

PhD project No. 23, Prof. Jeker

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| Scientific Area | Immune-related diseases |
| Two project titles | A) Understanding microRNA biology to improve T cell therapy B) Exploiting microRNAs for CAR T design |
| Host country | Switzerland |
| Supervisor, institution | Prof. Lukas Jeker, University of Basel, Switzerland |
| Co-Supervisor, institution | A) Prof. T. Cathomen, University of Freiburg, Germany B) Prof. T. Cathomen, University of Freiburg, Germany |
| Mentor, institution | A) and B) to be determined later |
| Secondment institution | A) and B) University of Basel, Switzerland and University of Freiburg, Germany |
| Short description of the supervisor's lab with introduction to the topic | |
| <p>The Jekerlab has studied microRNAs in cell types that affect immune regulation. Our current focus is on microRNAs that regulate T cell differentiation and function. We have demonstrated that the miR-17-92 cluster is important for multiple T cell types. More recently we have shown that miR-17-92 is a key regulator of CD4+ T cell activation. We have identified genes that are directly and indirectly regulated by the miR-17-92 cluster. We aim to deepen our mechanistic understanding and leverage this knowledge to improve T cell therapies. This goes in line with our research program on genome engineered cellular therapies.</p> | |
| Topic description, including techniques to be used | |
| <p>Project A) We plan to characterize the function of the identified miR-17-92 target genes. To this end we use genome engineering in murine T cells and we generate mouse models with new genetic modifications. We will study how alteration (deletion/modification) of miR-17-92 target genes affects T cell biology. We will study the genome engineered T cells in vitro and in mouse models in vivo. <u>Techniques:</u> Molecular biology, CRISPR/Cas9, gene targeting, mouse models, flow cytometry, transcriptome analysis, cell culture</p> <p>Project B) We plan to use genome engineering of microRNA target genes to improve T cell therapies including chimeric antigen receptor (CAR) T cells. In addition, we plan to exploit microRNAs for T cell therapies. We will generate mouse models and will validate our findings in human T cells. <u>Techniques:</u> Molecular biology, CRISPR/Cas9, gene targeting, mouse models, flow cytometry, transcriptome analysis, cell culture, mouse and primary human T cells</p> | |
| Recommended applicant's training (technical expertise and knowledge) | |
| <p><u>Techniques:</u> molecular biology, cell culture, flow cytometry <u>Knowledge:</u> cell biology, T cell biology, gene regulation including posttranscriptional gene regulation</p> | |
| Maximum two relevant publications | |
| <p>Kornete et al., 2018, Journal of Immunology: Highly Efficient and Versatile Plasmid-based Gene Editing in Primary T Cells Doelz et al., 2020, bioRxiv.org : The non-coding RNA miR-17-92 is a central mediator of T cell activation</p> | |

Ethics description

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| 1. Humans | |
| This research involves human participants. | YES <input type="checkbox"/> / NO <input checked="" type="checkbox"/> |
| This research involves physical interventions on the study participants. | YES <input type="checkbox"/> / NO <input checked="" type="checkbox"/> |
| 2. Human Cells /Tissues | |
| This research involves human cells or tissues, such as blood. | YES <input checked="" type="checkbox"/> / NO <input type="checkbox"/> |
| 3. Personal Data | |
| This research involves personal data collection and/or processing. | YES <input type="checkbox"/> / NO <input checked="" type="checkbox"/> |
| This research involves further processing of previously collected personal data (secondary use). | YES <input type="checkbox"/> / NO <input checked="" type="checkbox"/> |
| 4. Animals | |
| This research involves animals, such as mice. | YES <input checked="" type="checkbox"/> / NO <input type="checkbox"/> |