

Regulatory T cells in endogenous mouse lymphoma are antigen-specific nTregs and provide a target for immune checkpoint-inhibiting therapies

Vera Bauer^a, Fatima Ahmetlic^a, Tanja Riedel^a, Nico Trautwein^b, Tim Sparwasser^c, Stefan Stevanovic^b, Martin Röcken^d, Ralph Mocikat^{a,*}

^a Helmholtz-Zentrum München

^b Eberhard-Karls-Universität, Interfakultäres Institut für Zellbiologie, Tübingen, Germany ^c Institut für Infektionsimmunologie, Twincore, Zentrum für Experimentelle und Klinische Infektionsforschung, Hannover, Germany ^d Eberhard-Karls-Universität, Universitäts-Hautklinik, Tübingen, Germany

* Corresponding author, email: Mocikat@helmholtz-muenchen.de

© 2018 Ralph Mocikat; licensee Infinite Science Publishing

This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/4.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Keywords: Regulatory T cell, B-cell lymphoma, tumor escape, checkpoint inhibition

Purpose/Objectives: Foxp3+ regulatory T cells (Tregs) play an important role in maintaining immune homeostasis. In malignant disease, however, Tregs contribute to the generation of an immunosuppressive microenvironment. To establish therapeutic approaches it is necessary to uncover activating and suppressive mechanisms. Immune checkpoint-blocking (ICB) monoclonal antibodies (mAbs) have been shown to elicit cancer regression and long-lasting tumor control. Particular interest was focused to the mutual influence of Treg cells and therapy by ICB.

Materials/Methods: We used a c-MYC transgenic mouse model of spontaneously arising Bcell lymphoma, which mirrors key features of human Burkitt lymphoma. To assess treatment effects mice were intraperitoneally injected with mAbs against PD-1 and CTLA-4. Mice splenocytes and lymphocytes were analyzed by flow cytometry.

Results: An augmented fraction of CD4+Foxp3+Tregs was detected in spleens and lymph nodes of tumor-bearing c-MYC mice. Tregs were involved in suppressing antitumor response because specific ablation of Tregs significantly delayed tumor development. Compared to wildtype mice, c-MYC Treg cells showed an activated phenotype as evidenced by upregulated CD69 and CD137. The high expression of the costimulatory molecule CD137 indicated TCR-specific activation. As intratumoral Tregs were predominantly Nrp-1+Helios+nTregs, we suggested that Tregs recognized tumor-associated self-antigens. We identified MHC class II-restricted self-epitopes, which were prevalent in lymphoma as compared to normal B cells and could be recognized by Treg cells resulting in enhanced proliferation (Ki-67) in vitro. Interestingly, effector T cells (Teffs) were able to recognize the same epitopes. Antigen contact apparently led to upregulation of IL-10. In vitro suppression assays showed that Tregs from c-MYC and wt mice potently suppress proliferation of Teffs in a dose-dependent manner. When preventing cell contacts, Teff proliferation was increased and almost restored upon additional IL-10 neutralization. Tregs from c-MYC mice treated with anti-CTLA-4/anti-PD-1 mAbs revealed a lower suppressive capacity.

Conclusion: In malignant lymphoma, nTregs directed against self epitopes are involved in cancer immune escape. Taken together, the Treg population is a promising target in future immunotherapies and needs to be further investigated.