Genetic Control of T-Cell Proliferation with Synthetic RNA Regulatory Systems

Thesis by

Yvonne Yu-Hsuan Chen

In Partial Fulfillment of the Requirements

for the Degree of

Doctor of Philosophy

California Institute of Technology

Pasadena, California

2011

(Defended December 10, 2010)

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Acknowledgments

The following pages tell a tale that has been five years in the making. As much as I would like to claim this work for the countless hours spent pipetting alone in tissue culture rooms, singing to Frank Sinatra and Ella Fitzgerald on Pandora Radio, it is in fact the fruit of many individuals' labors. I only hope that the final product has reciprocated some of the efforts given to it by so many.

First, I would like to thank my tireless thesis advisor, Prof. Christina Smolke, for her unfailing support and infinite patience with my research and myself. Even with her abundant talents, Christina is a model of diligence and commitment. I thank her for the thoughtful care she puts into steering this laboratory and guiding its members. Even amidst my most draining experiments, I always had the comfort of knowing that I did not walk alone on this journey.

I must also thank Dr. Michael Jensen for his boundless energy and excitement about my research, his generosity in providing me with resources and advice, and his interest in my professional development. I could not have completed this work without the help of many friends at City of Hope. I thank Araceli Hamlett and Bessie Meechoovet for initiating me into the art of T-cell culturing, Brenda Aguilar for laboring many hours in the animal facility on my behalf, and Cherrilyn Bautista for being my reliable source of expert knowledge on T cells and beyond. I also thank Dr. Christine Brown, Wen-Chung Chang, Michelle Hunter, Renate Starr, Jamie Wagner, and Winnie Wong for being always ready teach me a technique or lend me a hand. I also must acknowledge Christine Wright for managing the most organized laboratory I have ever seen.

The Smolke Laboratory has been a wonderful home for the past five-and-a-half years. I would like to thank Dr. Chase Beisel and Dr. Stephanie Culler for showing me the ropes in tissue culture and for being so generous with their time and knowledge whenever I had questions or needed advice. I thank Dr. Maun Nyan Win for sharing his expertise on his beloved creation, the ribozyme switch platform. I will always remember Katie Galloway for our stimulating discussions on not-so-scientific subjects, Arwen Brown for sustaining a passion for the humanities in a sea of scientists and engineers, and Leo D'Espaux for his amazing culinary skills and unfettered pursuit of life, even if it leads to a broken bone or two.

Graduate school would not have been nearly as eventful or enjoyable without the many friends I met at Caltech. I thank Arthur Chan, Puneet Chhabra, Claire Farnsworth, and Andrew Metcalf for leading me out of the lab and onto hiking trails. Mike Rubel, Karen Lau, and my cousin Solomon kept me in a semblance of fitness by humoring me on the badminton court. The many hours spent first on transport and thermo problem sets and later on the equally intellectually rigorous game that is Mexican Train with Diana Smirnova, Havala O. T. Pye, and Mary Louie will remain some of my fondest memories of Caltech. Mary, my roommate of three years, kept me sane by being the patient ears for my happy and melancholy ramblings alike. I thank her for showing me a generosity of spirit that I am not sure I knew was possible.

Finally, I would like to thank my most constant supporters—my family. I thank my parents and grandparents for showing me the virtue of perseverance, and for providing a secure and nurturing environment in which to grow. 感謝阿嬤教我謹慎;感謝阿公教我好學;感謝外婆教我努力奮鬥;感謝外公教我在工作之餘不忘生活的

浪漫。爸爸媽媽給我的一切是篇幅再長也述不完的。在這裡只能說聲「實在真多謝。」And last but certainly not least, I would like to thank my sister, Alice, whose path I seemed destined to follow, at least until now. If the road I traveled has been smoother or more colorful, it is because you have blazed the trail and shown me the way.

Abstract

Adoptive T-cell therapy, or the use of autologous T cells to seek and destroy diseased cells, is a promising treatment option for opportunistic diseases, virus-associated malignancies, and cancers. However, the safety and efficacy of adoptive T-cell therapy depend, in part, on the ability to sustain and tightly regulate the proliferation of transferred T cells *in vivo*. The emerging field of synthetic biology provides powerful conceptual and technological tools for the construction of regulatory systems that can interface with and reprogram complex biological processes such as cell growth. Here, we present the development of RNA-based regulatory systems that can control T-cell proliferation in a ligand-dependent manner, and examine the construction of integrated control systems capable of fine-tuned programming of cellular behavior.

We systematically investigate the translation of ribozyme-based regulatory devices from yeast to mammalian cells and identify design parameters critical to the portability of regulatory devices across host organisms. We report the construction of ligand-responsive ribozyme switch systems capable of modulating the transgenic expression of growth-stimulatory cytokines in mammalian lymphocytes. We demonstrate the ability of ribozyme switch systems to regulate T-cell proliferation in primary human central memory T cells and in animal models. We further develop ligand-responsive, miRNA-based devices to regulate the endogenous expression of cytokine receptor chains and the functional output of cytokine signaling pathways, highlighting the ability to construct integrated T-cell proliferation control systems employing various regulatory mechanisms to modulate multiple components in relevant signaling pathways. Finally, we describe efforts in the generation of novel RNA aptamers to clinically suitable

molecules, which can serve as the molecular inputs for ligand-responsive, RNA-based control systems in therapeutic applications.

The regulatory systems developed in this work are designed to be modular and transportable across host organisms and application contexts, thus providing a template for future designs in RNA-based genetic regulation. This work demonstrates the capability of RNA-based regulatory systems to advance next-generation treatment options for critical diseases, and highlights the potential of synthetic biological systems to achieve novel and practical functions in diverse applications.

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