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## Genetic manipulation of periostin expression in the heart does not affect myocyte content, cell cycle activity or cardiac repair

Angela Lorts, Jennifer A. Schwanekamp, John W. Elrod, Michelle A. Sargent, and Jeffery D. Molkenin

Department of Pediatrics, Division of Cardiology (A.L., J.A.S.) and Division of Molecular Cardiovascular Biology (J.W.E., M.A.S., J.D.M.), and the Howard Hughes Medical Institute (J.D.M), University of Cincinnati, Cincinnati Children's Hospital Medical Center, Cincinnati, Ohio 45229.

### Abstract

Following a pathologic insult, the adult mammalian heart undergoes hypertrophic growth and remodeling of the extracellular matrix. While a small sub-population of cardiomyocytes can re-enter the cell cycle following cardiac injury, the myocardium is largely thought to be incapable of significant regeneration. Periostin, an extracellular matrix protein, has recently been proposed to induce re-entry of differentiated cardiomyocytes back into the cell cycle and promote meaningful repair following myocardial infarction. Here, we show that while periostin is induced in the heart following injury, it does not stimulate DNA synthesis, mitosis or cytokinesis of cardiomyocytes *in vitro* or *in vivo*. Mice lacking the gene encoding *periostin* and mice with inducible overexpression of full-length periostin were analyzed at baseline and after myocardial infarction. There was no difference in heart size or a change in cardiomyocyte number in either periostin transgenic or gene-targeted mice at baseline. Quantification of proliferating myocytes in the peri-infarct area showed no difference between periostin overexpressing and null mice compared with strain-matched controls. In support of these observations, neither overexpression of periostin in cell culture, via an adenoviral vector, nor stimulation with recombinant protein induced DNA synthesis, mitosis or cytokinesis. Periostin is a regulator of cardiac remodeling and hypertrophy and may be a reasonable pharmacological target to mitigate heart failure, but manipulation of this protein appears to have no obvious effect on myocardial regeneration.

### Keywords

Fibroblast; remodeling; periostin; cardiomyocyte proliferation; myocardial infarction

### Introduction

Myocardial infarction (MI) leads to an irreversible loss of cardiomyocytes with expansion and remodeling of the extracellular matrix (ECM) and scar formation. This “fibrous scar” lacks contractile capacity and results in adverse changes to chamber geometry, global systolic dysfunction and poor ventricular compliance. If adult cardiomyocytes could proliferate and repopulate the injured area, cardiac function may be rescued, although the ability of cardiomyocytes to proliferate in a physiological meaningful manner remains an area of ongoing

**Correspondence:** Jeffery D. Molkenin Howard Hughes Medical Institute Division of Molecular Cardiovascular Biology Cincinnati Children's Hospital Medical Center 240 Albert Sabin Way Cincinnati, Oh 45229–3039 E-mail: Jeff.Molkenin@cchmc.org.

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controversy.<sup>1-3</sup> Recent reports have proposed that a population of post-mitotic cardiomyocytes, if exposed to the proper environment, may be capable of proliferation.<sup>3-6</sup>

Although a targeted therapy to drive cardiomyocyte proliferation has not been reported, one may be possible given that detectable, albeit extremely small increases in cardiomyocyte cell cycle re-entry can occur after myocardial injury.<sup>3, 5, 7, 8</sup> The number of cardiomyocytes that enter the cell cycle at baseline, evident by markers for DNA synthesis, occurs at a very low frequency of approximately 0.008%.<sup>8</sup> The trigger that drives this small percentage of cells to proliferate has not yet been determined. A recent publication has proposed that the ECM protein periostin leads to the initiation of DNA synthesis, mitosis and ultimately cytokinesis.<sup>9</sup> These results have sparked controversy among those interested in the function of periostin since most other reports have only observed an effect on the ECM and collagen fibrillogenesis.<sup>10, 11</sup>

Periostin is a 90-kDa secreted protein that contains four fasciclin domains<sup>12, 13</sup> and is similar to the insect protein fasciclin I, which is involved in neuronal cell-cell adhesion.<sup>12</sup> Periostin is detectable in the developing heart but is not present in the adult ventricular myocardium at baseline.<sup>14-19</sup> It is re-expressed exclusively by fibroblasts or cells that adopt a fibroblast-like phenotype following an injurious event such as MI or pressure overload stimulation.<sup>20-23</sup> Using transgenic and gene-deleted mice, periostin was shown to affect cardiac hypertrophy and ventricular remodeling.<sup>15</sup> *Periostin* null (*Pn*<sup>-/-</sup>) mice are prone to rupture following MI, but if they survive the first 10 days of this insult, their cardiac morbidity is improved compared to wild-type mice.<sup>15</sup> These data support the hypothesis that periostin orchestrates cellular remodeling and collagen deposition in the heart. Periostin also functions in a similar fashion when secreted by neoplastic cells.<sup>24-26</sup> An increased concentration of periostin correlates with enhanced cellular migration and adhesion and is proportional to the cancer cells ability to invade and metastasize.<sup>24-28</sup>

Although the exact role of periostin has not been defined, the contention that it mediates physiological meaningful myocyte proliferation in the heart following MI is currently a matter of great interest considering the therapeutic ramifications. Here we used transgenic mice to determine if the presence of periostin leads to cardiomyocyte proliferation, or if its absence in gene-targeted mice antithetically reduces this index. We also examined a model of cellular proliferation in cultured neonatal cardiomyocytes, which have a finite capacity for proliferation. In both models we determined that periostin had no effect on any aspect of cell cycle re-entry in cardiomyocytes.

## Materials and Methods

### Animals

All mouse protocols were approved by the Institutional Animal Care and Use Committee of Cincinnati Children's Hospital Medical Center. *Pn*<sup>-/-</sup> and transgenic overexpressors (*Pn*<sup>tTA</sup>) were described previously.<sup>15</sup>

### Western Blot Analysis

Western blot analysis confirmed overexpression of periostin protein in *Pn*<sup>tTA</sup> mice and its absence in *Pn*<sup>-/-</sup> mice. A rabbit polyclonal antibody to periostin was used.<sup>24</sup> Western blotting procedures were described previously.<sup>29</sup>

### Myocardial Infarction and BrdU labeling

MI was performed by permanent ligation of the left coronary artery as previously described.<sup>30</sup> Due to the small size of the *Pn*<sup>-/-</sup> mice, body weight-matched 10–12 week-old *Pn*<sup>-/-</sup> mice (23.4 ± 0.6g) were used for MI while 8 week-old C57BL/6 mice (24.1 ± 0.5g) were used as

controls. Because the growth of the Pn<sup>tTA</sup> mice is not affected, 8–10 week old Pn<sup>tTA</sup> mice and FVB littermates were used for MI. Seven days post MI, mice received intraperitoneal (i.p.) injections with 5-bromodeoxyuridine (BrdU) labeling reagent (1 ml/100g) (Zymed). Four hours after injection organs were harvested, fixed in buffered 10% formalin, embedded in paraffin, and cut into 6 µm sections. Duodenum was harvested and fixed as a positive BrdU labeling control.

### Immunohistochemistry

Sections, 6 µm thick, were deparaffinized and incubated with primary antibody to BrdU (Sigma). Slides were stained with wheat germ agglutinin (WGA)-fluorescein-4-isothiocyanate (FITC) (Sigma) to distinguish the sarcolemma membrane. Nuclei were stained with TO-PRO-3 (Molecular Probes). Fluorescent secondary antibody was used (Molecular Probes) and slides were examined with confocal microscopy. To determine the number of cells positive for phosphorylated histone H3 (Cell Signaling), immunohistochemistry was performed following antigen retrieval with sodium citrate. These slides were also stained with WGA-FITC and TO-PRO-3. The peri-infarct area of four animals, >100,000 cells, were counted per cohort for both BrdU and phosphorylated histone H3. Data were presented as percent of positive cardiomyocyte nuclei versus total cardiomyocyte nuclei.

### Histology and Cell Surface Area Measurements

Hearts were harvested and fixed in 10% buffered formalin, embedded in paraffin and cut into 6 µm sections. Sections were stained for fibrosis with Masson's trichrome or with WGA for the sarcolemma membrane. Measurement of myocyte cross sectional area was counted from both the apex and the center region of the left ventricle, in a minimum of 3 separate hearts from each cohort; >2000 cardiomyocytes were measured. Cross sectional area was calculated using Image J software.

### Recombinant Adenovirus and *in vitro* Proliferation Experiments

A periostin recombinant adenovirus, AdPn, was generated using the full-length mouse periostin, subcloned into the shuttle vector for the Adeasy system (Invitrogen). All *in vitro* experiments were performed in primary neonatal rat cardiomyocytes and fibroblasts, which were isolated from 1- to 2- day old neonates and differentially plated as previously described.<sup>31</sup> The cardiomyocytes were initially plated in 2% fetal bovine serum (FBS) in M199 media on gelatin coated chamber slides and infected/stimulated on day 2, after which the serum was reduced for experimentation of proliferation rates. For adenoviral infection, cardiomyocytes and fibroblasts were incubated for 2-hr with AdPn and AdGFP. The positive control cells were grown in media supplemented with 15% FBS, all other conditions were performed in 0.2% FBS media. Similar methods and conditions were used for the neonatal cardiac fibroblasts. To determine the effect of periostin, cells were infected with AdPn or stimulated daily with recombinant human periostin, a C-terminal truncated version of the full-length protein (BioVendor, Czech Republic). The source of recombinant periostin is the same as used by Kühn and colleagues.<sup>9</sup> AdGFP and vehicle controls were also used under identical conditions. Infection efficiency was examined with AdGFP infection. Forty eight hours following infection, greater than 95% of the cells were GFP positive. A similar dosage of AdPn infection was used to achieve greater than 95% infectivity. To assess proliferation, cells were labeled with BrdU for the final 24 hours of experimentation. To detect DNA synthesis, immunohistochemistry for BrdU was performed, along with cell-type specific markers for cardiomyocytes;  $\alpha$ -actinin (Sigma) and troponin (Cell Signaling), while fibroblasts were detected with vimentin (Sigma) antibody. Further immunohistochemistry was performed with antibodies to phosphorylated histone H3 and aurora B kinase (Abcam) to detect mitosis and cytokinesis, respectively. Fluorophoreconjugated secondary antibodies were provided by

Invitrogen. Proliferating cells in culture were quantified (N>1000) and data presented as percent of positive nuclei versus total nuclei. Each experiment was repeated in quadruplicate.

## Statistics

All data are expressed as mean  $\pm$  SEM. Statistical significance was determined with a paired Student's t-test between 2 groups or one-way ANOVA where appropriate. Probability values of <0.05 were considered statistically significant.

## Results

### Periostin Does Not Induce Neonatal Cardiomyocyte or Fibroblast Proliferation *in vitro*

A previous report concluded that fibroblasts produce periostin to promote neighboring cardiomyocytes to re-enter the cell cycle in the heart.<sup>9</sup> To examine this potential phenomenon in greater detail, *in vitro* experiments were first performed with isolated neonatal cardiomyocytes since they possess some proliferative capacity at baseline, suggesting they would be permissive to cell cycle re-entry. Cardiomyocytes were pulse-labeled with BrdU to detect DNA synthesis. As a positive control, 15% FBS was used, which induced a 2.1-fold increase in BrdU positive cardiomyocyte nuclei after 48 hrs compared with 0.2% FBS cultured myocytes. However, overexpression of periostin by infection with a recombinant adenovirus, AdPn, did not increase DNA synthesis rates compared with AdGFP infection (Figure 1A,B). AdPn treatment for 3 days also did not increase DNA synthesis (data not shown). Treatment with recombinant periostin also did not induce a significant change in DNA synthesis (Figure 1A,B). As neonatal cardiomyocyte preparations are rarely free of fibroblasts, myocytes were identified by expression of  $\alpha$ -actinin and/or troponin. We further characterized cardiomyocyte proliferation by utilizing specific immunohistochemical markers of mitosis and cytokinesis. The percent of nuclei positive for phosphorylated histone H3, a marker for mitosis, was not significantly different between the AdPn infected cells or with recombinant periostin compared with their respective controls (Figure 1C,D). The percent of positive cells treated with recombinant periostin and vehicle were lower than the negative control, possibly due to the pH of the buffer required to reconstitute the protein. Cytokinesis, the final event of proliferation, was detected by aurora B kinase staining. Once again, neither infecting with AdPn or treatment with exogenous periostin significantly increased the percentage of cardiomyocytes undergoing cytokinesis, although serum stimulation was readily effective (Figure 1E,F). To rule out post translational processing as a requirement for bioactivity we also collected conditioned media of myocytes and fibroblasts infected with AdPn to incubate with freshly prepared neonatal myocytes for 48 hrs but, once again, no effect on DNA synthesis or other indexes of proliferation was observed (data not shown). As a control, AdPn infection of cardiomyocytes, or conditioned media from AdPn infected cardiomyocytes both showed expression of full-length periostin by Western blotting (Figure 1G).

To further delineate the role of periostin in inducing cell cycle re-entry, we repeated the above experiments with neonatal cardiac fibroblasts. AdPn and recombinant periostin did not stimulate DNA synthesis in fibroblasts, while the positive control of 15% FBS was efficacious (Figure 2A,B). Phosphorylated histone H3 staining revealed a 3-fold increase in the percent of cells undergoing mitosis in the 15% FBS group, while AdPn and recombinant periostin did not significantly alter this index of cellular proliferation (Figure 2C,D). Furthermore, AdPn and recombinant periostin had no effect on cellular mitosis, as detected by aurora B kinase immunohistochemistry (Figure 2E,F). In the myocardium, fibroblasts and cardiomyocytes co-exist, and are immersed in the same milieu of growth factors and proteins, hence additional co-culture experiments were performed. Neonatal rat cardiac fibroblasts were infected with AdPn and were subsequently cultured with neonatal rat cardiomyocytes. Markers of cell cycle re-entry, (BrdU, phosphorylated histone H3 and aurora B kinase) were examined and revealed

no significant change in the quantity of cells proliferating (data not shown). Collectively, these data indicate that periostin cannot induce cell cycle re-entry in two different cell types that are normally permissive to some level of proliferation.

### Periostin Null and Overexpressing Mice Do Not Have Altered Myocyte Numbers in the Heart at Baseline

We also reasoned that if periostin directly regulates myocyte proliferation it was most likely to do so throughout development when these cells are readily dividing. Indeed, a large number of genetically modified mouse models with altered cell cycle regulatory proteins have been created that display significant changes in total myocyte cell numbers due to a developmental effect of these proteins.<sup>32-35</sup> However, baseline characterization of the  $Pn^{-/-}$  and periostin overexpressing transgenic ( $Pn^{tTA}$ ) mice at 6 weeks of age showed no difference in total cell number in the heart. At 6 weeks of age periostin is not detectable in the heart of the wildtype or  $Pn^{-/-}$  mice (data not shown) but is abundantly present in the hearts of the  $Pn^{tTA}$  mice (Figure 3A). Importantly, the heart weight to body weight ratio was not altered in the  $Pn^{-/-}$  animals ( $4.3 \pm 0.5$  mg/g) versus age-matched controls ( $4.2 \pm 0.4$  mg/g) (Figure 3B) or  $Pn^{tTA}$  transgenic animals ( $5.1 \pm 0.2$  mg/g) versus appropriate controls ( $4.8 \pm 0.3$  mg/g).  $Pn^{tTA}$  mice at 6 months of age also showed no significant difference in heart size normalized to body weight ( $5.4 \pm 0.2$  mg/g) when compared to controls ( $5.3 \pm 0.3$ , Figure 3C). To calculate total myocyte number in the heart the cross sectional area of myocytes measured from ventricular histological sections was analyzed at 6 weeks of age and determined to be similar between the  $Pn^{-/-}$ ,  $Pn^{tTA}$ , and their controls (Figure 3D,E). Cell number and myocyte fraction was assessed by immunohistochemistry and once again no differences were noted between any of the groups (data not shown). Additionally, the degree of fibrosis was not significantly different in the hearts of 6-week-old,  $Pn^{tTA}$  mice and age-matched controls as assessed by Masson's trichrome staining of histological sections (data not shown).

### Periostin Does Not Induce Cardiomyocyte Proliferation *in vivo* Following MI

Following myocardial injury there is abundant re-expression and accumulation of periostin in the interstitial space within and surrounding the injured area.<sup>15</sup> Periostin protein is detected as early as 4 days after MI and peaks at 7 days, and remains detectable for up to 8 weeks.<sup>15, 16</sup> To examine if periostin expression and accumulation in the ECM stimulates cardiomyocyte cell cycle re-entry *in vivo*, MI injury was induced in both the  $Pn^{-/-}$  and  $Pn^{tTA}$  transgenic mice, and their appropriate wild type controls. There was no difference in infarct size between the  $Pn^{-/-}$  mice, the  $Pn^{tTA}$  mice, and their respective controls in agreement with our previous results (data not shown).<sup>15</sup> Histological analysis was performed 7 days after MI, when periostin expression is maximal (Figure 4A). Four hours prior to harvesting the heart, animals underwent an intraperitoneal injection of BrdU to label cardiomyocytes that were undergoing DNA synthesis. On examination of the peri-infarct area of control mice (C57BL/6), consistent with published data, there was a very small number, 0.035%, of BrdU positive cardiomyocyte nuclei.  $Pn^{-/-}$  and  $Pn^{tTA}$  transgenic mice did not show a significant change in cardiomyocyte DNA synthesis (Figure 4B). A representative immunohistochemical image from the infarct border zone is shown with cells outlined with WGA-FITC and BrdU labeling, which revealed nonmyocytes (asterisk) and a single labeled cardiomyocyte (arrow) as undergoing DNA synthesis (Figure 4B). To further support these results, immunohistochemical analysis was done to detect phosphorylated histone H3 staining as evidence of mitosis. Only 0.012% of myocyte nuclei were positive for phosphorylated histone H3 in the peri-infarct area. There was no significant increase in the percent of positive nuclei in either  $Pn^{-/-}$  or  $Pn^{tTA}$  mice (Figure 4C). A representative immunohistochemical image from the infarct border zone is shown with cells outlined with WGA-FITC and phosphorylated histone H3 labeling, which revealed a labeled non-myocyte (asterisk) and cardiomyocyte (arrow) (Figure 4C). A minimum of 4 mice

was analyzed in each group. These results indicate that overexpression of periostin or its loss does not appreciably affect myocyte cell cycle activity in the peri-infarct region.

## Discussion

While adult cardiac myocytes are not thought to proliferate *in vivo* given their highly differentiated state, a finite population are proposed to re-enter the cell cycle following experimental MI.<sup>8</sup> Such myocytes are typically located in the peri-infarct area and are presumably responding to external stimuli, such as growth factors or changes in ECM composition. Our rates of endogenous cardiomyocyte cell cycle activity in the peri-infarct region of hearts after MI were comparable to previous accounts in the literature (0.005% – 0.04%)<sup>5, 8, 36</sup>, which given the extremely low value is unlikely to contribute to cardiac regeneration in a meaningful manner. We also employed an *in vitro* model in which neonatal cardiomyocytes were isolated from 1–2 day old rat pups. This model was selected because neonatal myocytes inherently possess some degree of cell cycle activity, suggesting they are physiologically primed to respond to proliferative signals if they occur. Indeed, serum stimulation promoted a greater than 2-fold activation of cell cycle activity in these myocytes. However, in neither context did periostin alter myocyte cell cycle activity or proliferation.

As a secreted ECM protein that is associated with areas of fibrosis, remodeling and inflammation, periostin can directly interact with other ECM proteins such as fibronectin, tenascin-C, collagen I, collagen V, and heparin.<sup>10, 11, 23</sup> Periostin serves as a ligand for select integrins, such as  $\alpha_5\beta_3$ ,  $\alpha_v\beta_5$ , and  $\alpha_4\beta_6$ , where it can affect the ability of cells (fibroblasts or cancer cells) to migrate and/or undergo a epithelial-mesenchymal transition (EMT) in select tissues.<sup>24, 25</sup> MI or pressure overload stimulation to the adult heart induces abundant re-expression of periostin from resident fibroblasts located between myocytes within the heart parenchyma proper, where prior to such stimulation the only observed expression of periostin in the heart is within the collagen rich environment of the valves.<sup>14-19</sup> Periostin expression is also induced at sites of vascular injury,<sup>37, 38</sup> in the lung after injury/fibrosis,<sup>23, 39</sup> in and around tumors,<sup>39</sup> and at wound sites.<sup>40</sup> While not unequivocal, these various studies suggest a common biologic response whereby periostin becomes re-expressed at injury or inflammatory sites within the adult organism, often associated with a need for ECM remodeling and cellular migration, not unlike many developmental processes associated with periostin expression.

The function of periostin upon re-expression in the injured myocardium is currently an area of controversy given disparate accounts in the literature. We originally reported that  $Pn^{-/-}$  mice had an altered fibrotic response and were unable to form proper scars after MI injury, leading to greater rates of LV rupture and death.<sup>15</sup> This result was independently confirmed the following year in separately generated  $Pn$  null mice.<sup>16</sup> Conversely, we generated transgenic mice overexpressing full-length periostin in the heart, which were protected from wall rupture following MI injury.<sup>15</sup> Abundant periostin overexpression in the heart did not induce fibrosis, although it did eventually promote a mild hypertrophic response in aged mice (32 weeks). Another interesting aspect of  $Pn^{-/-}$  mice is that they are less susceptible to fibrotic heart disease associated with long-term pressure overload or MI, resulting in better ventricular performance.<sup>15</sup> These results suggest that loss of periostin is protective to the heart long-term, provided the initial insult is survived.

An entirely novel function for periostin was recently reported whereby cardiomyocyte proliferation and the repair of injured myocardium was shown to be regulated by this protein.<sup>9</sup> In addition to reporting increased proliferation, Kühn et al. showed that periostin expression prevented fibrosis and reduced cardiac hypertrophy after an injury event, results that are not consistent with other reports.<sup>15</sup> A potential reason for these differences may be that Kühn et

al. utilized a recombinant truncated form of periostin (amino acids 22–669 of 811) in ascertaining biological effects.<sup>9</sup> Using this abridged protein, as well as a full-length protein, the authors reported increased proliferation rates in adult rat cardiac myocytes cultured for a total of 12 days.<sup>9</sup> These later results may also be influenced by the known process of de-differentiation that occurs in adult myocytes cultured for this length of time. We failed to observe any induction of cell cycle activity in a model system that is known to be responsive to proliferative signals, neonatal myocytes in culture for 2 or 3 days.

More remarkably, Kühn et al. reported that application of the truncated form of periostin with a gelfoam patch onto infarcted rat hearts reduced fibrosis, reduced cardiac hypertrophy, increased myocyte proliferation, and improved cardiac function.<sup>9</sup> However, overexpression of full-length periostin in the mouse heart by transgenesis (via the well-characterized  $\alpha$ -myosin heavy chain promoter), which results in the proper accumulation of periostin protein in the ECM without detectable intracellular retention, did not increase myocyte number at baseline, nor did it augment indices associated with cardiac repair after MI injury.<sup>15</sup> However, one potential difference with our study is that the Pn transgene was used in a constitutive manner so that periostin protein was always present, while Kühn et al. applied recombinant periostin acutely. We could not mimic this acute affect because the inducible Pn transgene takes 4–6 weeks to achieve reasonable expression due to the time it takes doxycycline to clear the mouse's system. Also to be considered, we did not observe differences in infarct size in any of the experimental or control groups, suggesting that the presence of the tetracycline transactivator was not a confounding variable (data not shown). Another potential difference is that we used the mouse while Kühn et al used the rat for infarct studies. Finally, our transgenic mice expressed only one isoform of periostin, of which there are many alternately spliced versions present in the heart<sup>15</sup>. Thus, it is possible that another spliced form of periostin could have cell cycle promoting activity that is akin to the truncated version of periostin employed by Kühn et al.<sup>9</sup>

While our data are negative concerning a role for periostin in regulating myocyte proliferation, we believe that they are credible given the nature of the model systems employed and our consistent use of positive and negative controls. First, we carefully assessed the possibility that lack of periostin or constitutive expression of periostin throughout embryonic and postnatal development would alter myocyte numbers in the heart, although no effect was observed. By comparison, transgene mediated overexpression of calmodulin in the developing mouse heart promoted a 73% increase in myocyte number by embryonic day 19 and a 40% increase in myocyte number in the adult heart.<sup>32</sup> Overexpression of either insulin-like growth factor 1 in the heart or Bcl-2 increased myocyte number as measured by assessment of myocyte size normalized to heart weight and volume fractions.<sup>33, 34</sup> Overexpression of telomerase in the heart by transgenesis also produced greater myocyte content in the heart with some hypertrophy.<sup>35</sup> Second, neither  $Pn^{-/-}$  mice nor periostin overexpressing transgenic mice showed any difference in stimulated cell cycle activity in myocytes within the peri-infarct zone of the adult mouse heart. Third, application of periostin by multiple methods was without effect on proliferation and cell cycle activity in cultured neonatal cardiomyocytes. Forth, Shimazaki, et al. also noted no difference in Ki67 positive cells in the infarct border zone of  $Pn^{-/-}$  mice.<sup>16</sup> In conclusion, while real differences in experimental conditions may have underlied an inability to detect proliferation and regeneration in our mice, we favor the alternate hypothesis that periostin mostly functions in collagen fibrillogenesis, ventricular remodeling, and in facilitating cellular movements in areas of injury or ongoing EMT.

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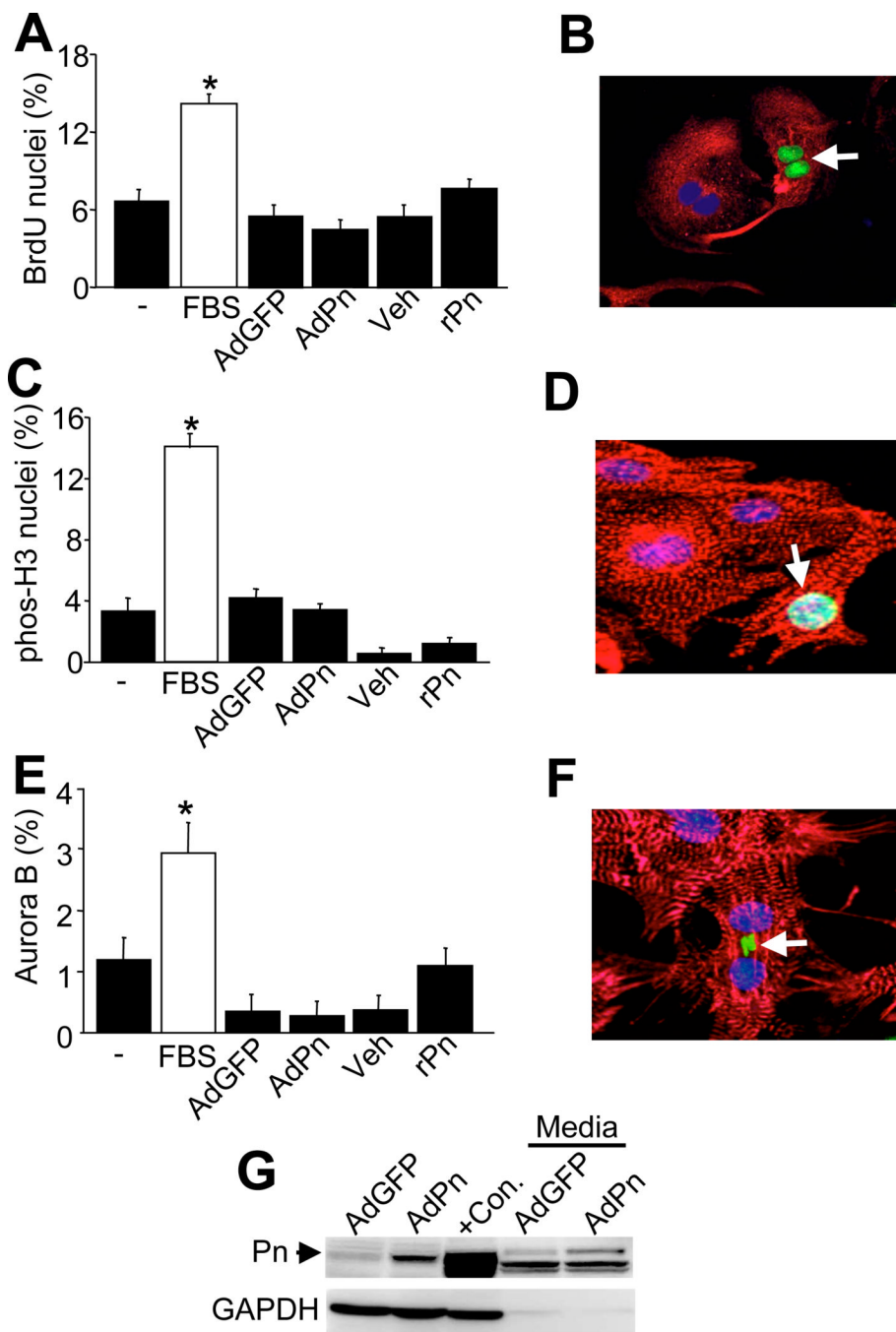
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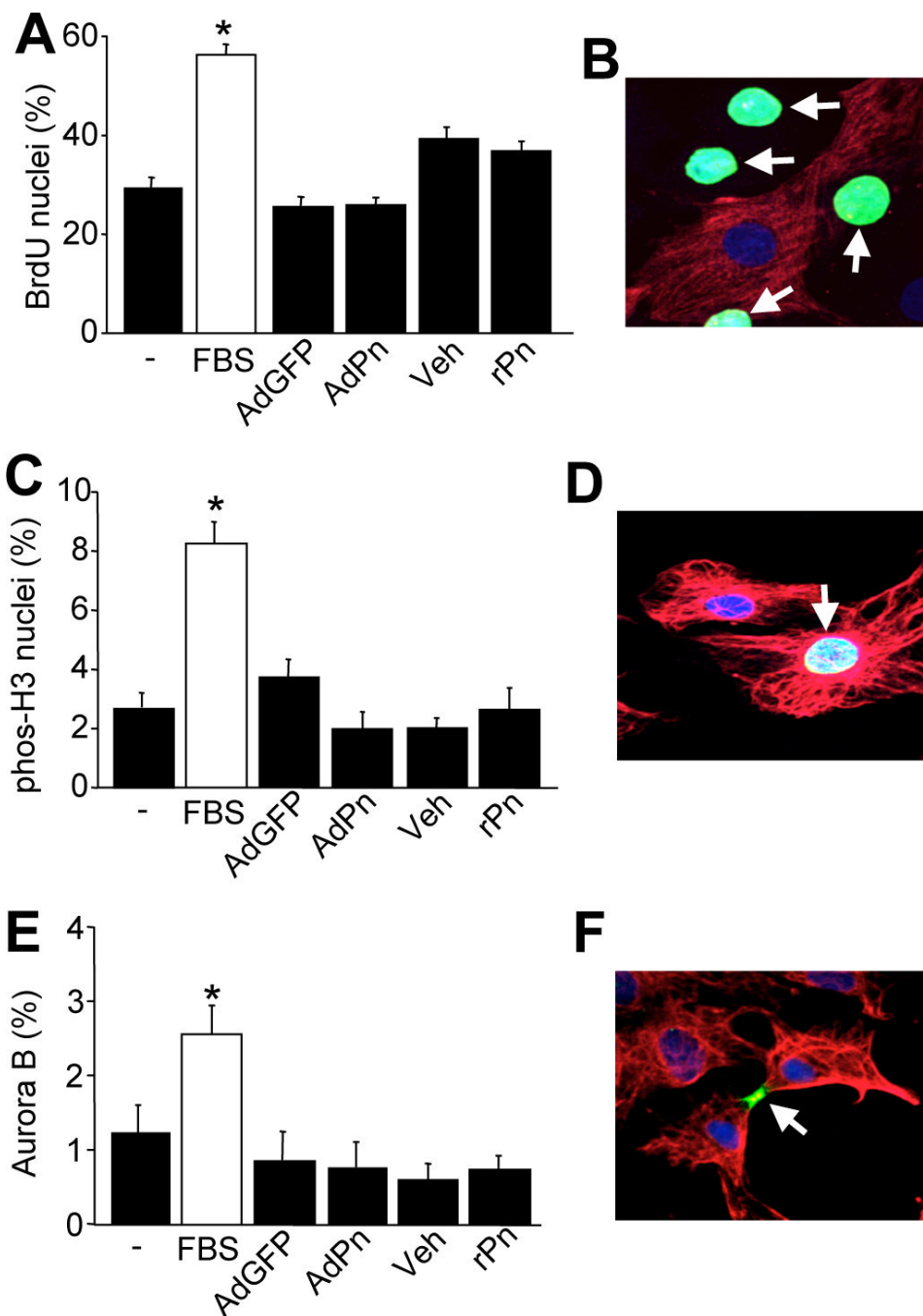
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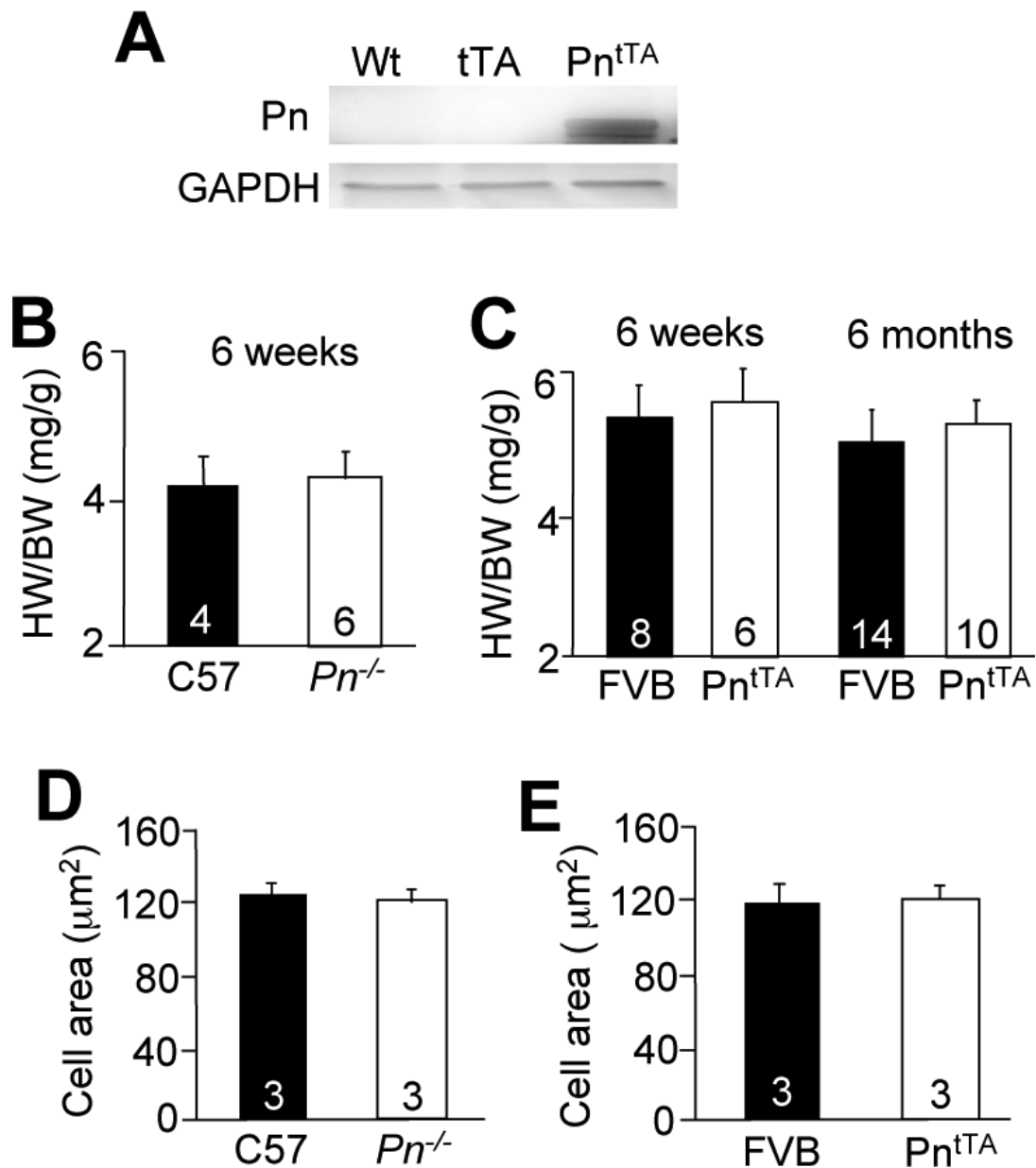
**Figure 1.** Neonatal cardiomyocyte proliferation is not altered by periostin. **A.** Immunocytochemistry detection of BrdU incorporation in neonatal cardiomyocytes versus total cardiomyocyte nuclei. Positive control was 15% FBS and the negative control (-) was no stimulation. **B.** Image of representative BrdU positive cardiomyocyte nuclei. Red is troponin (myocyte), blue is nuclei, and green is BrdU (arrow). **C** and **D,** Immunocytochemical quantification and representative image of cardiomyocyte nuclei positive for phosphorylated histone H3. Red is troponin (myocyte), blue is nuclei, and green is phosphorylated histone H3 (arrow). **E** and **F,** Immunocytochemical quantification and representative image of cardiomyocyte nuclei positive for aurora B kinase staining compared to total cardiomyocyte nuclei. Red is troponin

(myocyte), blue is nuclei, and green is aurora B kinase (arrow). G, Western blot for periostin protein from AdPn infected cardiomyocytes versus AdGFP infection, or concentrated conditioned media 48 hrs after AdPn or AdGFP infection. The positive control (+Con) is protein extract from 2 day-old neonatal feet. \*P<0.05 positive vs. negative control. Abbreviations: rPn, recombinant periostin; Pn, periostin; veh, vehicle.



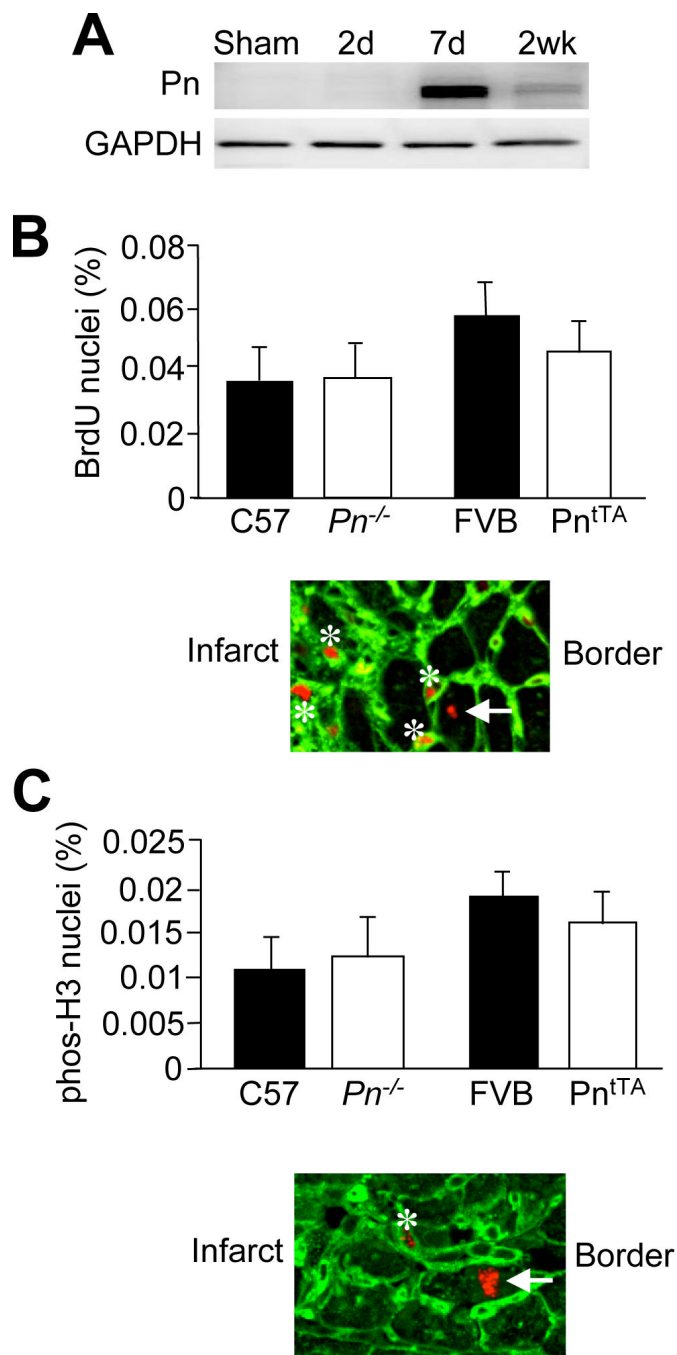
**Figure 2.** Neonatal fibroblast proliferation is not altered by periostin. A, Immunocytochemistry detection of BrdU incorporation in primary fibroblasts versus total fibroblast nuclei. The positive control was 15% FBS and the negative control (-) was no stimulation. B, Image of a representative BrdU positive fibroblast nuclei. Red is troponin to show contaminating myocytes, blue is nuclei, and green is BrdU (arrows). C and D, Immunocytochemical quantification and representative image of fibroblast nuclei positive for phosphorylated histone H3. Red is vimentin (fibroblast), blue is nuclei, and green is phosphorylated histone H3 (arrow). E and F, Immunocytochemical quantification and representative image of fibroblast nuclei positive for aurora B kinase staining compared to total fibroblast nuclei. Red is vimentin (fibroblast), blue is nuclei, and

green is aurora B kinase (arrow). \*P<0.05 positive vs. negative control. Abbreviations: rPn, recombinant periostin; Pn, periostin; veh, vehicle.



**Figure 3.**

Baseline myocyte content is not different in the hearts of *Pn*<sup>-/-</sup> and *Pn* transgenic mice. A, Western blot for periostin (*Pn*) from hearts of 6 week-old FVB, tTA and *Pn*<sup>tTA</sup> mice. GAPDH is a loading control. B, Assessment of heart weight (HW) normalized to body weight (BW) in *Pn*<sup>-/-</sup> versus wild type (C57BL/6) mice at 6 weeks of age. C, HW/BW in *Pn*<sup>tTA</sup> and wildtype (FVB) at 6 weeks and 6 months of age. D and E, Assessment of myocyte cross sectional cell areas from left ventricular histological sections of the indicated genotypes (n=3 hearts each, with a minimum of 2000 cells counted in total).



**Figure 4.** Assessment of cardiomyocyte proliferation 7 days after MI in *Pn*<sup>-/-</sup> and *Pn*<sup>tTA</sup> transgenic mice. A, Western blot analysis of periostin in the heart following MI. GAPDH is a loading control. B, Percent of BrdU positive cardiomyocyte nuclei when compared to total cardiomyocyte nuclei in the peri-infarct area of *Pn*<sup>-/-</sup>, *Pn*<sup>tTA</sup> mice and their appropriate controls. Below the graph a representative immunohistological image is shown with labeled non-myocytes (asterisk) and a labeled myocyte (arrow). C, Percent of phosphorylated histone H3 positive cardiomyocytes compared to total cardiomyocytes in the peri-infarct area of *Pn*<sup>-/-</sup>, *Pn*<sup>tTA</sup> mice and their appropriate controls. Below the graph a representative immunohistological image is shown with a labeled non-myocyte (asterisk) and a labeled myocyte (arrow). Experiments were



performed on a minimum of 4 hearts per cohort. Greater than 100,000 cells were counted per experimental group. None of the groups were significantly different.