

Design of Protein-DNA Dimerizers

Thesis by

Ryan Leonard Stafford

In Partial Fulfillment of the Requirements
for the Degree of
Doctor of Philosophy

California Institute of Technology

Pasadena, California

2008

(Defended August 8, 2007)

© 2008

Ryan Leonard Stafford

All Rights Reserved

for my family

Acknowledgements

It has been a long time since I started working at Caltech, so there are a lot of people to thank. First, I would like to thank my Mom and Dad for giving all of their kids, including me, the freedom to do what we want and support us no matter what. And thanks to my family-in-law, Annie, the Mom and Dad, and of course, my love, Kim. You have given me something else to do so I do not end up working every moment- only every other moment.

At a place like Caltech, it has been no surprise that I have been able to work with such smart people (although it may be surprising how fun they all are). In particular, Hans-Dieter (bear or beer) made a fantastic collaborator and mentor during my first couple years. I enjoyed the opportunity to work closely with Nick Nickols (risk:fun) and Mike Brochu (Palms Thai) my last couple years. Sometime in the middle I mentored Rachel Wang (Jonas) who always found a way to work hard when it mattered most and learned how to keep a notebook. I want to thank Carey Hsu (my old roomie), who showed me how to hustle on a basketball court and all the other core Freeballers: Justin Cohen (the 330 oasis), Dan Harki (outside jumper), Michelle Farkas (take care of my desk), John Phillips (the new Ray Doss), Anand Vedehra (pillow fight), Ray Doss (punched in the back), Mike Kesden (elbows), Tim Best (homestar runner), Adam (Tucker the turtle), and Katherine Poulin-Kerstien (lacrosse)-you guys are awesome. I have to especially thank Adam for guiding me to a good orthopedic surgeon. And I want to thank everyone else who should have played more ball: Jim Sanchez (fellow anteater), Claire Jacobs (Puckett's shoe), Christian Dose (something inappropriate), Jim Puckett (fried food), and Mike Marques (no team sports). And I want to thank all the other

Dervanites of the past and present, that I have not already mentioned, that have made my lab experience more enjoyable including Dave Chenoweth (racquetball), Steve Bartusiak (YouTube), Sherry Tsai (*really?*), Mareike Goeritz (cappuccino machine), Julie Poposki (flowers on campus), Anne Viger (*coffee?*), Ben Edeslon (acrobat slim shady), Shane Foister (Kentucky basketball), Eric Fechter (flying), Alex Heckel (German lessons), and Victor Rucker (footprinting stories). I want to wish the best of luck to Katy Muzikar (purple hair). I hope you, John, and Michelle have fun training all the new Dervan recruits.

And thanks to Peter for providing me with all the resources and real estate to do science. Thanks also to my committee, Bob Grubbs, who has been a pleasure to T.A. for; Doug Rees for being an inspiration and helping to support my future endeavors; and Steve Mayo. And I want to thank Greg Weiss for always supporting me since my days at UCI.

Abstract

Genes are regulated by proteins called transcription factors that bind to DNA in a sequence-specific manner and modulate the rate of transcription. Mutated transcription factors often lead to abnormal gene expression, developmental defects, and disease. This thesis describes the design of chemicals called protein-DNA dimerizers that mimic natural transcription factor protein-DNA complexes. In the long-term, it is hoped that these dimerizers will be able to engage or even replace mutant transcription factors and artificially regulate gene expression in living cells. Specifically, programmable DNA-binding pyrrole-imidazole polyamides conjugated to YPWM peptide motifs incorporating various linker domains facilitate the binding of a natural transcription factor, extradenticle, to DNA. From a design point of view, it has been explored what the minimum size and shape (branched or linear) is that will ultimately be optimal for cell uptake with adequate functional potency in the transcriptional apparatus. Branched dimerizers are shown to function with a minimal WM dipeptide protein-binding domain *in vitro* up to 37 °C, and linear dimerizers are shown to function with WMK tripeptides up to 20 °C. Collectively, branched and linear dimerizers can facilitate protein binding to DNA from 2 base pair overlap sites to ones that reach 6 base pairs apart. Polyamide-WM-fluorescein conjugates are also found to be cell permeable in several cell lines including HeLa, MCF-7, and PC3. These studies provide insight into the importance of linker length and composition, binding-site spacing and orientation, and the protein-binding domain content that are important for the optimization of protein-DNA dimerizers suitable for biological experiments.

Table of Contents

	Page
Acknowledgements.....	iv
Abstract.....	vi
List of Figures and Tables.....	viii
Chapter 1 Introduction to Protein-DNA Dimerizers.....	1
Chapter 2 Minimization of a Protein DNA Dimerizer.....	21
Chapter 3 Defining the Reach of Linear Protein-DNA Dimerizers.....	49
Chapter 4 Cell-Permeable Protein-DNA Dimerizers.....	77
Chapter 5 Toward a Synthetic HIF-1 α Mimic.....	97
Appendix A Protein-DNA Dimerizers Targeted to a Natural HOX Site.....	120
Appendix B Toward a Small Molecule YPWM Mimic.....	131
Appendix C Exd Expression Protocols.....	148
Appendix D Solid-Phase Synthesis of Polyamides Using Marshall-Liener Resin.....	155
Appendix E Dimerizer-Exd-DNA Crystallization Trials.....	161
Appendix F Supplemental Figures.....	161

List of Figures and Tables

Chapter 1			Page
Figure 1.1	A schematic of a gene promoter.....	3	
Figure 1.2	The structure of DNA.....	4	
Figure 1.3	Crystal structures of four transcription factor-DNA complexes.....	5	
Figure 1.4	An atomic model of the IFN- β enhanceosome.....	6	
Figure 1.5	Crystal structures of HOX/TALE/DNA ternary complexes.....	7	
Figure 1.6	Evolutionary conservation of the YPWM motif.....	9	
Figure 1.7	Structures illustrating DNA recognition by netropsin and distamycin.....	10	
Figure 1.8	The chemical structure and schematic of a typical pyrrole-imidazole hairpin polyamide.....	11	
Figure 1.9	A structure illustrating the DNA recognition by a synthetic polyamide...	12	
Figure 1.10	The chemical structure and model of a protein-DNA dimerizer.....	14	
Chapter 2			
Figure 2.1	Schematic of a protein-DNA dimerizer.....	23	
Figure 2.2	MPE footprinting of a protein-DNA dimerizer and Exd.....	24	
Figure 2.3	Dimerizer and control chemical structures.....	27	
Figure 2.4	Gel shift assays at 4 °C.....	28	
Table 2.1	Exd-DNA-dimerizer stabilities at 4 °C.....	29	
Table 2.2	Exd-DNA-dimerizer stabilities at 20 and 37 °C.....	30	
Figure 2.5	Gel shift assays at 20 and 37 °C.....	31	

Figure 2.6	DNase I footprinting of a minimal protein-DNA dimerizer.....	31
Table 2.3	Conjugate-DNA equilibrium association constants.....	32
Figure 2.7	An illustrative model of minimal protein-DNA dimerizers.....	35

Chapter 3

Figure 3.1	Design of linear protein-DNA dimerizers.....	52
Figure 3.2	Structures of linear conjugates.....	54
Figure 3.3	Proximal gel shift assays.....	56
Table 3.1	Summary of proximal gel shift assays.....	57
Figure 3.4	Distal gel shift assays.....	58
Table 3.2	Summary of distal gel shift assays.....	59
Figure 3.5	Gel shift assays with WMK conjugates.....	59
Figure 3.6	Estimation of linker distances.....	61
Figure 3.7	Models of linear protein-DNA dimerizers.....	62

Chapter 4

Figure 4.1	Summary of previous cell uptake results.....	73
Figure 4.2	Summary of more previous cell uptake results.....	75
Figure 4.3	Synthesis of polyamide-peptide-FAM conjugates.....	76
Figure 4.4	Uptake results with polyamide-dipeptide-FAM conjugates.....	77
Figure 4.5	Uptake results with polyamide-tetrapeptide-FAM conjugates.....	79
Figure 4.6	Synthesis of peptide-FAM conjugates.....	80
Figure 4.7	Uptake results with peptide-FAM conjugates.....	80

Figure 4.8	Structures of IPA conjugates.....	81
Figure 4.9	Gel shift experiments with cell-permeable protein-DNA dimerizers.....	81

Chapter 5

Figure 5.1	Small molecule binding sites on CBP.....	93
Figure 5.2	Design of a small molecule HIF-1 α mimic.....	93
Figure 5.3	Putative small molecule activation domains (ADs).....	94
Figure 5.4	Synthesis of AD1.....	95
Figure 5.5	Synthesis of AD3.....	95
Figure 5.6	Biotin-AD1 conjugate pull-down experiments.....	96
Figure 5.7	NMR structure of CBP with AD3.....	96
Table 5.1	Frequency of polyamide match sites in the VEGF promoter.....	97
Figure 5.8	Synthesis of polyamide-AD-FITC conjugates.....	98
Figure 5.9	Cell uptake results for polyamide-AD-FITC conjugates.....	99
Figure 5.10	Quantitative DNase I footprinting of polyamide-AD conjugates.....	100
Figure 5.11	Synthesis of branched polyamide-AD conjugates.....	101
Figure 5.12	Synthesis of linear polyamide-AD conjugates.....	101
Figure 5.13	RT-PCR measurements of VEGF induction.....	103

Appendix A

Figure A.1	Structures of polyamides targeted to natural and artificial sites.....	115
Figure A.2	DNase I footprinting of labial compounds.....	116
Figure A.3	Site-selective recruitment of Exd.....	117

Figure A.4	Cell uptake of a polyamide with β -C3-FAM C-terminus.....	119
Figure A.5	Synthesis of a labial polyamide-peptide-FAM conjugate.....	119
Figure A.6	Cell uptake results for a labial polyamide-peptide-FAM conjugate.....	120
Table A.1	DNA equilibrium association constants for labial compounds.....	120

Appendix B

Figure B.1	Design of a YPWM β -turn mimetic.....	126
Figure B.2	Synthesis of a polyamide- β -turn mimetic conjugate.....	127
Figure B.3	Gel shift experiment with a polyamide- β -turn mimetic conjugate.....	128
Figure B.4	Design of WM peptoid conjugates.....	129
Figure B.5	Synthesis of WM peptoids.....	130
Figure B.6	Gel shift experiment with a polyamide-WM peptoid conjugate.....	130
Figure B.7	Structure of a small molecule that binds Pbx.....	131
Figure B.8	Synthesis of a small molecule that binds Pbx.....	132
Figure B.9	Gel shift experiment with a polyamide-small molecule conjugate.....	133

Appendix C

Figure C.1	Amino acid sequence of Exd isoform C.....	143
Figure C.2	Purification and analysis of first Exd preparation.....	143
Figure C.3	MALDI-TOF MS analysis of Exd.....	144
Figure C.4	UV-Vis of Exd.....	146
Figure C.5	Purification and analysis of the second Exd preparation.....	146
Figure C.6	Additional purification of the second Exd preparation.....	147

Appendix D

Figure D.1	Scheme for solid-phase synthesis on Marshall-Liener resin.....	151
------------	--	-----

Appendix E

Figure E.1	Summary of HOX/PBC/DNA crystal structures.....	156
Figure E.2	Design of a complex for crystallization.....	157
Figure E.3	Pictures of select crystals.....	157

Appendix F

Figure F.1	Proximal gel shift assays with controls.....	159
Figure F.2	Distal gel shift assays with controls.....	160
Figure F.3	Gel shift assays with WM conjugates.....	161
Figure F.4	Gel shift assays with more WMK conjugates.....	162
Figure F.5	The dependence of complex stability on site spacing.....	163
Figure F.6	Proximal DNA duplexes.....	164
Figure F.7	Distal DNA duplexes.....	165