characterized by a proinflammatory state and high blood glucose levels induces the activation of several inflammatory pathways [1-3], besides being associated with disorders of the immune system [4]. Inflammation is pivotal in the development of insulin resistance in part via the activation of the toll-like receptors (TLRs), as TLR4 and TLR2 [5, 6]. TLR4 and TLR2 expression are increased in diabetes [7] and the activation of these receptors contributes to insulin resistance in muscle [8].

Upon activation, TLR4 and TLR2 recruit the transcription factor nuclear factor- $\kappa$ B (NF- $\kappa$ B) and the c-Jun NH2-terminal kinase (JNK), which initiates the transcription of proinflammatory cytokine genes [9-12], as chemokine ligand 1 (CXCL1/KC) and tumor necrosis factor (TNF $\alpha$ ). These chemokines are involved with decreased insulin sensitivity [13]. In skeletal muscle, both JNK [14] and NF- $\kappa$ B [15, 16] signal transduction pathways impair insulin signaling. In addition, JNK may induces the phosphorylation of serine residues in insulin receptor substrate (IRS-1) [17-19] while NF- $\kappa$ B induces, by transcriptional activation, the production of proinflammatory cytokines, including TNF $\alpha$ , IL-6, IL-1 $\beta$ , and resistin[20, 21]. These cytokines further activates JNK and NF- $\kappa$ B pathways through a feed-forward mechanism [22].

JNK and NF- $\kappa$ B are activated and inhibited by different mechanisms. Classically, JNK can be activated by both MKK4 and MKK7. These enzymes can phosphorylate Thr-Pro-Tyr residues induced by several stress stimuli and it is inhibited by JNK interacting protein (JIP) [23]. NF- $\kappa$ B in turn, is activated by the phosphorylation of inhibitor  $\kappa$ B (I $\kappa$ B), which liberates NF- $\kappa$ B to translocate into the cell nucleus stimulating the transcription of inflammatory mediators[24]. The lack of inhibition of these molecules can stimulate apoptotic processes, otherwise trigger inflammation. However, the mechanism of the interactions among these mediators remains unclear.

Evidence indicates that both JNK and NF- $\kappa$ B also participate of pro-apoptotic and anti-apoptotic pathways, respectively. JNK activation also promotes BAX translocation to mitochondria [25] and the phosphorylation of Bcl2, suppressing its function [26]. NF- $\kappa$ B also activates Bcl-2 expression [27]. BAX and Bcl2 are members of Bcl-2 family of proteins, which can be divided into pro-apoptosis (BAX, Bid, Bad, Bak, Bcl, Bok, Bcl-XS, and Hrk) or anti-apoptosis (Bcl2, Bcl-XL, Bcl-W, A1 and Mcl-1) proteins [28].

Metformin is a drug that has a hypoglycemic action, broadly used in the treatment of diabetes, and with a variety of other applications[29]. For example, in the inflammatory pathway, metformin inhibited the nuclear translocation of NF- $\kappa$ B and the phosphorylation of STAT3 in cancer stem cells [30]. Moreover, this drug promotes the inhibition of TNF $\alpha$ -induced I $\kappa$ B kinase phosphorylation and IL-6 production in endothelial cells through PI3K-dependent AMP-activated protein kinase (AMPK) phosphorylation [31, 32] as well as, suppresses TLRs of the myocardial infarction in rat [33]. In apoptotic signal, metformin activates AMPK,

which stimulates JNK1–Bcl-2 signaling and disrupts the Beclin 1–Bcl-2 complex, normalizing the cardiac autophagy and attenuating high glucose–induced apoptosis in H9c2 cultured cells [34]. These findings suggest that metformin may attenuate the inflammatory damage caused by hyperglycemia and improve insulin sensitivity in skeletal muscle.

Thus, the aim of the present study was to investigate the metformin role on both inflammatory and apoptotic signaling pathway in skeletal muscles of streptozotocin-diabetic rats before and after insulin tolerance test.

## **Material and methods**

### *Animals*

Male Wistar rats (7 weeks and ~230 g of body weight) were kept in standard conditions (22±1 °C, humidity 60%±5 and 12h light/12h darkness cycle) with *ad libitum* food and water. The procedures for handling and use of animals followed the resolutions proposed by the Brazilian Society of Laboratory Animal Science and by the Ethics Committee for Animal Research of the Federal University of Uberlandia, Brazil (CEUA / UFU 107/14).

### *Diabetes Induction*

Animals were fasted 12 hours and rendered diabetic by a single intraperitoneal injection of streptozotocin (STZ) (45 mg/kg BW). Diabetes was defined as a non-fasting glucose > 250 mg/dL in tail vein blood (Accu-Chek Performa) 10 days after STZ injection.

### *Groups and treatment*

Diabetic animals were divided randomly into 2 groups (n=6): untreated diabetic (D) and diabetic treated with metformin (DM). Metformin ((metformin hydrochloride, Merck) was given daily by oral gavage at 500 mg/kg body weight. Ten days after diabetes induction, DM rats were subjected to a 7 day treatment with metformin and blood samples and soleus muscles were obtained from anesthetized rats (sodium thiopental, 60 mg/kg, ip) before and 16 min after intravenous injection of regular insulin (0.75U/kg BW) to intravenous insulin tolerance test. Then, animals were euthanized by sodium thiopental overdoses (80 mg/kg body weight).

### *Intravenous insulin tolerance test*

Glucose levels were measured on samples obtained from the tail vein in *ad libitum* fed animals before (0min) and 4, 8, 12, and 16 min after intravenous injection of regular insulin (0.75U/kg BW). The constant rate for blood glucose disappearance during insulin tolerance test (KITT) was calculated based on the linear regression of the Neperian logarithm of blood glucose concentrations (test strips, Advantage, Roche). The tests were performed between 0900 to 1100h [35].

### *Sample collection and Tissue Preparation*

The soleus muscles were quickly removed before (right side) and 16 min after (left side) regular insulin injection to intravenous insulin tolerance test. After that, the

tissues were washed with chilled saline (NaCl 0.9%) and immersed in liquid nitrogen. For ELISA and western blotting analyses, soleus muscles were homogenized in buffer (25 mM Tris-HCl pH 7.4, cocktail protease inhibitor, 10 mM EDTA and 10 mM EGTA) during 3 min. The homogenized was centrifuged at 15.000xg for 40 min. at 4°C. The pellet was resuspended in the same homogenization buffer. The supernatant were used for the WB analysis.

### *ELISA*

CXCL1/KC, TNF $\alpha$  proteins detection was performed with commercially available sandwich ELISA kits with capture and detection antibodies according to manufacturer's instructions (R&D Systems, Minneapolis, MN). Each sample and standard was analyzed in triplicate. The plates were read on a microplate reader (Molecular Devices, Sunnyvale, CA) and analyzed (SOFTmax software; Molecular Devices).

### *SDS-PAGE and Western blotting*

Protein samples (10  $\mu$ g) were applied to 5–22% polyacrylamide gradient gels under denaturing conditions [36] and stained with Coomassie brilliant blue. For immunoblotting, proteins were transferred to nitrocellulose membranes in Tris–glycine buffer as described[37]. The membranes were incubated with 5% dried milk in Tris-buffered saline (TBS-T) (50 mM Tris–HCl, pH 8.0, 150 mM NaCl, 0.05% Tween 20) and immunodetected using primary antibodies (anti-TLR4, anti-TLR2, anti NF- $\kappa$ B, anti-p-AMPK, anti, p-JNK, anti-Bcl2 and, anti-BAX ), diluted to 0.2  $\mu$ g/mL in TBS-T [37]. The reaction was detected using an ECL kit (Amersham Biosciences, Buckinghamshire, UK) according to the manufacturer's instructions, and Amersham Imager 600 was used for exposure. The intensities of the protein bands were analyzed and quantified using the ImageQuant TL software, version 8.1 (GE Healthcare Life Science), and the results were expressed as a percentage.

### *Statistical analysis*

All results were presented as mean  $\pm$  SD. The Shapiro-Wilk test was used to evaluate the normality of data distribution. Data were normalized with untreated diabetic group before regular insulin injection as 1.00. Comparisons of means were performed by unpaired Student's t-test and One-way ANOVA, Student–Newman–Keuls *post hoc* test. Differences were considered significant at  $p < 0.05$ .

## **Results**

Table 1. Morphometric and metabolic parameters of diabetic rats untreated and treated with metformin.

Metformin doses was 500 mg/kg body weight/day. Basal glycemia was measured in blood collected at the beginning (time 0) of the insulin tolerance test. Data are mean $\pm$ SD. of 6 rats. One-way ANOVA, Student–Newman–Keuls *post hoc* test. # Statistical difference between untreated and treated with metformin. \* Statistical difference between before and after insulin injection  $p < 0.05$ .

	untreated		metformin treatment	
	initial	final	initial	final
<b>Body weight</b>	265.3±23.8	283.9±27.9	244.9±14.2	312.6±17.9 *#
<b>Soleus muscle weight</b>	0.150±0.009	0.146±0.012	0.149±0.012	0.145±0.02
	before insulin	after insulin	before insulin	after insulin
<b>Blood glucose</b>	465.8±68.5	443.3±25.9	360.2±55.8 #	284±20.5 *#

Table 1 shows that the untreated diabetic rats have a less weight gain compared to metformin treatment and, no change was observed in soleus muscle weights. As expected, higher blood glucose level was observed in untreated diabetic rats and metformin treatment decrease the glycaemia. In addition, after insulin injection we observed a decrease of blood glucose level in metformin treatment compared with each other groups.

*Representative figure of the role of metformin and insulin in soleus muscle of diabetic rats.*

Supplementary figure 1 shows the protein profile and representative images of apoptotic and pro-inflammatory proteins WB in soleus muscle of diabetic rats untreated and treated with metformin as well as, before and 16 min after regular insulin injection. The statistical analyses of the quantification of these proteins are demonstrated below.

*Metformin role on inflammatory markers expression in soleus muscle of diabetic rats*

To verify whether metformin treatment decreases inflammatory signal in skeletal muscle of diabetic rats and to analyze if inflammatory pathway is dependent of insulin, soleus muscles were removed from rats untreated and treated with metformin before, and after (16min), regular insulin injection. Figure 1 shows that metformin treatment suppressed both TLR2 and TLR4 concentration in soleus muscle of diabetic rats. However, only TLR4 concentration was increased 16 min after regular insulin injection in rats treated with metformin (Figure 1.A and 1.B). In addition, metformin treatment decreased NF-κB and increased IκB concentrations. After regular insulin injection, NF-κB decreased and IκB increased in untreated diabetic rats. These changes in NF-κB and IκB parameters were not observed in insulin-treated rats (Figure 1.C and 1.D).

Then we checked if the metformin alters the expression of two inflammatory cytokines (TNFα and CXCL1/KC) in soleus muscle and the role of insulin in this process. Metformin treatment decreased TNFα and CXCL1/KC concentrations in soleus muscle compared with untreated diabetic group. Moreover, the insulin injection promoted the increase of TNFα in both untreated and treated with metformin diabetic rats (Figure 1.E and 1.F). However, CKCL1/CK increased only

in untreated diabetic group after insulin injection (Figure 1.F). Furthermore, metformin treatment decreased p-JNK concentration (Figure 1.G) before regular insulin injection. As expected, metformin treatment increased p-AMPK expression prior to insulin administration (Figure 1.H). After regular insulin injection, p-AMPK increased in untreated diabetic rats and these changes were not observed in metformin-treated rats.

#### *Metformin role on insulin sensitivity of diabetic rats*

To verify whether metformin treatment improves the insulin sensibility, decay glucose and plasma glucose disappearance rate were measured. Metformin treatment increase decay glucose and increase the KITT compared to untreated diabetic rats. The slope of plasma glucose concentration during 0-16min. after insulin administration in the metformin-treated group was higher than the untreated group, indicating an improving in the insulin sensitivity (Figure 2).

#### *Metformin role on apoptotic markers expression in soleus muscle of diabetic rats*

To understand how metformin treatment alters the dynamic of apoptotic signaling in the skeletal muscle and whether this mechanism is dependent of insulin, we probe the same samples described previously. Metformin treatment increased BAX concentration before regular insulin injection compared to untreated group (Figure 3.A) and, interestingly, similar results were observed for Bcl2 concentration. However, after insulin injection an increase of Bcl2 was verified only in the untreated group (Figure 3.B). No difference was observed on BAX:Bcl2 ratio in both untreated and treated diabetic rats, before and after insulin injection (Figure 3.C).

### **Discussion**

A large number of evidences suggest that metformin acts on different signaling pathways [38]. However, there are few data of this action on inflammatory/apoptotic signaling pathway in skeletal muscle. This tissue has a central role on peripheral glucose uptake and the role of inflammation and metformin treatment in skeletal muscle, especially in soleus, remains undetermined. Hyperglycemia is associated with an increase of the pro-inflammatory pathway [1-3] and disorders of the innate immune system [4]. Studies in other tissues showed that metformin acts on TLRs activity [33] and NF- $\kappa$ B/I $\kappa$ B signal [31, 39] attenuating the transcription of pro-inflammatory regulators. In the regulation of apoptotic pathway, AMPK activation by metformin also stimulate JNK-Bcl2 signaling [34]. These findings suggest that metformin may attenuate the inflammatory injuries promoted by hyperglycemia. To our knowledge, we demonstrate for the first time that metformin treatment suppresses inflammatory pathways (TLR2/TLR4 and NF- $\kappa$ B/ I $\kappa$ B/ p-JNK, TNF $\alpha$  and CXCL1/KC) as well as, attenuates apoptotic signaling (BAX:Bcl2 ratio) independent of p-AMPK in soleus muscle of diabetic rats.

In the inflammatory pathway, TLRs can initiate a complex signaling cascade that activates inflammatory markers leading to cellular injury (more details [40]). It is

know that diabetes stimulates the TLRs activation [41, 42] and that TLR2 and TLR4 are expressed in insulin target tissues [40, 43]. Here, we show a higher concentration of these receptors in soleus muscle of untreated diabetic rats compared to metformin treatment. Some studies have shown that metformin treatment suppressed TLRs pathway [33] and decreased the TLR4 concentration by attenuating the TLR4 mRNA [44]. This finding in other tissues corroborate with the present results in soleus muscle, which showed the attenuation of both TLR4 and TLR2 concentrations in diabetic rats treated with metformin. We also observed an increase of TLR4 concentration after regular insulin injection, this is in agreement with other study which showed the role of insulin in the transcription of TLRs [45].

Furthermore, TLR2 and TLR4 are central to the regulation of inflammatory responses through activation of NF- $\kappa$ B [9-12]. Some previous studies suggested that metformin may inhibit NF- $\kappa$ B via activation of I $\kappa$ B in vascular endothelial cells [39] and suppress the expression of NF- $\kappa$ B induced by high blood glucose level via AMPK activation in glomerular mesangial cells *in vitro* [46, 47]. As expected, metformin treatment increased phosphorylated AMPK in soleus muscle of diabetic rats. The reduction of phosphorylated AMPK in soleus after acute effects of insulin (16 minutes) are in agreement with previous studies that suggest that insulin, via activation of Akt, inhibits AMPK activity by phosphorylation of Ser 485 of  $\alpha$ -AMPK [48, 49]. The present set of data confirmed that metformin was also able to reduce NF- $\kappa$ B associated with the increase in I $\kappa$ B concentration in soleus muscles of diabetic rats. Besides, the association of metformin with acute effects (16 minutes) of insulin do not show additional effects on NF- $\kappa$ B and I $\kappa$ B concentrations. To our knowledge, the present study showed for the first time that acute effects of insulin (16 minutes) promotes decrease in NF- $\kappa$ B concentration and increase in I $\kappa$ B concentration.

NF- $\kappa$ B role in transcriptional activation to production of inflammatory cytokines is well established [20, 21]. Several evidences indicate that TNF $\alpha$  is a cytokine involved in systemic inflammation. Other studies showed that metformin inhibited TNF $\alpha$  expression in bladder smooth muscle cells [50] as well as, inhibit TNF $\alpha$ -induced mRNA expression and NF- $\kappa$ B activation via AMPK in endothelial cells [31, 39]. However, this mechanism of inhibition has not yet been described in skeletal muscle. In the present study, we demonstrate that metformin treatment suppressed TNF $\alpha$  expression in skeletal muscle of diabetic rats which may be associated with decreased of NF- $\kappa$ B [51]. Surprisingly, insulin treatment promotes an increase in TNF $\alpha$  levels in soleus of diabetic untreated rats and diabetic metformin-treated rats. We suggest that these acute effects (16 minutes) of insulin was reduced due to metformin treatment. In addition, we showed a decrease of CXCL1/KC after metformin treatment. This chemokine is involved in the recruitment and activation of neutrophils [52] linked with the increase of TNF $\alpha$  [53, 54]. However, the mechanisms by which metformin inhibit this cytokine inflammatory response remains unclear.

Metformin also attenuate the activation of JNK in Alzheimer's disease-like neuropathology [55]. In addition, JNK can be activated by TNF $\alpha$  [56, 57] which is



increased in hyperglycemic condition [58] impairing insulin pathway. These results are consistent with our findings that showed a decrease of p-JNK levels related with the reduction in TNF $\alpha$  levels after metformin treatment. However, the regulation of these proteins is not correlated after acute effects of insulin in soleus muscle of diabetic rats. Some studies also showed that JNK pathway impairs insulin signal transduction directly through IRS inhibitory phosphorylation [17-19]. Although our study has not analyzed IRS substrate, we verified that metformin treatment improves glucose uptake as demonstrated by the increase of insulin sensitivity. This evidence indicated that metformin promotes the reduction of the inflammatory markers in muscle cells, which improves insulin pathway and contributes to glucose homeostasis.

Recently was reported that metformin also inhibit JNK phosphorylation in rat hepatocyte against oxidative stress-induced apoptosis [59]. Inflammatory pathway and apoptotic signaling may be interconnected by JNK/ AMPK-induced activation [60, 61]. It is well documented that JNK promotes BAX translocation to mitochondria releasing cytochrome c, which leads to apoptosis [25], as well as suppress the function of the anti-apoptotic protein - Bcl2 [26]. The ratio of BAX:Bcl2 is detrimental to the life/death signaling following apoptotic stimuli which was significantly increased in metformin-treated tumors. The antitumor effect of metformin appears to be dependent on the inhibition of NF- $\kappa$ B and mTOR signaling pathways [62]. Our results are in agreement with those presented in other studies, once we observed an increase in BAX concentration after the treatment with metformin. However, we also show that metformin decreases p-JNK concentration and increases the concentration of Bcl2 independent of p-AMPK. Other studies have also observed that the treatment with metformin decreased the JNK concentration, modulating the expression of Bcl2 and attenuating apoptotic processes [63, 64]. Considering that BAX:Bcl2 ratio was unchanged and p-JNK decreased after metformin treatment, the present data also revealed an attenuation of the apoptotic signaling *in vivo*.

Taken together, our data point out that metformin may attenuate the activation of the inflammatory pathway TLRs/NF- $\kappa$ B/TNF $\alpha$ /CXCL1/KC and the apoptotic signaling BAX:Bcl2 ratio/p-JNK independent of p-AMPK, which could be accompanied by a reduction of the inflammatory damage caused by hyperglycemia in skeletal muscle of diabetic rats. These alterations may explain the improvement of insulin sensitivity observed in diabetic rats treated with metformin. Furthermore, the treatment with metformin was not capable of attenuating the changes in the inflammatory pathway triggered by insulin injection.

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**Author contributions**

Conceived and designed the experiments: L.G.P., R.R.T., R.S.S., F.S.E.  
Performed the experiments: L.G.P., R.R.T., D.D.V., L.N.B., S.R.D., Analyzed the data: L.G.P., R.R.T., R.S.S., F.S.E., Wrote the paper: L.G.P., L.N.B., R.R.T., R.S.S., Supervised the entire project: F.S.E.

**Declaration of interest**

The authors report no conflicts of interest. The authors alone are responsible for the content and writing of the paper.

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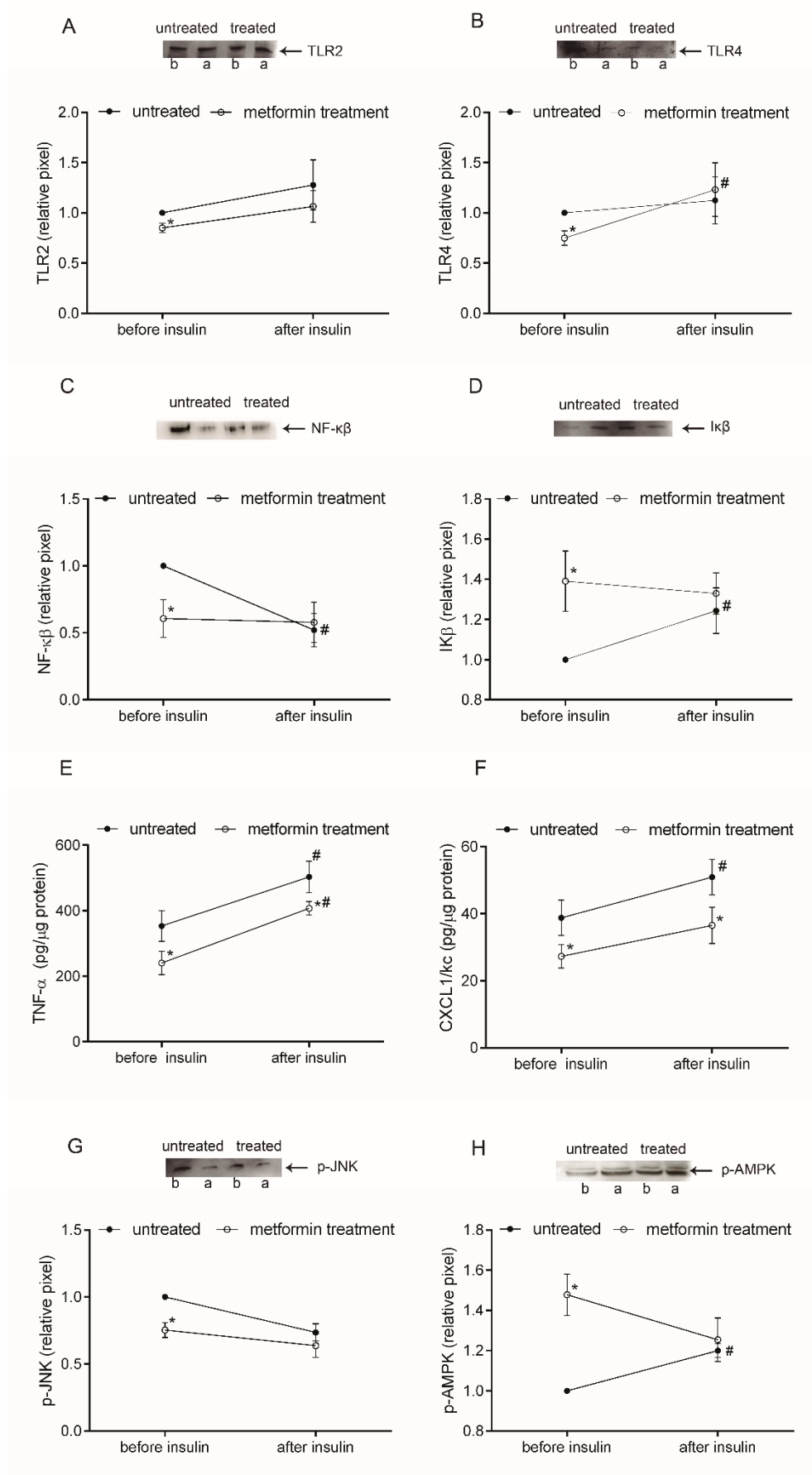
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## Figures and legends

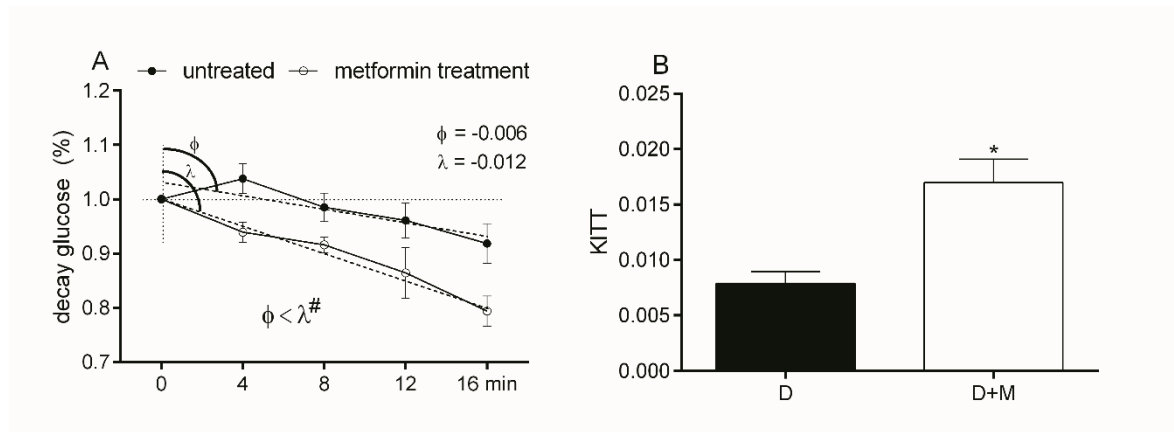
Figure 1. Metformin role on inflammatory pathway





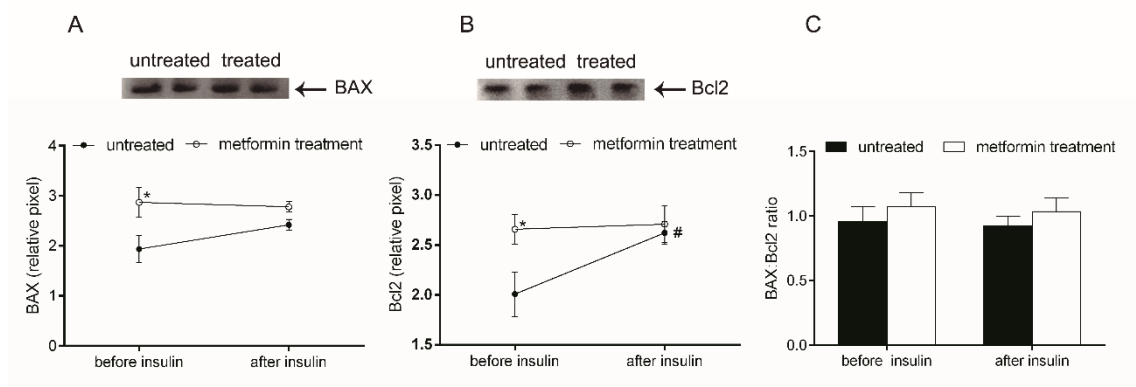
**Figure 1.** Effect of metformin on inflammatory signaling triggered by TLRs in soleus muscle. The tissue was collected before and 16 min after regular insulin injection (0.75U/kg BW). Variation of **A-** TLR2; **B-** TLR4; **C-** NF- $\kappa$ B; **D-** IK $\beta$ , **E-** TNF $\alpha$ ; **F-** CXCL1/KC; **G-** p-JNK; **H-** p-AMPK concentration before and after insulin injection. Results are means  $\pm$  SD of 6 rats/group. \* $p < 0.05$  vs. untreated diabetic rats; #  $p < 0.05$  vs. before insulin injection; paired Student's.

**Figure 2. Glucose disappearance rate.**



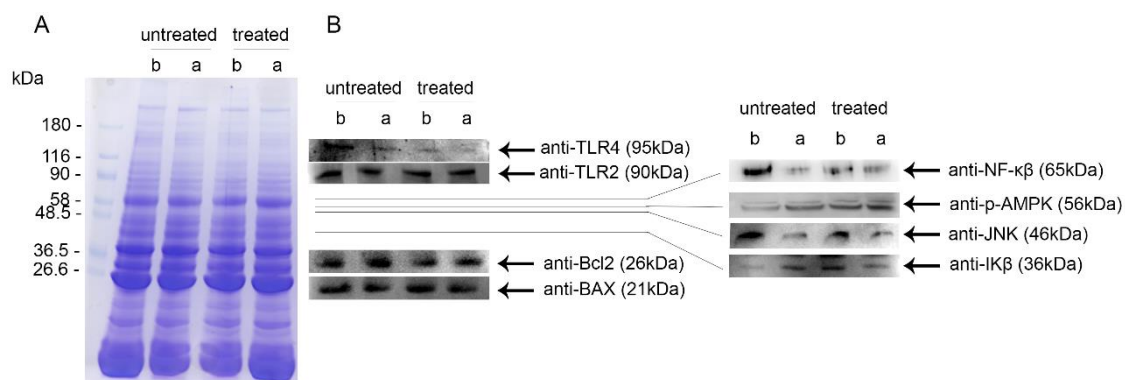
**Figure 2:** The effects of metformin on the decay glucose (A) and plasma glucose disappearance rate (KITT) (B) in diabetic rats. Glycaemia was measured (0min) and 4, 8, 12, and 16min after i.v. injection of regular insulin (0.75U/kg BW). Results represent the mean $\pm$ SD (n=6),  $p \leq 0.05$ , (#) statistical difference before and 16 min after regular insulin injection using t-test.

**Figure 3. Metformin role on apoptotic pathway**



**Figure 3:** Effect of metformin on apoptotic pathway in soleus muscle. The muscle was collected before and 16 min after regular insulin injection (0.75U/kg BW). Dynamic of **A**-BAX; **B**- Bcl2; **C**- BAX:Bcl2 ratio; concentration. Results are means  $\pm$  SD of 6 rats/group. \*p<0.05 vs. untreated diabetic rats; # p<0.05 vs. before insulin injection; paired Student's.

## Supplementary figure. Summary of SDS PAGE and WB analysis.



Supplementary figure: Apoptotic and pro-inflammatory signaling pathways of soleus muscle before and after insulin injection in diabetic rats untreated and treated with metformin. **A-** SDS-PAGE 5-22% stained with coomassie blue (b: before insulin injection; a: after insulin injection). **B-** WB of apoptotic and pro-inflammatory pathway.