

# ***IGF-1* Deletion Affects Renal Sympathetic Nerve Activity, Left Ventricular Dysfunction, and Renal Function in DOCA-Salt Hypertensive Mice**

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## **Summary**

The objective of the paper is to determine the influence of *IGF-1* deletion on renal sympathetic nerve activity (RSNA), left ventricular dysfunction, and renal function in deoxycorticosterone acetate (DOCA)-salt hypertensive mice. The DOCA-salt hypertensive mice models were constructed and the experiment was classified into WT (Wild-type mice) +sham, LID (Liver-specific *IGF-1* deficient mice) + sham, WT + DOCA, and LID + DOCA groups. Enzyme-linked immunosorbent assay (ELISA) was used to detect the serum *IGF-1* levels in mice. The plasma norepinephrine (NE), urine protein, urea nitrogen and creatinine, as well as RSNA were measured. Echocardiography was performed to assess left ventricular dysfunction, and HE staining to observe the pathological changes in renal tissue of mice. DOCA-salt induction time-dependently increased the systolic blood pressure (SBP) of mice, especially in DOCA-salt LID mice. Besides, the serum *IGF-1* levels in WT mice were decreased after DOCA-salt induction. In addition, the plasma NE concentration and NE spillover, urinary protein, urea nitrogen, creatinine and RSNA were remarkably elevated with severe left ventricular dysfunction, but the creatinine clearance was reduced in DOCA-salt mice, and these similar changes were obvious in DOCA-salt mice with *IGF-1* deletion. Moreover, the DOCA-salt mice had tubular ectasia, glomerular fibrosis, interstitial cell infiltration, and increased arterial wall thickness, and the DOCA-salt LID mice were more serious in those aspects. Deletion of *IGF-1* may lead to enhanced RSNA in DOCA-salt hypertensive mice, thereby further aggravating left ventricular dysfunction and renal damage.

## **Key words**

*IGF-1* • DOCA-salt hypertension • Renal sympathetic nerve activity • Left ventricular dysfunction • Renal impairment

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## **Introduction**

Hypertension, as a crucial risk factor for cardiovascular diseases, refers to a clinical syndrome featured by the persistent increase of blood pressure (Tapia-Castillo *et al.* 2014). Unfortunately, no special symptoms are shown at the early stages of hypertension, but when observed, the blood pressure has often reached the standard level for clinical hypertension (Lake-Bruse and Sigmund, 2000). To our knowledge, hypertension is a multifactorial disease, which is possibly related to patients' renal function, renal sympathetic nervous system, high-salt intake, and other factors (Fujita, 2014). Salt sensitivity has been estimated to be exhibited in approximately 51 % of the world's adult population suffering from hypertension (Armando *et al.* 2015, Zhang *et al.* 2016). Thus, it is valuable to explore the unknown etiology of hypertension via an animal model, like the deoxycorticosterone acetate (DOCA)-salt model of hypertension.

Up till now, increasing attention has been

focused on the insulin-like growth factor system (IGFs) in the development and progression of cardiovascular diseases (Heald *et al.* 2003). As a group of polypeptides with the structure highly homologous to insulin, IGFs includes IGF-1, IGF-2, and receptors on the cell membrane (IGF-1R and IGF-2R), as well as insulin-like growth factor-binding proteins (IGFBP) (Kota *et al.* 2012). As for IGF-1, it is a single-chain polypeptide consisting of 70 amino acids, mainly synthesized and secreted by liver cells, and circulating in the blood (Anderle *et al.* 2002). IGF-1, one of the most crucial regulators of growth, has vasodilatory effects, and causes partial reversion of hypertension-induced changes in cardiac function (Casella *et al.* 2010, Tanaka *et al.* 2018). It can promote the proliferation of vascular smooth muscle cells (VSMCs) (Shuang *et al.* 2018), consequently resulting in hypertension (Perros *et al.* 2007). Likewise, some researchers also revealed a significantly reduced expression of IGF-1 in the myocardial tissue and serum of patients with hypertension (Fernandez-Sola *et al.* 2015, Zhang *et al.* 2011). This is inconsistent with the above mentioned results, indicating a controversial role of IGF-1 in hypertension pathogenesis. In this paper, we used the liver-specific IGF-1 deficient (LID) mice model subjected to DOCA-salt treatment, a commonly used intervention to induce hypertension (Imenshahidi *et al.* 2014) and we investigated the effect of IGF-1 deletion on the left ventricular dysfunction, RSNA and renal function in hypertensive mice.

## Materials and Methods

### Ethics statement

This study was in line with the Guide for the Care and Use of Laboratory Animals published by National Research Council (2011) and all animal experiments were carried out under the supervision of the Ethics Committee for Medical Laboratory Animals in Hebei Medical University (IACUC-Hebmu-2018056).

### Laboratory animal

Male C57BL/6 WT (Wild type) mice (4 months of age) with a homozygous floxed exon 4 of the IGF-1 gene ( $\text{IGF-1}^{\text{flx/flx}}$ ) were bought from Jackson Laboratories (Bar Harbor, ME, USA). The LID mice in this experiment were obtained by Cre/loxP recombination strategy, as described in a previous study (Tarantini *et al.* 2017). All mice were housed under specific pathogen-

free conditions with access to standard rodent chow and water on a 12-h light-dark cycle.

### *DOCA-salt sensitive hypertension model construction and experimental grouping*

There were 4 groups (with 20 mice in each group) in this experiment, including WT + sham, LID + sham, WT + DOCA, LID + DOCA groups. Mice in DOCA groups were anesthetized by an intraperitoneal injection of chloral hydrate (400 mg/kg). Next, the left kidney of mice was exposed from the abdominal cavity, and the left artery, vein and ureter were carefully separated with microsurgical forceps, followed by the electrocoagulation of the left kidney artery and vein. After that, the eye scissors were used to cut the ligated renal artery, vein and ureter along the distal end of the ligation thrum for left kidney resection. When post-operative bleeding stopped, the muscle and skin of mice were sutured in turn, and the mice received postoperative intraperitoneal injection of penicillin (100,000 units/day) for 3 days to prevent infection. One week later, the mice were subcutaneously injected with DOCA by 50 mg/kg (once a week for 5 consecutive weeks), and drank water containing 1 % NaCl and 0.2 % KCl. By contrast, mice in sham groups only received uninephrectomy and given normal tap water for drinking. The systolic blood pressure (SBP) was determined via a non-invasive tail cuff method (Kent Scientific, Torrington, CT; model XBP 1000) once a week for 4 consecutive weeks. Before experiment, mice were acclimated for 2 weeks to the blood pressure measuring device.

### *Detection of serum IGF-1 levels*

After the determination of SBP on the 28<sup>th</sup> day of the experiment, blood samples were immediately collected from the submandibular vein of mice and centrifuged at 4 °C with the centrifugal force of 2500 g for 20 min. The obtained serum was preserved at -80 °C for later use. IGF-1 concentration in the serum samples was measured by the enzyme-linked immunosorbent assay (ELISA) (R&D Systems, Minneapolis, MN, USA).

### *Norepinephrine (NE) spillover*

The blood sample (1 ml) was immediately obtained and preserved at -80 °C after the determination of SBP on the 28th day of the experiment. Solid phase extraction and high-performance liquid chromatography with electrochemical detector (HPLC-ECD) (ESA Biosciences, Inc, Chelmsford, MA, USA) were employed

to determine the NE concentration in plasma. Levo-[ring-2, 5, 6-<sup>3</sup>H]NE (Perkin-Elmer, Boston, MA, USA, 0.13  $\mu\text{Ci} \cdot \text{min}^{-1} \cdot \text{kg}^{-1}$ ) were infused intravenously at the rate of 16  $\mu\text{l}/\text{min}$  for 90 min to maintain the stable plasma concentration of <sup>3</sup>H-NE (King *et al.* 2008). NE clearance and spillover were calculated as described previously (Kandlikar and Fink 2011).

#### *Urine biochemical test*

After the determination of SBP, mice in each group were accessed to water and food for 24 h, and then the urine samples were collected with the toluene anticorrosion (1%). Next, 1.5 ml of urine samples was determined for urinary protein, urea nitrogen and creatinine by using an automatic biochemical analyzer.

#### *Echocardiography examination*

After the determination of SBP on the 28th day of the experiment, five mice were randomly selected from each group for intraperitoneal anesthesia with chloral hydrate (400 mg/kg), fixed on the operation table, and a horizontal skin incision was made in the midline of the neck to separate the subcutaneous tissue. The right common carotid artery was exposed and ligated at the distal end, with a vascular clamp at the proximal end to block the bleeding. Then, the Vevo 2100 ultrasonic machine (Visual Sonics, Toronto, Canada) was utilized to measure the interventricular septal thickness in diastole (IVSd), interventricular septal thickness in systole (IVSs), left ventricular posterior wall thickness in diastole (LVPWd), left ventricular posterior wall thickness in systole (LVPWs), left ventricular anterior wall thickness in diastole (LVAWd), left ventricular anterior wall thickness in systole (LVAWs), left ventricular internal diameter in diastole (LVIDd), and left ventricular internal diameter in systole (LVIDs).

#### *Measurement of renal sympathetic nerve activity (RSNA)*

On the 28<sup>th</sup> day of the experiment, another five mice were randomly selected from each group to expose their left kidney by cutting the skin along an extraperitoneal tunnel running from the incision to the left flank of the abdominal cavity. In the abdominal aorta area, left kidney sympathetic nerves were carefully separated with a glass needle, which was immersed in warm liquid paraffin. Meanwhile, a bipolar silver electrode was used to record RSNA beneath the renal nerve, which was amplified by a differential amplifier and recorded by Powerlab® 8/30 to obtain the raw RSNA

and the integrated RSNA. The RSNA was evaluated based on the previous study (Fardin *et al.* 2012).

#### *HE staining*

Further five mice were selected from each group for intraperitoneal anesthesia with 10 % chloral hydrate (400 mg/kg) and cut the skin along the median line to open the abdomen and thoracic cavity. The kidney was fully exposed and taken out, rinsed with normal saline, fixed in 10 % formaldehyde solution, dehydrated routinely, embedded with paraffin, and cut into 4  $\mu\text{m}$  thick slices. After the slices were deparaffinized, they were stained in hematoxylin for 5–10 min, immersed in 0.5 % hydrochloric acid in the presence of 70 % ethanol for 30–60 s, sunk in alkaline solution, washed with distilled water for 1 min, and stained in eosin for 30–60 s. Then, slices were dehydrated with gradient alcohol, hyalinized with xylene, mounted in neutral gum, and observed under a microscope.

#### *Statistical analysis*

All statistical data were analyzed with the statistical software SPSS. The measurement data were presented by mean  $\pm$  SD. The independent sample *t*-test was applied to compare difference between two groups, while among multiple groups compared by One-Way ANOVA. Besides, the inter-group difference was analyzed by the least significant difference (LSD) test.  $P<0.05$  indicated the statistical significance.

## **Results**

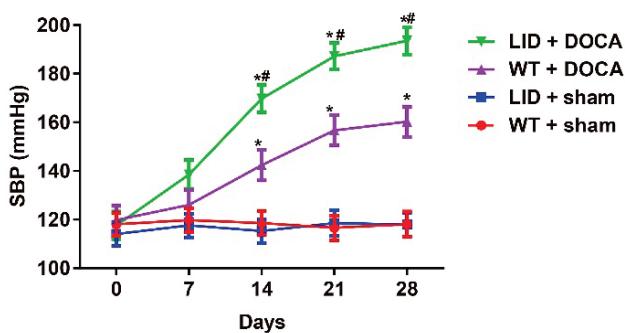
#### *Deletion of IGF-1 gene resulted in the increased SBP in DOCA-salt hypertensive mice*

According to our results, DOCA-salt treatment could time-dependently increase the SBP of mice. SBP was significantly higher in mice from WT + DOCA group and LID + DOCA group than in those from WT + sham group and LID + sham group on the 14<sup>th</sup>, 21<sup>st</sup>, and 28<sup>th</sup> day of the DOCA-salt treatment, respectively (all  $P<0.05$ ). Moreover, mice in the LID + DOCA group had remarkably elevated SBP, as compared with those in the WT + DOCA group on the 14<sup>th</sup>, 21<sup>st</sup>, and 28<sup>th</sup> day (all  $P<0.05$ , Fig. 1).

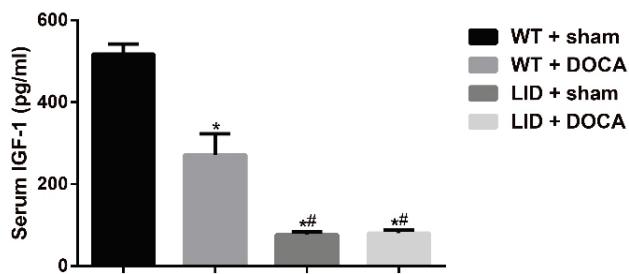
#### *Serum IGF-1 levels of mice in four groups*

Figure 2 shows that WT + DOCA group had lower serum IGF-1 level compared to WT + sham group ( $P<0.05$ ). Meanwhile, there were very low serum levels

of IGF-1 in LID mice but no difference between the LID + sham group and the LID + DOCA group ( $P > 0.05$ ).



**Fig. 1.** Deletion of IGF-1 gene can significantly increase SBP in mice induced by DOCA-salt. \*,  $P < 0.05$  compared with sham groups; #,  $P < 0.05$  compared with WT + DOCA group.



**Fig. 2.** Serum IGF-1 levels of mice among four groups detected by ELISA. \*,  $P < 0.05$  compared with WT + sham group; #,  $P < 0.05$  compared with WT + DOCA group.

#### *Deletion of IGF-1 gene led to enhanced RSNA and NE Spillover in DOCA-salt hypertensive mice*

RSNA was significantly enhanced in WT and LID mice after DOCA-salt treatment when compared to sham mice, but it was clearly higher in mice from LID + DOCA group than in those from WT+DOCA group (all  $P < 0.05$ , Fig. 3A). Besides, the plasma NE

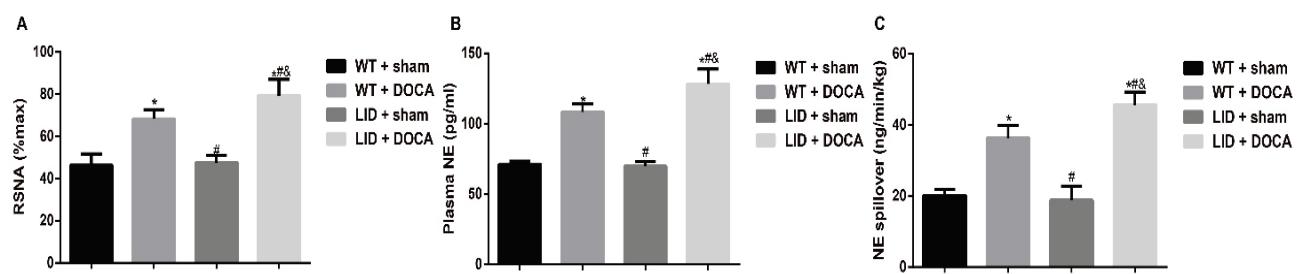
concentration and NE spillover were significantly increased in mice from the WT + DOCA group, as compared with sham mice (all  $P < 0.05$ ), and these elevations were even more obvious in mice from the LID + DOCA group, although no significant differences were found among sham mice (all  $P > 0.05$ , Fig. 3B-C).

#### *Deletion of IGF-1 gene aggravated left ventricular dysfunction in DOCA-salt hypertensive mice*

The DOCA mice presented the higher left ventricular function-related indexes (IVSd, IVSs, LVPWd, LVAWd, LVAWs, LVIDd and LVIDs) than sham mice (all  $P < 0.05$ , Table 1). These indexes were remarkably increased in mice from LID + DOCA group in comparison with those in WT + DOCA group (all  $P < 0.05$ ), whereas no differences were exhibited between sham groups (all  $P > 0.05$ ).

#### *Deletion of IGF-1 gene aggravated renal dysfunction in DOCA-salt hypertensive mice*

The DOCA mice had higher levels of urinary protein, urea nitrogen and creatinine, but lower creatinine clearance than the respective sham mice (all  $P < 0.05$ ). Furthermore, mice in the WT + DOCA group had significantly reduced urinary protein, urea nitrogen and creatinine, while they had elevated creatinine clearance as compared with mice in the LID + DOCA group (Fig. 4A-D). On the other hand, HE staining demonstrated that (Fig. 4E) the DOCA mice showed obvious renal histological changes (including tubular ectasia, glomerular fibrosis, interstitial cell infiltration, and increased arterial wall thickness) than sham mice. The renal dysfunction was more serious in mice from the LID + DOCA group.

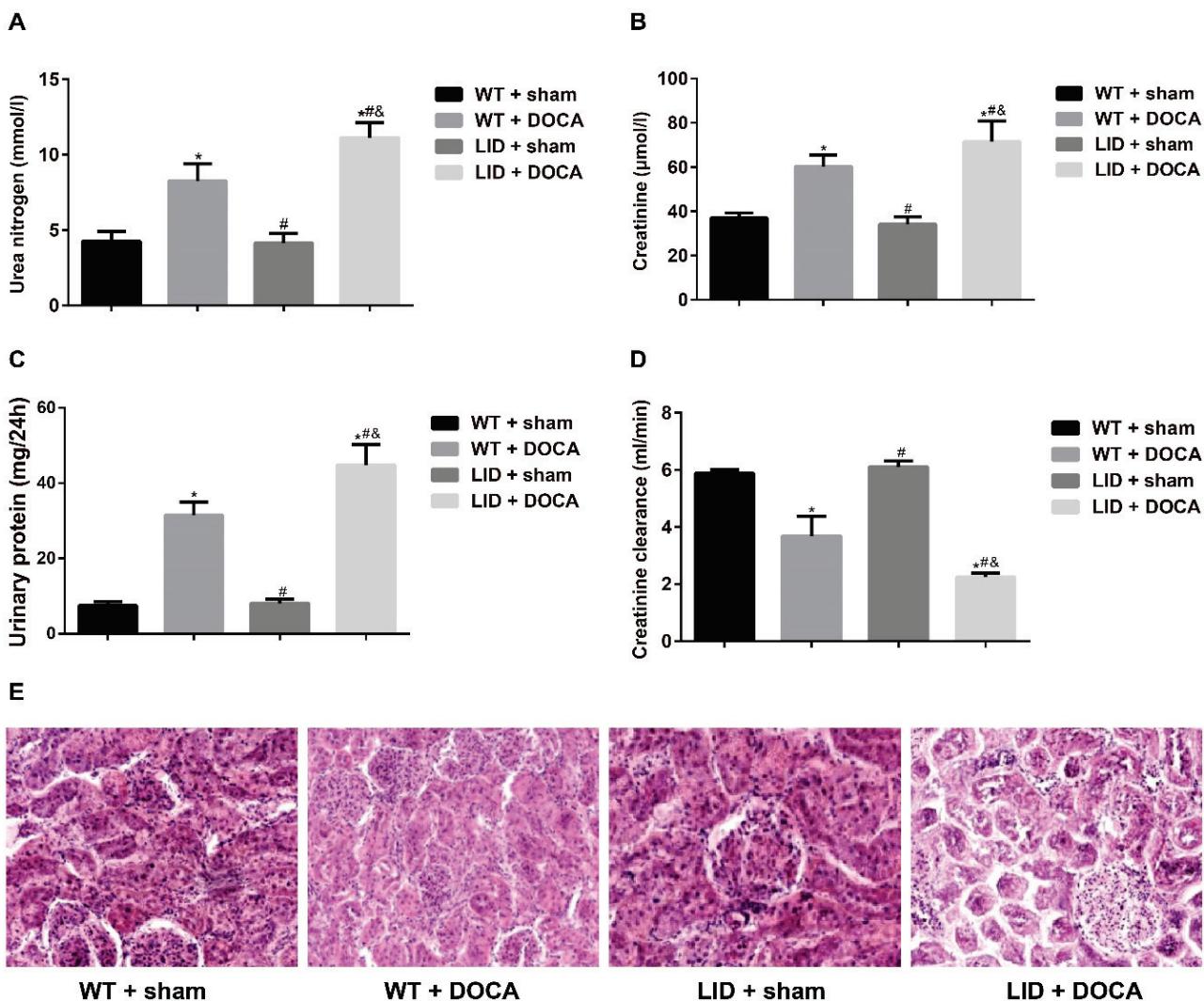


**Fig. 3.** Deletion of IGF-1 gene led to enhanced RSNA (A), plasma NE concentration (B) and NE Spillover (C) in DOCA-salt hypertensive mice. \*,  $P < 0.05$  compared with WT + sham group; #,  $P < 0.05$  compared with WT + DOCA group; \*\*,  $P < 0.05$  compared with LID + sham group.

**Table 1.** Deletion of IGF-1 gene significantly aggravated the left ventricular dysfunction in DOCA-salt hypertensive mice

	WT + sham	WT + DOCA	LID + sham	LID + DOCA
IVSd (mm)	0.85 ± 0.11	1.02 ± 0.05*	0.83 ± 0.05	1.29 ± 0.09**#
IVSs (mm)	1.31 ± 0.15	1.50 ± 0.04*	1.29 ± 0.11	1.70 ± 0.07**#
LVPWd (mm)	0.75 ± 0.06	0.88 ± 0.07*	0.74 ± 0.05	1.04 ± 0.03**#
LVPWs (mm)	1.11 ± 0.07	1.05 ± 0.13	1.04 ± 0.10	1.08 ± 0.05
LVAWd(mm)	0.78 ± 0.05	0.87 ± 0.03*	0.77 ± 0.05	0.96 ± 0.03**#
LVAWs(mm)	1.17 ± 0.05	1.31 ± 0.01*	1.19 ± 0.03	1.45 ± 0.09**#
LVIDd(mm)	3.47 ± 0.11	3.82 ± 0.15*	3.33 ± 0.17	4.38 ± 0.07**#
LVIDs(mm)	2.55 ± 0.10	2.95 ± 0.25*	2.53 ± 0.23	3.43 ± 0.12**#

IVSd, interventricular septal thickness in diastole; IVSs, interventricular septal thickness in systole; LVPWd, left ventricular posterior wall thickness in diastole; LVPWs, left ventricular posterior wall thickness in systole; LVAWd, left ventricular anterior wall thickness in diastole; LVAWs, left ventricular anterior wall thickness in systole; LVIDd, left ventricular internal diameter in diastole; LVIDs, left ventricular internal diameter in systole; \*, P<0.05 compared with sham groups; #, P<0.05 compared with WT + DOCA group.



**Fig. 4.** Deletion of IGF-1 gene on renal dysfunction in DOCA-salt hypertensive mice. (A-D) Comparison of the levels of urine protein, urea nitrogen, creatinine and creatinine clearance of mice among each group; \*, P<0.05 compared with WT + sham group; #, P<0.05 compared with WT + DOCA group; &, P<0.05 compared with LID + sham group; (E) The pathological changes of renal tissues in mice among each group evaluated by HE staining ( $\times 10$ , scale bar = 50 $\mu$ m)

## Discussion

Salt is the main cause of elevated blood pressure in DOCA-salt hypertensive mice, because in the absence of salt, the blood pressure of mice would not increase significantly even after removing the kidney or subcutaneously given DOCA (Krishnan *et al.* 2016). Thus, in the current study DOCA-salt hypertension was induced in mice subjected to subcutaneous injection of DOCA after uninephrectomy and receiving 1% NaCl solution to drink. As a consequence, the SBP in mice was significantly increased and renal damage (like proteinuria) was observed, which was consistent with the previous study (Jennings *et al.* 2012). Importantly, IGF-1 has been identified as a protective factor for cardiovascular system through a variety of ways, such as inhibiting the vascular endothelial cell injury, which was negatively linked to some cardiovascular risk factors (Yang *et al.* 2012), including hypertension (Vickers *et al.* 2001). For example, the cavernous IGF-1 bioavailability and levels were found to be dramatically reduced in patients with hypertension (Zhang *et al.* 2011, Zhou *et al.* 2016). A similar decreased serum level of IGF-1 was also observed in DOCA-salt hypertensive mice. A possible explanation is that there might be a mutation in the IGF1 gene in patients with hypertension, causing the down-regulation of circulating IGF1 levels (Yang *et al.* 2012). Moreover, we also confirmed that the mice in LID + DOCA groups and LID + sham groups had the lower serum IGF-1 levels than those in WT + DOCA and WT + sham groups, since the LID mice had a complete deficiency of IGF-1 knocked out by the Cre/loxP homologous recombination, leading to the significant decreased serum levels of IGF-1. Besides, the reduction of IGF1 levels caused by the hypertension was not stronger than the complete deficiency of IGF-1, therefore the mice in WT + DOCA group had higher serum levels of IGF-1 than those in LID + sham group. In addition, the LID mice and WT mice are comparable owing to the same LoxP structure and genetic stability (Ealey *et al.* 2008). Furthermore, hypertension is the clinical syndrome featured by increased arterial blood pressure, namely, the SBP above 140 mm Hg (Jones, 2017). However, SBP (measured in this study by the tail-cuff method) increased time-dependently in DOCA-salt hypertensive mice, SBP increase being higher in LID mice. This suggests that IGF-1 gene deletion enhanced blood pressure response to DOCA-salt treatment, possibly due to the reduced vascular bioavailability of

nitric oxide (NO), eventually promoting the progression of hypertension (Yang *et al.* 2012). However, no variation of the SBP was found between the LID + sham group and WT + sham group in our study. In addition, pulse pressure was not correlated with serum IGF-1 levels in healthy women as reported by JH Kim *et al.* (2006), indicating the *IGF-1* deletion may not affect SBP of the healthy mice or individuals.

Over-enhancement of sympathetic nerve activity (SNA) is accepted as an important characteristic of hypertension, thus the reduction of increased SNA constitutes one of the major targets for the prevention and treatment of hypertension (de Almeida Chaves Rodrigues *et al.* 2013). As we know, the renal sympathetic nerve has higher sympathetic nerve activity (SNA) than any other nerves, which could better reflect the cardiovascular SNA level and the central SNA level (Baum, 2018). Indeed, we observed the apparent increase of RSNA in mice from the DOCA groups (WT + DOCA group and LID + DOCA group) as compared with the sham groups (WT + sham group and LID + sham group), and the increase was higher in mice from LID + DOCA group than those from WT + DOCA group. Consistently, IGF-1 has been found to reduce the lumbar sympathetic nerve activity (LSNA) and RSNA (Duanmu *et al.* 1997), which was related to the activity of GH/IGF-1 axis (Sverrisdottir *et al.* 2010). On the other hand, the long-term stimulation has been reported to lead to the increase of NE spillover in peripheral blood, which faithfully reflects the state of RSNA (Noshiro *et al.* 1991). Thus, the plasma NE concentration and NE spillover detected in this experiment were clearly higher in LID mice with DOCA-salt treatment. However, the RSNA, plasma NE concentration and NE spillover in the LID + sham are lower than in the WT + DOCA, which indirectly indicated that IGF-1 gene deletion may cause the enhancement of RSNA and NE spillover in the progression of hypertension, but having no effects in sham mice. Recent study reported that an increase in SNA may be responsible for left ventricular dysfunction (Guizar-Mendoza *et al.* 2006). In accordance, the left-ventricular function related indicators, such as IVSd, IVSs, LVPWd, LVAWd, LVAWs, LVIDd, and LVIDs were elevated in DOCA-salt hypertensive LID mice. A previous study also discovered the reduction of serum IGF-1 levels also in patients with left ventricular dysfunction (Abe *et al.* 2006). Notably, Enoki *et al.* reported that mesenchymal stem cells loaded with IGF-1 can effectively improve the left ventricular function in

rats undergoing myocardial infarction (Enoki *et al.* 2010), indirectly indicating that IGF-1 gene deletion can aggravate the condition of left ventricular dysfunction in hypertensive patients or animals.

Finally, the kidney, as one of the major target organs in hypertensive disease, is responsible for removing waste products and excess water (Mule *et al.* 2017). The pathological renal change in hypertension mainly include tubular ectasia, glomerular fibrosis, interstitial cell infiltration, and increased arterial wall thickness (Wang *et al.* 2013). Similar results were disclosed by HE staining in the kidneys of our WT + DOCA mice. It is worth mentioning that the LID mice had more severe renal impairment than WT mice among the mice treated by DOCA-salt. It was demonstrated that IGF-1 was dramatically decreased in patients with acute renal failure (ARF) and reflux nephropathy (RN) (Kornhauser *et al.* 2002). There is a possibility that the poor expression of IGF-1 may promote the deposition of interstitial collagen and the apoptosis of renal tubular cells to deteriorate the renal function (Chertin *et al.* 2004). As previous studies demonstrated, urinary protein and creatinine levels of hypertensive patients were significantly higher than the healthy controls, which could diagnose renal damage in patients with

hypertension (Wang *et al.* 2015, Zhu *et al.* 2015). In our experiment, we also found that as compared with sham groups, the DOCA-salt hypertensive mice had the apparent elevation in urinary protein, urea nitrogen and creatinine, and the reduction of creatinine clearance, which changed more obvious in mice from LID + DOCA group. H Yasuda and his group demonstrated that injection of recombinant human IGF-1 (rhIGF-1) reduced serum creatinine and urea nitrogen levels in cisplatin (CDDP) group but not in CDDP-untreated rats (Yasuda *et al.* 2004). All mention above further suggested that IGF-1 deletion may cause a more severe renal damage in DOCA-salt hypertensive mice.

In summary, we found that deletion of IGF-1 gene may promote the increase in RSNA and up-regulate the levels of plasma NE concentration and NE spillover, eventually aggravating the left ventricular dysfunction and renal damage in DOCA-salt hypertensive mice.

### Conflict of Interest

There is no conflict of interest.

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