

## Visfatin reduces gap junction mediated cell-to-cell communication in proximal tubule-derived epithelial cells

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**Aim:** Associated with obesity, visfatin may have a role in the pathogenesis of diabetic nephropathy in type II diabetes. This study identifies a link between the visfatin, TGF- $\beta$ 1 and reduced cell-to-cell coupling in proximal tubule derived epithelial cells.

**Methods:** Western blot analysis confirmed changes in expression of connexins Cx26, Cx40 and Cx43 in HK2-cells +/- visfatin &/or TGF- $\beta$ 1. Visfatin evoked increases in TGF- $\beta$ 1 secretion were determined by ELISA, and functional intercellular communication assessed via Lucifer yellow dye transfer and paired-whole cell patch clamp electrophysiology.

**Results:** Visfatin (10, 100 and 200ng/mL) decreased expression of Cx26 to  $58\pm 11\%$ ,  $40\pm 3\%$  and  $21\pm 2\%$  and Cx43 to  $73\pm 5\%$ ,  $44\pm 4\%$  and  $29\pm 6\%$  as compared to control at 48hrs ( $n=3$ ;  $P<0.01$ ). The effects were not dependent on changes in membrane integrity, cytotoxicity or cell viability as assessed by MTT, crystal-violet and lactate-dehydrogenase assays. Expression of Cx40 was unaffected by the adipocytokine. Visfatin (200ng/ml) increased TGF- $\beta$ 1 secretion by 154% of control ( $n=3$ ;  $p<0.01$ ). Visfatin-evoked changes in Cx26 and Cx43 expression were mimicked by exogenous application of TGF- $\beta$ 1 (2-10ng/ml,  $p<0.001$   $n=3$ ). Visfatin reduced dye transfer between coupled-cells and decreased functional conductance by 63% as compared to control ( $n=6$ ;  $p<0.01$ ).

**Conclusions:** Visfatin reduced connexin expression in HK2-cells, an effect mirrored by a loss in functional conductance between coupled cells. Visfatin increased TGF- $\beta$  secretion and the pattern of change for connexin expression was mimicked by exogenous application of the pro-fibrotic cytokine. These data suggest that visfatin reduces connexin-mediated intercellular communication in proximal tubule-derived epithelial cells via a TGF- $\beta$  dependent pathway.

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