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## Gas6 - Axl receptor signaling is regulated by glucose in vascular smooth muscle cells

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### Abstract

**Objective**—The receptor tyrosine kinase Axl and its ligand Gas6 are involved in the development of renal diabetic disease. In vascular smooth muscle cells (VSMC) Axl is activated by reactive oxygen species and stimulates migration and cell survival, suggesting a role for Axl in the vascular complications of diabetes.

**Methods and Results**—We investigated the effect of varying glucose concentration on Axl signaling in VSMC. Glucose exerted powerful effects on Gas6-Axl signaling with greater activation of Akt and mTOR in low glucose, and greater activation of ERK1/2 in high glucose. Plasma membrane distribution and tyrosine phosphorylation of Axl were not affected by glucose. However, co-immunoprecipitation studies demonstrated that glucose changed the interaction of Axl with its binding partners. Specifically, binding of Axl to the p85 subunit of PI3-kinase was increased in low glucose, whereas binding to SHP-2 was increased in high glucose. Furthermore, Gas6-Axl induced migration was increased in high glucose, while Gas6-Axl mediated inhibition of apoptosis was greater in low glucose.

**Conclusion**—This study demonstrates a role for glucose in altering Axl signaling through coupling to binding partners, and suggests a mechanism by which Axl contributes to VSMC dysfunction in diabetes.

### Keywords

Axl; glucose; vascular smooth muscle migration; signaling

### Introduction

Hyperglycemia alters vascular smooth muscle cell (VSMC) signaling, which is a major factor in the development of vascular complications of diabetes such as atherosclerosis and hypertension<sup>1-3</sup>. The receptor tyrosine kinase Axl is expressed in several cell types including VSMC<sup>4,5</sup>. Axl is activated by Growth Arrest Gene 6 (Gas 6), which is a homologue of Protein S<sup>6,7</sup>. This leads to stimulation of downstream signaling cascades including the PI3K-Akt pathway, and ERK1/2<sup>4,8-10</sup>. Gas6 activation of Axl in VSMC increases directed migration and inhibits apoptosis<sup>11,12</sup>. There are several reports that suggest a role for Axl in the pathogenesis of vascular and diabetic disease. Expression of Axl and Gas6 are increased in the glomerulus of diabetic rats. Furthermore, diabetic nephropathy is less severe in Gas6 knockout mice and when Gas6 activity is inhibited by warfarin treatment<sup>13</sup>. Axl is highly upregulated

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in balloon injured carotid arteries with a time course paralleling that of neointima formation<sup>14</sup>. Axl mRNA and protein level in cultured VSMC are also increased by the G-protein-coupled receptor agonists thrombin and angiotensin II<sup>14</sup>. Axl is activated by H<sub>2</sub>O<sub>2</sub>, which is increased in vascular injury and hyperglycemia, in both VSMC and *ex vivo* vessels<sup>15</sup>. In addition, neointima formation induced by injury is decreased in Axl knockout mice<sup>15</sup>. These studies suggest Gas6-Axl represents an important pathogenic mechanism for cardiovascular and renal complications associated with diabetes.

Recently, we demonstrated that high glucose enhanced phosphorylation of Akt and ERK1/2 by angiotensin II through alterations in epidermal growth factor receptor (EGFR) N-glycosylation<sup>16</sup>. Based on this study, we hypothesized that glucose would modulate Axl signaling and thereby VSMC function. Interestingly, while Gas6-stimulated ERK1/2 signaling was greater in high glucose than low glucose, the opposite was true for Akt. This suggests that glucose modulates the downstream coupling of Axl to its effectors. Here we show increased interaction between PI3K and Axl in low glucose and increased interaction between the protein tyrosine phosphatase SHP-2 and Axl in high glucose. Furthermore, Gas6-Axl signaling increased cell survival in low glucose and increased migration in high glucose. Our data demonstrate that glucose modulates Axl signaling via different cell signaling mechanisms and this may contribute to the vascular complications of diabetes.

## Methods

### Materials

Antibodies to Axl and ERK1/2 were from Santa Cruz Biotechnology; antibodies to phospho-ERK, phospho-Akt (Ser-473), Akt, mTOR, phospho-mTOR (Ser-2448); p85 PI3K antibody and Akt antibody were from Upstate, LiCor fluorescent secondary antibodies were from Molecular Probes. Sulfo-NHS-SS-biotin and streptavidin agarose were from Pierce. Gas6 was kindly provided by Brian Varnum (Amgen). All other reagents and chemicals were obtained from Sigma, unless specifically indicated.

### Cell Culture

Cultured VSMC were obtained from rat aorta as described<sup>17</sup>. VSMC were grown in Dulbecco's modified Eagle Medium supplemented with 25 mM NaHCO<sub>3</sub>, 10 mM HEPES, pH 7.4, 50 IU/ml penicillin, 50 µg/ml streptomycin, 10 % fetal bovine serum (FBS) containing 5.5 mM glucose (low glucose (LG)) in a 5 % CO<sub>2</sub>/95% O<sub>2</sub> incubator at 37 °C. For high glucose (HG), cells were grown in 27.5 mM glucose and controls received 22.5 mM mannitol and 5 mM glucose (total).

### Preparation of cell lysates and Immunoprecipitations

Cell monolayers were rinsed with ice-cold phosphate-buffered saline (PBS; 150 mM NaCl, 20 mM Na<sub>2</sub>PO<sub>4</sub>, pH 7.4) and then scraped in 1 ml of PBS. After a brief centrifugation, the cells were solubilized in 1 ml of cell lysis buffer (10 mM HEPES, pH 7.4, 50 mM Na Pyrophosphate, 50 mM NaF, 50 mM NaCl, 5 mM EDTA, 5 mM EGTA, 1 mM Na<sub>3</sub>VO<sub>4</sub>, 0.5 % Triton plus 1:1000 protease inhibitor cocktail). Cells were sonicated for 20 s, agitated on a rotating rocker at 4 °C for 30 min and centrifuged at 12,000 g for 30 min to remove insoluble cellular debris.

For immunoprecipitation studies, lysates were precleared for 1 hr with protein G agarose (Invitrogen) followed by incubation with anti-Axl antibody for 3 hr, and protein G agarose for a further 1 hr. Immunoprecipitates were then washed 4 times with 1 ml cell lysis buffer before the addition of Laemmli sample buffer. After heating at 95 °C for 3 min, proteins were resolved on SDS-PAGE and transferred to nitrocellulose membranes for Western analysis.

Immunoreactive bands were detected with LiCor fluorescent secondary antibodies and the

LiCor Odyssey infrared Imaging system. Analysis of blots was performed using the LiCor densitometry software.

### Glycosidase Digestion

Peptide N-glycosidase F (PNGase-F, New England Biolabs) digestion was performed as described<sup>16</sup>.

### Cell surface biotinylation

Cells were biotinylated as described<sup>16</sup>. Cells were then lysed in 1 ml of N+ buffer (in mM: 60 HEPES, pH 7.4, 150 NaCl, 3 KCl, 5 EDTA trisodium, 3 EGTA, and 1% Triton X-100) and sonicated to clarity on ice with a Branson 450 probe sonicator. NHS-SS-biotin labeled proteins were precipitated using streptavidin agarose. The avidin-agarose beads were washed five times in N+ buffer, and bound proteins were solubilized in sample buffer yielding the surface fraction. Samples were separated on a 9% gel by SDS-PAGE, and Western analysis was then performed.

### Subcellular Fractionation

Subcellular fractionation was performed as described<sup>16</sup> and an equal volume of each fraction was analyzed by SDS-PAGE.

### Caspase-3-Enzyme Activity

Caspase-3-activity was detected by the color absorbance of the chromophore *p*-nitroanilide (pNA) obtained after proteolytic cleavage from the labeled substrate Asp-Glu-Val-Asp-pNA (DEVD-pNA) and pNA as standard using a spectrophotometer 405 nm (Clontech ApoAlert Caspase-3 assay kit) according to the manufacturers protocol.

### Migration Assays

VSMC migration was measured using a Boyden chamber assay. Cells were incubated in LG or HG serum free media for 24 hr. Control medium (0% serum DMEM), Gas6 medium (0% serum DMEM with 200 ng/ml Gas6) or serum medium (10 % serum DMEM) was placed in the lower chamber. A collagen-coated polyvinylpyrrolidone-free polycarbonate membrane was placed on top, and  $2 \times 10^5$  cells suspended in 50  $\mu$ l 0% serum DMEM/0.1 % BSA were seeded in the upper chamber. After 6 hr incubation at 37 °C in 95 % air/5% CO<sub>2</sub>, cells attached to the lower side were fixed and stained using a Diff-Quik stain set (Dade Behring). The number of migrating cells was quantified using densitometry (NIH Image).

**Statistical analysis**—All experiments were carried out at least 3 times. The difference between LG and HG was assessed by analysis of variance.

## Results

### Glucose modulates Akt and ERK1/2 phosphorylation upon stimulation with Gas6

Previously, we found that that HG altered transactivation of the EGFR by angiotensin II; with increased ERK1/2 and Akt phosphorylation in HG as compared to LG<sup>16</sup>. To explore the effect of glucose on Gas6-Axl signal transduction, we studied Akt and ERK1/2 activation. The phosphorylation of ERK1/2 by Gas6 was dramatically increased in HG as compared to LG (Figure 1A). In contrast, the phosphorylation of Akt by Gas6 was opposite, dramatically increased in LG as compared to HG (Figure 1B). (Please see Figure I supplement at [www.ahajournals.org](http://www.ahajournals.org) for quantification of A and B). This demonstrates that the ability of Axl to stimulate Akt and ERK1/2 is differentially regulated by glucose concentration.

### Glucose does not modulate phosphorylation of Axl or cell surface expression of Axl

Previously, our lab demonstrated that the molecular mass of the EGFR varies depending on the glucose concentration due to N-glycosylation<sup>16</sup>. We hypothesized that glucose might also affect Axl N-glycosylation and therefore studied the effect of LG vs. HG on Axl molecular mass. In HG Axl was present predominantly as a 140 kDa form ( $62.2 \pm 1.7\%$ ), with a small proportion present as 120 kDa ( $16.1 \pm 1.4\%$ ) and 114 kDa ( $21.7 \pm 0.85\%$ ) forms (Figure 2A, B, C). In contrast, in LG the 114 kDa form of Axl was the major form (Figure 2A,B, C). To show that the Axl isoform change was due to glucose levels and not osmotic stress, the metabolically inactive sugar mannitol (22.5 mM) was added to LG media. This had no effect on the appearance of the 114 kD isoform in LG (data not shown).

Glucose concentration has previously been shown to alter the level of membrane protein N-glycosylation<sup>16, 18-20</sup>. Therefore, we used N-glycosidase F (PNGase F), which removes carbohydrates from glycopeptides at N-glycosylation sites, to determine whether the difference in Axl molecular mass was due to Axl N-glycosylation. PNGase-F treatment yielded a 114 kDa form of Axl from lysates of HG and LG treated cells demonstrating that the difference in molecular mass is due to N-glycosylation (Figure 2A). An important function for N-linked glycosylation is to target membrane proteins to the plasma membrane. Lack of proper glycosylation can cause proteins to be improperly folded and retained in the endoplasmic reticulum<sup>21</sup>. We used cell surface biotinylation to determine whether the 114 kDa form of Axl was present in the plasma membrane at similar levels to the 140 kDa Axl. Membrane proteins were labeled with a membrane impermeant form of biotin, then after cell lysis the biotinylated proteins were extracted using streptavidin agarose. As shown in Figure 2B, in both LG and HG the same amounts of 114 kDa Axl and 140 kDa Axl were present in the plasma membrane demonstrating that glucose concentration does not alter cell surface levels of Axl. Finally, in response to Gas6 both 114 kDa and 140 kDa were tyrosine phosphorylated with similar time courses (Figure 2C), consistent with cell surface localization.

### Glucose modulates the interaction between p85-PI3-kinase (p85-PI3K) and Axl

To determine if enhanced activation of Akt by Gas6 in LG was due to an increased interaction between Axl and p85-PI3K, co-immunoprecipitation of Axl with p85-PI3K was determined. Gas6 increased interaction between p85-PI3K and Axl, as previously reported<sup>22</sup>. However, in LG, the interaction between Axl and p85-PI3K was significantly increased as compared to in HG (Figure 3A). This suggests that the increased phosphorylation of Akt in LG after Gas6 stimulation is due to increased binding to p85-PI3K.

### Glucose modulates the interaction between SHP-2 and Axl

The protein tyrosine phosphatase SHP-2 associates with tyrosine kinase receptors including Axl<sup>23</sup> and positively regulates ERK activation<sup>24-26</sup>. Therefore we determined whether glucose concentration modulated SHP-2 interaction with Axl. SHP-2 co-immunoprecipitated with Axl in HG. However, an interaction between SHP-2 and Axl could not be detected in LG (Figure 3B). Therefore increased phosphorylation of ERK1/2 in HG may be due in part to increased association of SHP-2 with Axl. Gas6 induces Axl activation in LG leads to Akt activation via interaction of Axl with p85-PI3K (Figs 1 and 3). To characterize further the anti-apoptotic mechanism of Axl in LG conditions, modulation of downstream signaling molecules was determined. The PI3K-Akt pathway antagonizes apoptosis in VSMC as well as other cell types, in part via activation of mTOR. Because mTOR is expressed in VSMC<sup>27</sup>, we tested the hypothesis that mTOR activation by Gas6 is stimulated in LG. There was a slight increase in Gas6 induced phosphorylation of mTOR at Ser 2448 in LG. However, the baseline level of mTOR phosphorylation was much less in HG and upon stimulation with Gas6 there was a significant time dependent decrease in phospho-mTOR (please see Figure II supplement at [www.ahajournals.org](http://www.ahajournals.org)).

To determine whether the altered interactions with p85-PI3K and SHP-2 were due to altered distribution of Axl within the plasma membrane in LG vs. HG, by subcellular fractionation sucrose gradient density centrifugation was performed (Figure 4). This demonstrated that Axl was predominantly present in caveolin enriched fractions both basally (Figure 4A) and after Gas6 (Figure 4B) stimulation, and this was not altered by glucose concentration (Figure 4).

### The anti-apoptotic effect of Axl is enhanced in LG conditions

To determine whether the Gas6-Axl mediated increase in Akt signaling in LG has physiological consequences, the ability of Axl to protect VSMC from apoptosis was determined by measuring caspase-3 activity. VSMC were incubated in 0% serum media containing LG or HG for 24 hr. Apoptosis was induced by serum deprivation alone or serum deprivation and TNF $\alpha$  (10 ng/ml) for 24 hr in LG or HG media. Caspase-3 activity was increased by 22-fold vs. control in LG +0% serum in contrast to a 1.6-fold increase vs. control in HG+0% serum. After TNF $\alpha$  treatment, caspase-3 activity was 35-fold higher than control in LG in contrast to 2.4-fold higher than control in HG (Figure 5A). This is consistent with previous studies, which demonstrate that high glucose is anti-apoptotic<sup>28-30</sup>. After treatment with Gas6, caspase-3 activity in LG +0% serum was reduced from 22-fold to 11-fold, but was unchanged in HG+0% serum (1.6-fold vs. 1.5-fold; Figure 5A). After TNF $\alpha$  treatment in LG, caspase-3 activity was reduced from 35-fold to 17-fold, but was unchanged in HG (2.4-fold vs. 3.4-fold; Figure 5A). Because the baseline level of apoptosis in HG was so low it would be difficult to achieve significant differences, but the fact that the trend shows no change toward less apoptosis is consistent with the Akt data in Fig. 1. The 114 kDa Axl is the Axl isoform that predominates in the signaling of this anti-apoptotic effect based on the data in Figure 2. These results are consistent with the greater Akt activation observed in LG (Figure 1B-C).

### Migration is enhanced in response to Gas 6 in HG

Since Gas6 stimulation of Axl increases VSMC migration<sup>12</sup>, we determined the effect of LG and HG on Gas6 stimulated VSMC migration. VSMC were incubated in 0% serum DMEM containing LG or HG. Baseline migration in LG and HG was similar as was migration in response to 10% serum (Figure 5B). However, in response to 200 ng/ml Gas6, migration in HG was  $27 \pm 2\%$  higher than in LG ( $p < 0.005$ ; Figure 5B). This demonstrates that the pro-migratory effect of Axl is increased in HG and suggests that the 140 kDa Axl plays a more significant role in migration than the 114 kDa Axl.

## Discussion

In this study we investigated the effect of glucose concentration on Axl signal transduction in VSMC. The major finding is that glucose alters Gas6-Axl signaling by modulating interaction of Axl with specific signaling proteins. In HG, Gas6-Axl stimulation increased ERK1/2 activation while in LG Gas6-Axl stimulated PI3K-Akt-mTOR. It appears that the 140kDa Axl is primarily responsible for ERK 1/2 activation while the 114kDa Axl is responsible for Akt activation. Increased interaction of Axl with SHP-2 in HG and increased interaction with PI3-K in LG correlated with these alterations in Axl downstream signals. Functionally, this resulted in increased migration in response to Gas6 in HG and increased protection against apoptosis in LG.

Glucose has previously been shown to modulate signaling in VSMC. HG activates protein kinase C and this inhibits VSMC apoptosis<sup>30</sup>. Increased activation of ERK1/2 and p38 also occurs in HG<sup>31, 32</sup>. Recently, we demonstrated that transactivation of the EGFR by angiotensin II was increased in HG, which increased both Akt and ERK1/2 phosphorylation<sup>16</sup>. Interestingly, the activation of Akt by Axl was higher in LG than in HG and in contrast, Axl activated ERK1/2 more in HG as compared to LG. Axl is anti-apoptotic and pro-migratory in

VSMC<sup>11, 12</sup>. Previous studies have shown that the anti-apoptotic effect is predominantly through PI3K-Akt signaling<sup>11</sup>. The present study demonstrates that in LG Axl predominantly activates Akt signaling and therefore mediates cell survival, via a mechanism that includes caspase 3 inhibition. The anti-apoptotic effect of Gas6 stimulation in LG is further supported by the activation of signals downstream of Akt, including mTOR. The effects of Akt and mTOR interaction have been shown to decrease apoptosis in endothelial cells<sup>33</sup> and the Gas6-Axl and Akt/mTOR pathways play a crucial role in mesangial and glomerular hypertrophy<sup>13</sup>. Thus we propose a similar signaling mechanism in VSMC. We also demonstrated increased migration in HG as compared to LG in response to Gas6, suggesting a key role for ERK1/2 signaling in Gas6-Axl mediated migration in HG.

We propose a model for the downstream cell signaling mechanisms of Gas6 mediated activation of Axl in HG and LG (please see Figure III supplement at [www.ahajournals.org](http://www.ahajournals.org)). The mechanism by which glucose modifies Axl signaling appears to be altering Axl binding with signaling molecules. Activation of p85-PI3K and its downstream target Akt by Gas6-Axl requires interaction between PI3K and Axl<sup>9, 34, 35</sup>. This study suggests increased association of Axl with p85-PI3K mediates increased activation of Akt and mTOR in LG. Previous studies demonstrate that the protein tyrosine kinase SHP-2 is required for ERK1/2 activation by growth factors<sup>24-26</sup>. Gas6 activates SHP-2 and increases SHP-2 association with Axl<sup>23</sup>. We demonstrated that there is increased association of SHP-2 with Axl under HG conditions and this may mediate the increased ERK1/2 activation. The mechanism by which PI3K associates with 140kDa Axl and SHP2 with 114kDa Axl selectively remains to be determined.

In the case of the EGFR, N-glycosylation appears to change the distribution of EGFR within plasma membrane microdomains thereby modulating angiotensin II transactivation of the EGFR<sup>16</sup>. However, this cannot explain the effect of glucose on Axl since plasma membrane distribution was predominantly in the caveolae fraction in both LG and HG conditions. N-glycosylation of the EGFR also alters ligand binding properties and receptor dimerization, thus modifying EGFR downstream signaling<sup>36-38</sup>. This seems unlikely for Gas6-Axl since signaling to ERK1/2 and Akt occurred at high levels. A likely possibility is that cellular distribution of p85-PI3K and SHP-2 is altered depending on glucose concentration. Further studies will elucidate the mechanism by which glucose concentration and N-glycosylation state of Axl regulate interaction of Axl with signaling molecules.

This novel finding that Gas6-Axl-mediated Akt and ERK1/2 activation differs depending on glucose concentration has important implications for Axl function in vivo. Several studies point to an important role for Axl in the pathogenesis of diabetic cardiovascular disease. Expression of Axl is increased in injured VSMC, in diabetic mesangial cells and in mesangial cells stimulated with high glucose<sup>13, 14, 39</sup>. Diabetic nephropathy is less severe in Gas6 knockout mice and after treatment of rats with the Gas6 inhibitor warfarin<sup>13</sup>. Axl is also activated by H<sub>2</sub>O<sub>2</sub> in VSMC<sup>15</sup>. Therefore under conditions of oxidative stress, as occurs in diabetes, Axl is likely to be activated. In rat vasculature, Axl is present as 140 kDa, and 114 kDa forms in approximately equivalent ratio<sup>14, 15</sup>. Studies in cultured VSMC suggest that activation of the 140 kDa Axl in HG will result in increased ERK1/2 signaling and VSMC migration. VSMC migration is increased in patients with diabetes and diabetes accelerates the accumulation of VSMC in atherosclerotic lesions<sup>40, 41</sup>. Therefore, this study suggests that altered Gas6-Axl signaling will contribute to vascular dysfunction in diabetes.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgments

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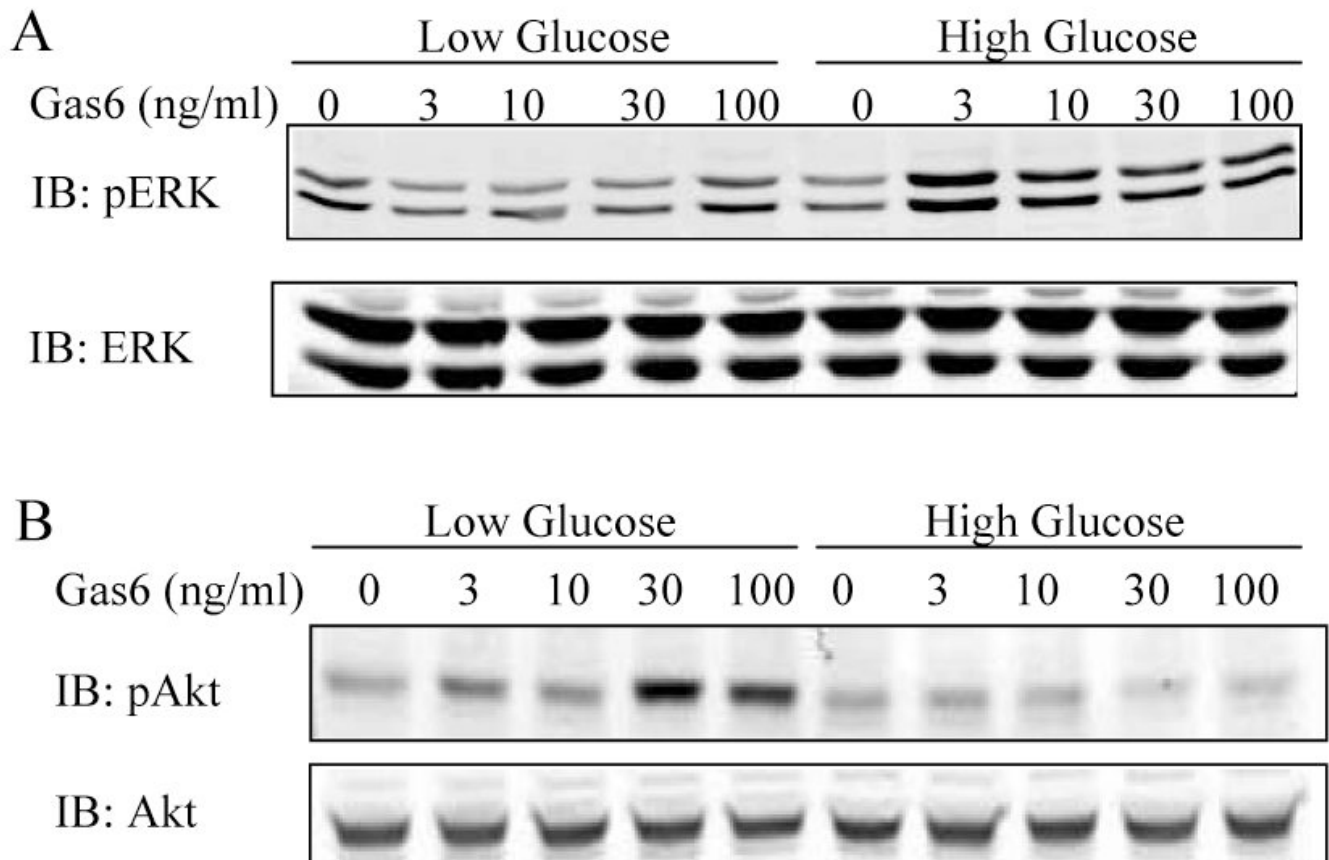
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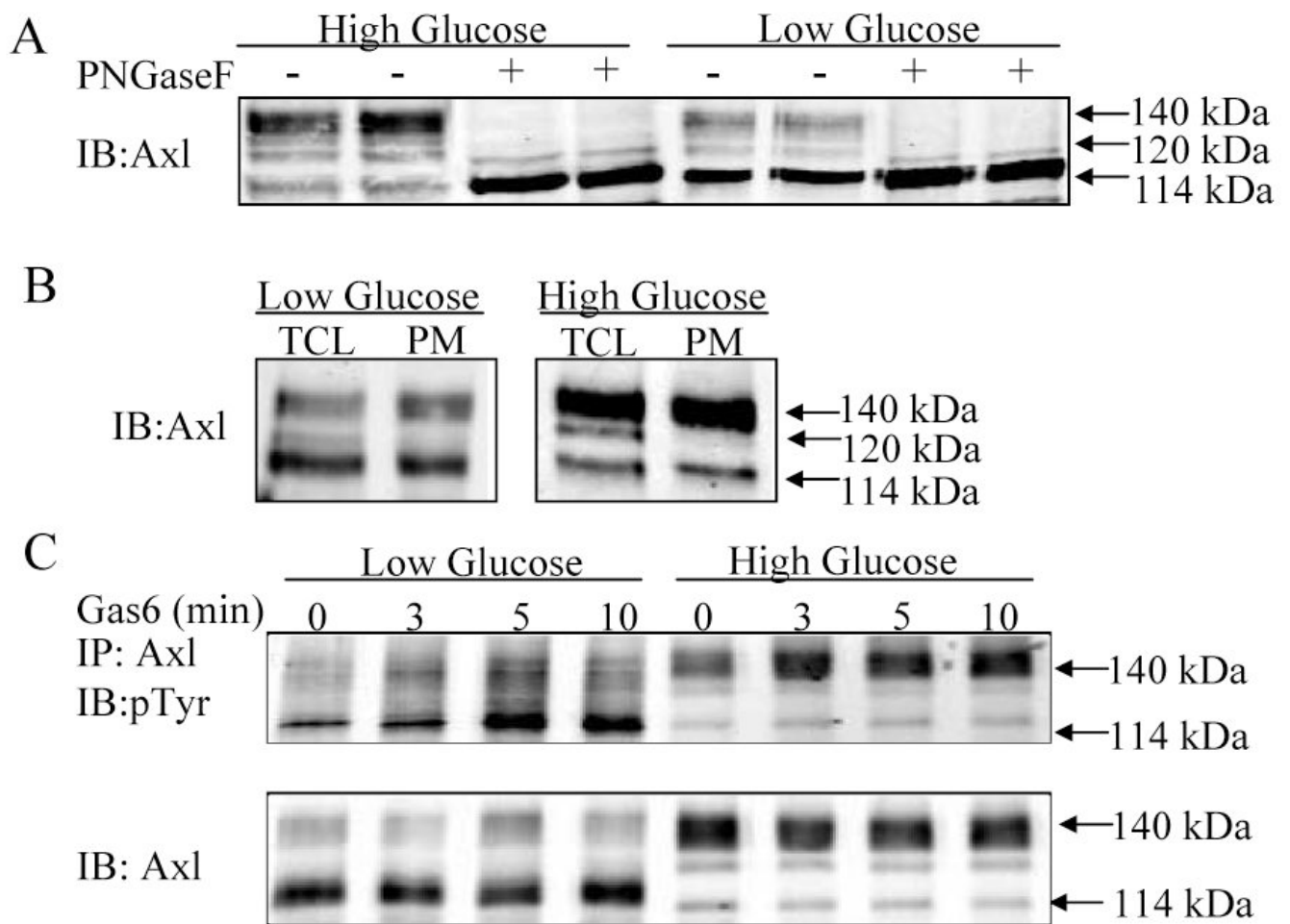
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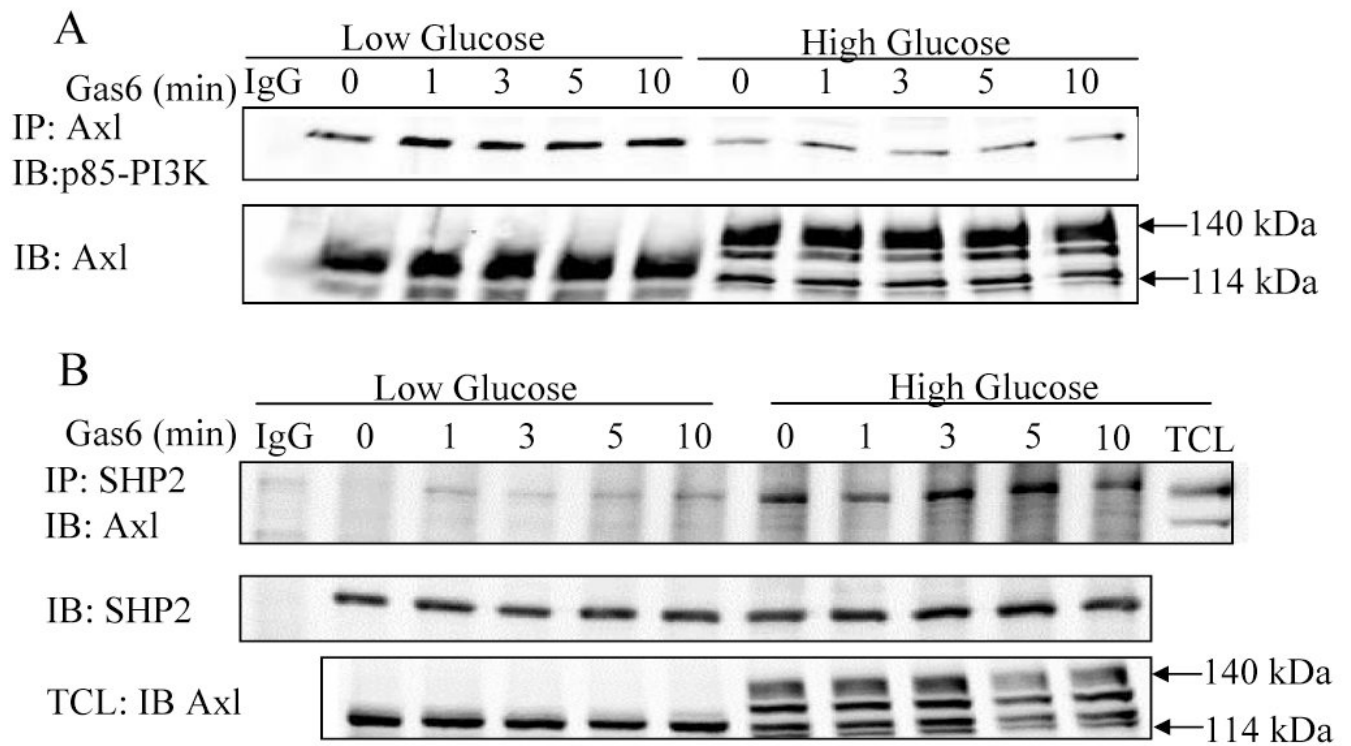
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**Figure 1. Glucose modulates Akt and ERK1/2 phosphorylation upon stimulation of Axl with Gas6**  
VSMC were cultured in serum free LG or HG DMEM for 24hr and treated with increasing Gas6 concentrations for 10 mins. Lysates were immunoblotted with: A. phosphospecific ERK1/2 antibody and reprobred with ERK1/2 antibody; B. phosphospecific Akt antibody and reprobred with Akt antibody.



**Figure 2. Glucose does not alter Axl cell surface localization or phosphorylation by Gas6**  
 VSMC were cultured in serum free LG or HG DMEM for 24hr. A. Axl was immunoprecipitated, treated with or without PNGase-F and immunoblotted with Axl antibody. B. Proteins were biotinylated with NHS-SS-biotin and precipitated with streptavidin agarose. Total cell lysates (TCL) and plasma membrane fractions (PM) were immunoblotted with Axl antibody. C. Following Gas6 treatment Axl was immunoprecipitated. Immunoprecipitates were immunoblotted with phospho-tyrosine antibody (4G10). Results represent n=3.



**Figure 3. Glucose modulates the interaction between Axl and PI3K and Axl and SHP-2**  
 VSMC were cultured in serum free LG or HG DMEM for 24hr. A. Axl was immunoprecipitated from total cell protein and immunoblotted with p85 PI-3 kinase and Axl antibodies. B. SHP-2 was immunoprecipitated from total cell protein and immunoblotted with Axl and SHP-2 antibodies. Lower panel shows total cell lysates (TCL) immunoblotted with Axl antibody. Results represent  $n \geq 2$ .

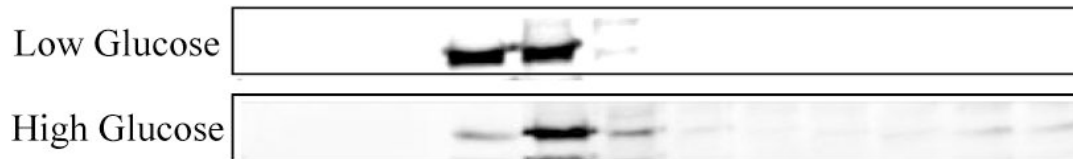
## A Control

IB: Axl

Sucrose % 5 10 25 30 35 45



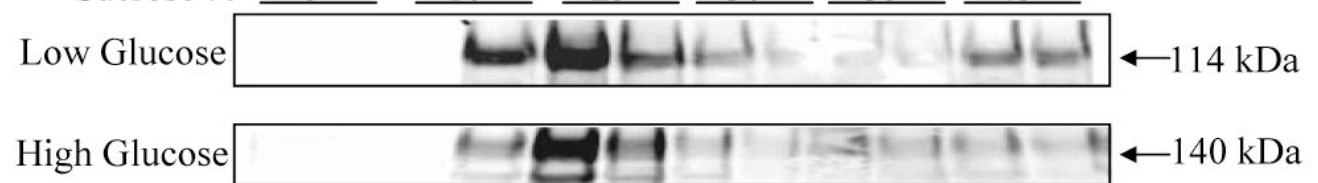
IB: caveolin1



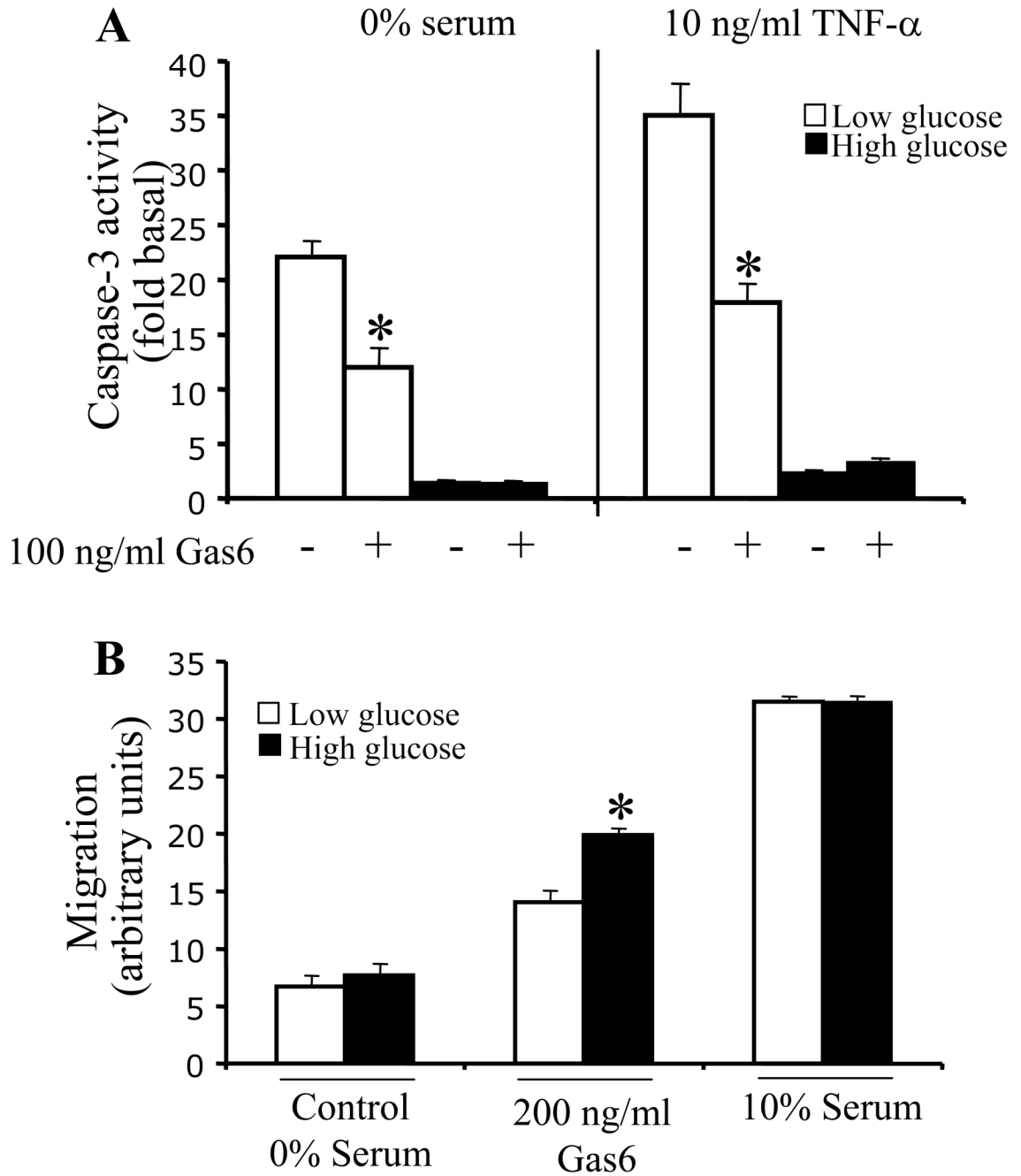
## B Gas6 100 ng/ml (10 min)

IB: Axl

Sucrose % 5 10 25 30 35 45



**Figure 4. Subcellular fractionation of Axl on a sucrose gradient is not altered by glucose**  
 VSMC were cultured in serum free LG or HG DMEM for 24hr. Post-nuclear supernatant was prepared and loaded on a 5-45% sucrose density gradient. Following centrifugation, fractions were collected. 30  $\mu$ l of each fraction was subjected to SDS-PAGE and blotted with the indicated antibodies. Results represent n=3. Cofractionation under (A) basal conditions and (B) after stimulation with Gas6.



**Figure 5. The anti-apoptotic effect of Gas6-Axl is increased in low glucose and the pro-migratory effect of Gas6-Axl is increased in high glucose**

VSMC were cultured in serum free LG or HG DMEM for 24hr. A. Apoptosis was induced using 0%-serum or 0%-serum + TNF- $\alpha$  with and without Gas6 for 24hr. In LG caspase-3 activity was significantly less than control;  $*=p<0.05$ ,  $n=4$ . B. 0%-serum, Gas6 or serum was placed in the lower Boyden chamber. VSMC were seeded in the upper chamber for 6hr. In response to Gas6 in HG VSMC migration was significantly greater compared to LG;  $*=p<0.005$ ,  $n=7$ .