

# KSN 2016 Abstract Submission

*Clinical & Experimental Pathology & Cell Biology*

KSN2016ABS-1459

## Two isoforms of Matrix Metalloproteinase-2 in diabetic nephropathy

Sang Heon Song\*<sup>1</sup>, Min Young Lee<sup>1</sup>, Nari Shin<sup>2</sup>, Harin Rhee, Il Young Kim<sup>3</sup>, Eun Young Seong<sup>1</sup>, Dong Won Lee<sup>3</sup>, Soo Bong Lee<sup>3</sup>, Ihm Soo Kwak<sup>1</sup>

<sup>1</sup>Internal Medicine, Pusan National University Hospital, Busan, <sup>2</sup>pathology, <sup>3</sup>Internal Medicine, Yangsan Pusan National University Hospital, Yangsan, Korea, Republic Of

**Background:** Recent study uncovered the intracellular form called N-terminal truncated MMP-2 (NTT-MMP-2), which was generated by activation of an alternative intronic promoter unlikely full length MMP-2 (FL-MMP-2). The aim of this study is to explore the status of two isoforms of MMP-2 the expression in vitro, in vivo and human biopsy tissue, respectively and propose the new fundamental basis for diabetic nephropathy study related with MMP-2.

**Methods:** HK2 cells were cultured with different concentrations of D-glucose (5mM, 30mM) for 2, 24 and 48h, respectively and we also use 4-hydroxy-2-hexenal (HHE) as an another stimuli on HK2 cells. Two isoform of MMP-2 transcripts were measured using qPCR and the change of two isoforms of MMP-2 was checked after pyrrolidine dithiocarbamate (PDTC) treatment as an antioxidant and NF-kB inhibitor. Also, two isoforms of MMP-2 were tested in streptozotocin-induced diabetic mice at 12 weeks and 24 weeks, respectively. Finally, using human kidney biopsy tissue, the expressions of FL-MMP-2 and NTT-MMP-2 were analyzed by qPCR and immunohistochemical stain [controls (N=10) and diabetic kidney disease group (N=25)].

**Results:** Both FL-MMP-2 and NTT-MMP-2 transcripts significantly elevated by high glucose in HK2 cells. FL-MMP-2 was increased according to exposed time and has a peak expression by 15 times at 48h ( $15.4 \pm 0.7$ ,  $p < 0.001$ ). Unlike FL-MMP-2, NTT-MMP-2 was more earlier peaked at 24h and approximately four times greater than control ( $3.9 \pm 0.2$ ,  $p < 0.001$ ). By HHE stimulation, both FL-MMP-2 and NTT-MMP-2 transcripts also significantly upregulated in HK2 cell. PDTC inhibited NTT-MMP-2 transcription by high glucose, but FL-MMP-2 was not changed by PDTC. In STZ-induced diabetic mice model, FL-MMP-2 and NTT-MMP-2 transcripts were upregulated compared with control. Especially, NTT-MMP-2 expression was more strikingly increased compared with FL-MMP-2 in 24 weeks diabetic mice (7.2 fold change,  $p = 0.003$  vs. 1.8 fold change,  $p = 0.002$ ). NTT-MMP-2 was intensely increased in diabetic mice compared with control ( $p = 0.001$ ). NTT-MMP-2 was not stained at all in control mice. FL-MMP-2 and NTT-MMP-2 expression were found mainly within tubular epithelial cellular compartment, not glomeruli. Like animal model, two isoforms of MMP-2 were highly upregulated in human diabetic kidney disease. Specifically, FL-MMP-2 and NTT-MMP-2 transcripts were increased about by 12.5 and 3.1 times compared with control, respectively ( $p = 0.001$ ). Immunohistochemical staining grade was higher in diabetic kidney disease both in case of FL-MMP-2 and NTT-MMP-2, respectively (FL-MMP-2,  $1.8 \pm 0.9$  vs.  $2.5 \pm 0.5$ ,  $p = 0.03$ ; NTT-MMP-2,  $0.9 \pm 0.7$  vs.  $2.4 \pm 0.6$ ,  $p < 0.001$ )

**Conclusion:** Two isoforms of MMP-2 is related with diabetic injury and NTT-MMP-2 is entirely induced by diabetes unlike FL-MMP-2. Further validated study is needed to support the pathogenic role of two isoforms of MMP-2 in diabetic nephropathy

**Keywords:** chronic kidney disease, Diabetic Nephropathy, matrix metalloproteinase-2, mitochondria