



MEETING ABSTRACT

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The role of the lymphotoxin- β receptor (LT β R) in hepatocyte-mediated liver regeneration

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From 1st International Conference of Collaborative Research Center 974: Liver Damage and Regeneration Düsseldorf, Germany. 15-16 November 2013

Background

The lymphotoxin beta receptor (LT β R) is a prototypic member of the TNF/TNFR superfamily and has diverse functions in regulating immune responses against pathogens, in organogenesis and maintenance of structural integrity of secondary lymphoid tissues [1]. Mice deficient in LT β R (LT β R^{-/-}) show impaired resistance to intracellular pathogens and defects in peripheral lymphoid tissues [2,3]. Importantly, LT β R^{-/-} mice also exhibit markedly reduced survival after partial (70%) hepatectomy (PHx) [4], compared to WT controls. Liver mass is tightly regulated to 5% of the body weight and the liver retains a remarkable capacity for regeneration in response to acute injury. Loss of at least 30% of liver mass leads to synchronized proliferation of normally quiescent mature hepatocytes (compensatory hyperplasia) until physiologic liver mass is regained. Recently, it has been shown that lymphotoxins required for liver regeneration are produced by T cells immigrating into the liver during the regeneration process [5]. During this regeneration process, it is mandatory that the liver continues to perform its essential functions, such as protein synthesis, glycogen storage and bile secretion. There is evidence, that nuclear receptor-dependent bile acid signaling is required to initiate efficient liver regeneration [6] and that signaling through the TNFRp55, another core member of the TNF/TNFR superfamily is an essential part of early liver regeneration [7]. Interestingly, signaling pathways through TNFRp55 and LT β R share downstream signaling components. However, the molecular mechanisms orchestrated by LT β R and TNFRp55 signaling remain incompletely defined.

Results

Our data confirms that following PHx LT β R^{-/-} mice have a decreased survival rate compared to WT mice (62 % vs. 90 %, respectively). Surviving LT β R^{-/-} mice show a rate of liver regeneration comparable to WT animals. Post PHx H&E stained liver sections of LT β R^{-/-} animals showed higher numbers and larger areas of necrosis in the liver tissue, increased vacuolisation of hepatocytes, oedematous changes in the portal fields and dilated lymphatic vessels. ELISAs and clinical chemistry analysis show an altered cytokine expression profile in LT β R^{-/-} animals and several serum proteins appear to be deregulated, e.g. a significant increase in alkaline phosphatase in LT β R^{-/-} animals 24 and 48 h post PHx could be observed. On the other hand, ALT and AST levels do not differ between LT β R^{-/-} and WT animals. Total bile acid levels as well as the bile acid profile, as determined by UPLC-MS/MS, change markedly in both cohorts after PHx and differ between WT and LT β R^{-/-} animals, with the latter showing indications of cholestasis. This difference in bile acid metabolism is supported by quantitative RT/PCR data which shows variations in expression levels of various transporters and regulatory proteins involved in bile acid homeostasis between the two cohorts. Interestingly, expression of the LT β R gene on *in vitro* outgrown cholangiocytes of untreated WT animals could be demonstrated. Microarray analysis of WT and LT β R^{-/-} mice identified a panel of differentially expressed genes; most prominent among these was a markedly decreased expression of murinoglobulin-2, a proteinase inhibitor of the α 2-macroglobulin family, in LT β R^{-/-} mice.

Conclusions

Our findings demonstrate that a deficiency in LT β R signalling leads to substantial changes in liver morphology, gene expression, cytokine levels and other serum parameters after PHx. The various bile acids analyzed are

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differentially regulated after PHx in WT animals and this regulation is disturbed in $LT\beta R^{-/-}$ animals after PHx. This markedly different bile acid profile in $LT\beta R$ animals after PHx that seems to be associated with cholestasis indicates that bile acid homeostasis may play a key role in determining survival after 70 % PHx in rodents. It is not yet known how $LT\beta R$ signaling pertains to bile acid metabolism, and the detailed mechanisms of this interaction are in the focus of further studies. In conclusion, these results emphasize the importance of $LT\beta R$ signaling for efficient liver regeneration and may provide further insights into this complex process.

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Published: 19 June 2014

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doi:10.1186/2047-783X-19-S1-S3

Cite this article as: Behnke et al.: The role of the lymphotoxin- β receptor ($LT\beta R$) in hepatocyte-mediated liver regeneration. *European Journal of Medical Research* 2014 **19**(Suppl 1):S3.

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