

Using CRISPR to improve T cell cancer therapies

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Introduction: Cytokines of the TGF-beta superfamily (TGFb-sf) are responsible for the immunosuppressive microenvironment of many cancer entities. Overcoming this barrier constitutes a major goal in the therapy with adoptively transferred immune cells to target cancer, especially solid tumors. Thus, strategies to counteract signaling induced by factors of the TGF-beta superfamily could help to improve adoptive cell therapy. TGF-beta subfamily (TGFb-f) signaling is mediated by phosphorylation of R-Smads (Smad2, Smad3) and formation of complexes with the Co-Smad (Smad4). BMP-subfamily (BMP-f) signaling is mediated by R-Smads 1, 5 and 8 together with Smad4. Thus, Smad4 acts as common integrator of TGFb-sf cytokines, while inhibitory factors such as Smad6 and Smad7 act as physiological antagonists. In order to target all cytokines of the TGFb-sf by only one genetic modification, we aimed to overexpress inhibitory Smads in primary human CD4⁺ and CD8⁺ T-cells and assess the effect of this manipulation on T-cell function and polarization. In parallel, we abrogated TGF-beta superfamily effects by CRISPR/Cas9 mediated knockout of Smad4.



Conclusion: The role of TGFb and cytokines of the TGFb-sf on the immune response against tumors is well documented and intensively investigated. Adoptive cell therapies, such as CAR T-Cells, represent successful therapeutics in lymphoma treatment. Several factors however limit the success of these therapies in solid tumors, among them a TGFb-sf driven tumor microenvironment. Thus, we aim to adoptive immunotherapy by intrinsically rendering T-cells resistant against all cytokines of the TGFb-sf. While targeted Smad7 overexpression and SMAD4 KO reverts TGFb mediated effects on proliferation and T cell differentiation, these approaches did not affect TGFb induced upredulation of

