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Abstract (English)

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| Title (in capitals) | THE ROLE OF IL-20 CYTOKINE SUBFAMILY IN THE PATHOGENESIS OF CHRONIC KIDNEY DISEASE |
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| Text | <p>Background: Regardless of the etiology kidney fibrosis is a common outcome of progressive chronic kidney diseases. Our recent study showed that levels of interleukin (IL)-20 subfamily members, including IL-19 and IL-24 significantly increased in kidneys underwent unilateral ureteral obstruction (UUO). However, their precise role in the pathomechanism of renal fibrosis is not clearly understood.</p> <p>Methods: To study the role of IL-20 cytokine subfamily we applied a mouse model of UUO induced kidney fibrosis on wild type and IL-20 receptor beta gene knockout (IL-20Rβ KO) mice. Masson's trichrome and Picro-Sirius Red staining were used to investigate the renal accumulation of extracellular matrix proteins. Real-time RT-PCR and western blot method were performed to measure the renal expression of fibrosis associated molecules. We also investigated the <i>in vitro</i> effect of IL-24 treatment on transforming growth factor beta (TGF-β) and platelet derived growth factor B (PDGF-B) expression of human proximal tubular epithelial (HK-2) cells by real-time RT-PCR and flow cytometry.</p> <p>Results: We found elevated level of IL-19, IL-24 and IL-20Rβ in the fibrotic kidneys. IL-20Rβ KO mice showed reduced extracellular matrix deposition and decreased α-smooth muscle actin expression compared to wild-type mice following UUO. Treatment of renal epithelial cells with IL-24 increased their TGF-β and PDGF-B production.</p> <p>Conclusion: Our study provides direct evidence of the pathogenic role of IL-20 cytokine subfamily in the development of renal fibrosis, possibly through the IL-24 mediated production of pro-fibrotic factors. Therefore, inhibition of IL-24 may have therapeutic effect in treatment of chronic kidney diseases.</p> <p>Grants: OTKA K116928 and VKE-2017-00006. This project was supported by the János Bolyai Research Scholarship of the Hungarian Academy of Sciences.</p> |
| Structure | |
| Background Methods Results Conclusion | |
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