BMC Pharmacology



Meeting abstract Open Access

Evaluation of Stat5 as a potential drug target in bcr/abl-induced leukemias

Andrea Hölbl¹, Christian Schuster¹, Boris Kovacic², Maria Hölzl¹, Sabine Fajmann¹, Florian Grebien³, Lothar Hennighausen⁴, Yongzhi Cui⁴, Richard Moriggl⁵, Hartmut Beug² and Veronika Sexl*¹

Address: ¹Centre of Biomolecular Medicine and Pharmacology, Medical University of Vienna, 1090 Vienna, Austria, ²Institute of Molecular Pathology (IMP), 1030 Vienna, Austria, ³Research Center for Molecular Medicine of the Austrian Academy of Sciences (CeMM), 1090 Vienna, Austria, ⁴NIDDK, National Institute of Health, Bethesda, MD 20892, USA and ⁵Ludwig-Boltzmann Institute for Cancer Research (LBI-CR), 1090 Vienna, Austria

Email: Veronika Sexl* - veronika.sexl@meduniwien.ac.at

* Corresponding author

from 14th Scientific Symposium of the Austrian Pharmacological Society (APHAR) Innsbruck, Austria. 21–22 November 2008

Published: 5 November 2008

BMC Pharmacology 2008, 8(Suppl 1):A4 doi:10.1186/1471-2210-8-S1-A4

This abstract is available from: http://www.biomedcentral.com/1471-2210/8/S1/A4 © 2008 Hölbl et al; licensee BioMed Central Ltd.

The Stat5 transcription factors Stat5a and Stat5b have been implicated in lymphoid development and transformation. Using the complete Stat5 knockout mice, we have previously shown that Stat5a/bnull/null cells were resistant to transformation and leukemia development induced by Abelson oncogenes, whereas Stat5a/b^{ΔN/ΔN} cells readily transformed. So far, these findings showed distinct susceptibility to Abelson-induced transformation of Stat5a/ $b^{\Delta N/\Delta N}$ and $Stat5a/b^{null/null}$ mice and defined Stat5 as key regulator of initial transformation. In this study, we tested whether Stat5a/b is also essential for the maintenance of a transformed state. Therefore we developed a system, where Stat5a/b could be deleted at will. Abelson-transformed B lymphoid cells were generated from Stat5a/bfl/fl gene targeted mice that had been crossed with Mx-Cre transgenic animals. These leukemic Stat5a/bfl/flMxCre cells were then used to test effects of Stat5a/b ablation in vitro and in vivo. In vitro, Stat5a/b deletion resulted in a cell cycle arrest followed by apoptosis. Nine days after deletion, no viable cells could be detected. In line with that, a down-regulation of Stat5 target genes mediating G1/S transition within the cell cycle and viability, such as cyclin D2 and cyclin D3, c-myc and bcl-x₁ was found. When leukemic Stat5a/bfl/flMxCre cells were injected into wild type or immuno-compromised mice leukemia rapidly

developed. Again, deletion of Stat5a/b in vivo within the leukemic cells significantly counteracted disease progression as indicated by an increase of leukemia latency from 16 to 49 days. Eventually, all animals succumbed to a Stat5a/b-positive leukemia indicating that a few residual cells escaped deletion. Moreover, p53 abruption or overexpression of the oncogene did not alter the susceptibility to Stat5 loss of established leukemic cell lines. Taken together our data define a key role for Stat5a/b not only for lymphoid development but also for lymphoid transformation. Stat5a/b is necessary for the initial transformation as well as for leukemia progression. This absolute necessity for the proliferation and viability of Abelsontransformed cells puts Stat5a/b into the spotlight of new therapeutic strategies for the treatment of bcr/abl-induced leukemias.