

Impact of Fluorinated Amino Acids on Artificial Protein Block Copolymers of Two Self-assembling Domains

Carlo Yuvienco¹, Jennifer S. Haghpanah¹, Richard Hwang and Jin Kim Montclare^{1,2}

Chemical & Biological Sciences, Polytechnic Institute of NYU, Brooklyn, NY, 11201¹
Department of Biochemistry, SUNY Downstate Medical Center, Brooklyn, NY, 11203²

1 Abstract

The requirement for smart protein-derived biomaterials to change in macromolecular structure in response to external stimuli necessitates the design of controllable modes of self-assembly. The recent advances in unnatural amino acid incorporation enables the integration of chemical diversity into such proteins, further expanding the level of control and materials properties. In particular, fluorinated amino acids have been of the biomaterials. Specifically, we have incorporated para-fluorophenylalanine and trifluoroleucine into three block polymers that consist of a β -spiral elastin-mimetic protein (E) and an α -helical coiled-coil region of cartilage-oligomeric matrix protein (C). These proteins, synthesized as the block sequences – EC, CE, and ECE – are chosen for their distinct structures, functions, and modes of self-assembly. We demonstrate successful incorporation of the non-natural amino acids as well as characterization emphasizing their structural and functional distinction relative to the non-fluorinated constructs.

4 Protein Block Polymers

MRGSH₆GSKPIASA–Elastin–LEGSELA(AT)₆AACG–**COMPcc**–LQA(AT)₆AVDLQPS
MRGSH₆GSACELA(AT)₆AACG–**COMPcc**–LQA(AT)₆AVDKPIASA–Elastin–LESGSGTGAKL
MRGSH₆GSKPIASA–Elastin–LEGSELA(AT)₆AACG–**COMPcc**–LQA(AT)₆AVDKPIASA–Elastin–LESGSGTGAKL

Elastin = [(VPGVG)₂VPGFG(VPGVG)₃]VP
COMPcc = DLAPQMLREQETNAALQDVRELLRQVKEITFLKNTVMESDASG

7 Expression

• Small-scale (5 mL) and large-scale (100 mL) expressions of
• Phe auxotrophic AF1Q cells
• Leu auxotrophic LAM1000 cells
• Residual natural amino acids removed prior to induction with IPTG by washing with 2 cycles of washing with 0.9% NaCl
• All lanes in gels below normalize to OD₆₀₀ = 1.00

10 Comparison of Melting Curves of Wild-type and Fluorinated Constructs

EC pFF and wt T_m = 33C \rightarrow 34C
CE pFF and wt T_m = 44C \rightarrow 45C
ECE pFF and wt T_m = 47C \rightarrow 33C

• Negligible effect on melting temperature for EC and CE diblocks
• Notable shift in T_m for ECE triblock
• Enhanced cooperativity of ECE with incorporated fluorinated Phe
• Though melt temperatures are the same as wild-type, the CE shows an increase in transition cooperativity versus the EC diblock upon incorporation

2 Cartilage Oligomeric Matrix Protein Coiled-Coil (C)

Hydrophobic pore 73 Å long 2-6 Å wide
Binding small molecules
Self-assembles into a pentameric bouquet-like structure α -Helical secondary structure 10.9 kDa

V. N. Malashkevich, R. A. Kammerer, V. P. Elmov, T. Schallies, and J. Engel, *Science*, (1996) 274, 761-765.
Sue Ozkay, Jürgen Engel, Jürg Stiefeld, *EMBO J* 2002, 21, 5993-8.

5 Non-canonical Amino Acids

• Incorporated non-canonical amino acids have been shown to enhance structural stability some proteins
• Photo-crosslinking possible with the incorporation of p-azidoPhe

• We hypothesize that the incorporation of fluorinated amino acids into our fusion sequences will result in an alteration of temperature-sensitive behavior
• Separate incorporation of p-fluoroPhe and trifluoroLeu

Tang, Y. et al. Fluorinated Coiled-Coil Proteins Prepared In Vivo Display Enhanced Thermal and Chemical Stability. *Angew. Chem. Int. Ed. Engl.* 40, 1484-1487 (2001).
[1] James Link, David A Tirrell. *Methods* 2005, 36, 291-298.

8 Purification of pFF Constructs

• Purification of whole cell lysate under denatured conditions (6M urea)
• Co²⁺ affinity chromatography using a 5mL HiTrap IMAC FF column (GE)
• Gels showing elution fractions from purification
• MALDI-TOF analysis confirming non-canonical amino acid incorporation

EC pFF
CE pFF
ECE pFF

EC wt
768 Da
750 Da
772 Da

CE pFF
768 Da
750 Da
772 Da

ECE pFF
768 Da
750 Da
772 Da

3 Elastin (E)

(Val/Ile)-Pro-Xaa-Yaa-Gly

Lower critical solution temperature (LCST) – from liquid to solid by adding heat

• Temperature-sensitive aggregation and self-assembly
• Coupling to other functional domains has been shown to affect both aggregation and temperature-sensitivity
• Focus of potential use lies in drug (small molecule) delivery and release

Idealized β -spiral (VPGVG)₆
10° C, 6 ns MD
42° C, 6 ns MD

B. U. V. Duggett, *J Muscle Res Cell Motil* 2002, 23, 561-73.
Bumjoo Kim, Ashutosh Chikoti, *J Am Chem Soc* 2008, 130, 17867-73.

6 Residue-specific incorporation of non-natural amino acids

Introduction of non-canonical amino acid upon induction
Aminoacyl tRNA synthetase
tRNA^{aa}
mRNA
ECFAM

• Random-like structure at low T
• Shift to α -helical

CE • different structure vs. EC at 10
• More α -helical at 4C

ECE

9 CD of pFF Constructs

Temperature Scale: °C
65
55
45
35
25
15
10
4

EC
CE
ECE

EC pFF
CE pFF
ECE pFF

• Different wavelength scan
• Initial structure at 4C

ACKNOWLEDGEMENTS

WE WOULD LIKE TO THANK POLYTECHNIC UNIVERSITY START-UP FUNDS, THE OTHMER INTITUTE, THE WECHLER AWARD, AIR FORCE OFFICE OF SCIENTIFIC RESEARCH, SOCIETY OF PLASTIC ENGINEERS, ACS CHEMISTRY INSTITUTE, ACS ENVIRONMENTAL CHEMISTRY DIVISION, UNILEVER, THE NATIONAL SCIENCE FOUNDATION GK-12 FELLOWS GRANT DGE-0741714