

## **PROFESSIONAL BIOGRAPHY: Fred J. Stevens**

Bioscience Division  
BIO-202 A-141  
Argonne National Laboratory  
6700 S Cass Ave  
Argonne, IL 60439

e-mail: <fstevens@anl.gov>  
phone: 630-252-3837  
fax: 630-252-3777

## **EDUCATIONAL BACKGROUND:**

|      |      |            |                         |
|------|------|------------|-------------------------|
| 1976 | Ph.D | Biophysics | Northwestern University |
| 1974 | M.S. | Physics    | Northwestern University |
| 1971 | B.A. | Physics    | Hamline University      |

## **PROFESSIONAL EXPERIENCE:**

- 1/01-present: Senior Biophysicist; Biosciences Division  
Argonne National Laboratory
- 8/87-12/00: Biophysicist; Biosciences Division,  
Argonne National Laboratory.
- 1/84-7/87: Assistant Biophysicist; Division of Biological and  
Medical Research (BIM), Argonne National Laboratory.
- 8/81-12/83: Project Manager, Research Biochemist; Diagnostics  
Division, Abbott Laboratories.
- 2/81-7/81: Assistant Scientist, BIM, Argonne National Laboratory  
(Term Appointment).
- 11/77-1/81: Postdoctoral Appointee, BIM,  
Argonne National Laboratory.
- 1/76-10/77: Research Associate, Department of Microbiology and  
Public Health, Michigan State University.

## **CURRENT RESEARCH INTERESTS:**

Analyses of protein primary structures to identify correlates of pathological properties with emphasis on the immunoglobulin superfamily and related beta-domain proteins. Protein:protein interactions; protein structure and function; physical principles in biological systems; computer modeling for enzyme modification and computer simulation of macromolecular interactions. Development of chromatography systems and computer simulations for the analysis of the kinetics and affinity of macromolecular interaction including antibody:antigen, anti-idiotope:idiotope, rheumatoid factor:IgG, antibody:peptide and protein:nucleic acid interactions. Development of chromatographic methods for the identification and characterization of pathological immunoglobulin light chains formed during multiple myeloma and related diseases. Recognition of protein structural and functional similarities at minimal levels of amino acid sequence identity.

## **HONORS:**

- Editorial Board: *Amyloid: The International Journal of Experimental and Clinical Investigation*  
(1999-present)
- NIH Postdoctoral Fellowship. 1978. Awarded for project initiated at  
Michigan State University. Declined.
- Fellowships from Physics Dept., Northwestern University (1971-1973)
- National Merit Scholarship (1966-1971)

## **CONSULTANTSHIPS AND ADJUNCT APPOINTMENTS:**

- 4/93--present: Affiliate, University of Chicago Cancer Institute
- 2/90--present: University of Tennessee Medical Center at Knoxville.  
Visiting Scientist
- 4/85--4/88: Abbott Laboratories. Consultant, Chemical and  
Agricultural Products Division.
- 1/82--12/83: Consultant, BIM, Argonne National Laboratory.
- 12/78--12/80: Guest Research Associate. Division of Biology,  
Brookhaven National Laboratory.

## **TEACHING EXPERIENCE:**

Argonne National Laboratory maintains several formal programs that support education and training of students at the undergraduate, graduate, and post-graduate levels. Undergraduate students participate in research projects in my laboratory on an annual basis. Training includes formal and informal lectures that cover fundamentals of the methods that are used, as well as presentation of the larger context and significance of the studies in which they participate. A graduate student recently completed her PhD (IIT) under my supervision. I have participated on the graduate committee of two graduate students who have obtained their PhD. I have supervised or participated in the supervision and development of 10 postdoctoral students. One of these has been recently hired onto the scientific staff of Argonne National Laboratory.

## **COMPETITIVE EXTRAMURAL FUNDING:**

Source: NIH

Project: "Biophysics of Myeloma Pathology" (05/01/92 - 03/31/06)

Total: \$3,117,489 (direct + indirect); (+ supplemental \$217,812)

Source: DOE (ER/LTT)

Project: "Exploratory Research to Identify  
Anti-Amyloid Drugs" (CRADA) (9/01/93 - 12/30/96)

Total: \$510,000 (direct + indirect)

Source: DOE (NN-20)

Project : "Ultrasensitive Actinide Detection Through  
Bio-based Receptors" (12/1/96 -09/30/00)

Total: \$1,500,000 (est, direct + indirect)

Source: NIH

Project: "Protein Structure, Stability, and Aging" (04/15/00 – 04/14/04)

Total: \$1,555,000 (direct + indirect); (+ supplemental \$83,952)

## **MEMBERSHIPS IN PROFESSIONAL SOCIETIES:**

Biophysical Society (active)

Protein Society (active)

American Association of Immunologists (active)

American Association for the Advancement of Science (active)

International Society of Amyloidosis (active)

International Interest Group in Biorecognition Technology

American Society for Cell Biology

American Society for Microbiology

## **PATENTS:**

1. Stevens, F.J. 1988. Size-exclusion chromatography system for macromolecular interaction analysis. **No. 4,762,617**
2. Stevens, F.J. 1991. A method of field flow fractionation wherein the polarity of the electric field is periodically reversed **No. 5,133,844**
3. Stevens, F.J. and S.-P. Tsai. Preparation of acrylic acid from lactic acid via modified fumarase. Abandoned.
4. Stevens, F.J., A. Solomon, and E. Myatt. 2000. Method for detecting and diagnosing disease caused by pathological protein aggregation. **No. 6,063,636**
5. Stevens, F.J. Method for producing fabrication material for constructing micrometer-scaled machines, fabrication material for micrometer-scaled machines.

Pending.

6. Stevens, F.J., R. Raffen, P. Wilkins-Stevens, and M. Schiffer. 2002. Method for altering antibody light chain interactions. **No. 6,485,943**
7. Stevens, F.J. and P. Wilkins-Stevens. Device for detecting actinides, method for detecting actinides. Pending
8. Stevens, F.J., M. Schiffer, P. Wilkins Stevens, W. Carey Hanly, and S.L. Tollaksen. Device for detecting molecules, method for detecting molecules. Pending.
9. Stevens, F.J., Y. Argon, D. Davis, and R. Raffen. A fibril-blocking peptide, a method for preventing fibril formation. **No. 6,878,521**
10. Stevens, F.J. 2003. System and method for a parallel immunoassay system. 2003 **No. 6,489,120**

#### INVENTION REPORTS:

1. Stevens, F.J. 1986. Size-exclusion chromatography system for macromolecular interaction analysis. ANL-IN-86-018, DOE: S-64,667.
2. Stevens, F.J. 1987. One-step viral immunoassay. ANL-IN-87-107, DOE: S-67,394.
3. Stevens, F.J. and D.A. LeBuis. 1988. Pulsed-field field-flow fractionation (PF4). ANL-IN-88-072, DOE: S68,647.
4. Stevens, F.J. 1990. Nephromimetic chromatography for identification and diagnosis of pathological proteins. ANL-IN-90-011, DOE: S-71,196.
5. Stevens, F.J. 1990. Antibody characterization workstation. ANL-IN-017, DOE: S-71,319.
6. Stevens, F.J. 1990. Nonlinear electroimmuno detector device. ANL-IN-038, DOE: S-71,992.
7. Stevens, F.J. 1991. Micrometer design and construction by self-assembly of nanometer-scale engineered proteins. ANL-IN-91-101, DOE: S-75,872.
8. Stevens, F.J. 1992. Automated analytical assay involving magnetic separation. ANL-IN-92-045, DOE: S-76,933.
9. Stevens, F.J. and S.-P. Tsai. 1992. A process for conversion of lactic acid into acrylic acid catalyzed by a modified fumarase. ANL-IN-92-060.
10. Stevens, F.J. and M. Schiffer. 1993. Protein engineering strategies for optimized production of recombinant antibodies. ANL-IN-93-108
11. Stevens, F.J. 1994. Engineering of proteins to create detectors for actinides and other heavy metals. ANL-IN-94-144, DOE: S-83,527.

12. Stevens, F.J. 1994. Drug design strategies to inhibit amyloid fibril formation. ANL-IN-94-145, DOE S-83,528.
13. Stevens, F.J. 1995 Method to evoke a cellular immune response to *in situ* amyloid deposits. ANL-IN-95-019, DOE S84,206
14. Stevens, F.J. 1995. A novel biomaterial substance derived from a biological fibril. ANL-IN-95-040, DOE S-84,207
15. Stevens, F.J. 1995. PROSPECDS (Protein SequencePoly-Evaluation, Comparison, & Display Systems) ANL-SF-95-068
16. Stevens, FJ. 1996. 30 Second Immunoassay. ANL-IN-96-062, DOE S-86,391
17. Stevens, F.J. 1996. Use of genetically engineered plants to produce human biomolecules. ANL-IN-96-075, DOE S-86,844
18. Stevens, F.J. 1996. Immunassay for detection of biological agents. . ANL-IN-96-122, DOE-S87,875
19. Stevens, F.J. 1997. Massively Parallel Immunoassay (MPIA). ANL-IN-97-055, DOE S89,003
20. Stevens, F.J., M. Schiffer, K. Nash, and M. Jensen. 1997. Designs for actinide binding sites at the interface of a protein dimer. ANL-IN-97-056, DOE S89,002
21. Stevens, F.J. and M. Schiffer. 1998. Janusbody: a compact molecule with antibody-antigen binding characteristics. ANL-IN-98-021, DOE S90,947.
22. Stevens, F.J., and R. Raffen. 1998. In vitro protein fibril production: potential high-throughput method to screen for anti-amyloid drug leads. ANL-IN-98-046, DOE S91,201
23. Stevens, F.J., C.S. Giometti, and A. Joachimiak. 1998. The anti-western blot.
24. Stevens, F.J. and P. Wilkins Stevens. 1998. Immunological assays for actinides. ANL-IN-98-086,  
DOE S91,835
25. Stevens, F.J., Y. Argon, D. Davis, and R. Raffen. 1999. Composition of a fibril-blocking peptide; strategy for development of peptides to inhibit conformational disease resulting from protein domain swapping. ANL-IN-99-019, DOE S93,203
26. Argon, Y., D.P. Davis, F.J. Stevens, and R. Raffen. 2000. A method for inhibiting amyloid-like protein aggregation with peptides. UCHI # 820.
27. Stevens, F.J. 2000. A method for high-throughput screening of protein nucleic acid (PNA) libraries for ligand (protein-binding) activity. ANL-IN-00-13
28. Stevens, F.J. and F. Collart. 2000. Universal Genomic Detector: A generic method for

identification and comparison of genomic material. ANL-IN-00-014

29. Stevens, F.J. 2000. Cyber cell: Molecular beacon methodology for a live-cell/microprocessor interface. ANL-IN-00-15
30. Stevens, F.J. and J.F. Carpenter. 2000. Lethal ligands: From conformational disease to conformational therapy. ANL-IN-00-22
31. Stevens, F.J. 2000. Biosentinel: Device for autonomous monitoring and evaluation of environmental samples. ANL-IN-00-52
32. Stevens, F.J. and M. Schiffer. 2000. Process for customizing functional stability of antibodies.
33. Stevens, F.J. 2000. Protein nucleic acids: methodology for generation and detection of forensic signatures.
34. Stevens, F.J. 2001. Differential PCR PNA-clamped SNP diagnostics. ANL-IN-01-101.
35. Stevens, F.J. 2001. Peptidyl chaperones. ANL-IN-01-101.
36. Stevens, F.J. 2003. Method for producing antibodies to proteins from select against without access to select agents; method for producing antibodies to proteins that cannot be produced for experimental reasons. ANL-IN-03-015
37. Vilim, R., and F.J. Stevens. 2003. Parsimonious fold-specific protein classifier from attribute-constrained genetic algorithms. ANL-IN-03-003.
38. Stevens, F.J. 2003. A thermophilic organophosphate detoxification enzyme. ANL-IN-03-051
39. Stevens, F.J., L. Chen, B. Kay. 2003. Functionalized three-dimensional protein lattices formed by self assembly. ANL-IN-03-52
40. Stevens, F.J. 2003. A systematic strategy for the disruption of adhesion protein-based biofilms of medical and commercial significance. ANL-IN-03-115

#### **TECHNICAL REPORTS:**

1. Levine, D., M. Facello, P. Hallstrom, G. Reeder, B. Walenz, and F.J. Stevens. 1996. STALK Users Guide. ANL/MCS-TM-214
2. Levine, D., M. Facello, P. Hallstrom, G. Reeder, B. Walenz, and F.J. Stevens. 1996. STALK Programmers Guide. ANL/MCS-TM-215
3. Stevens, F.J. 1999. Protein structure, stability, and conformational disease: human antibody light chains 1999. ANL/BIO/99-1

#### **ABSTRACTS/PRESENTATIONS:**

1. Stevens, F.J. and T.T. Wu. 1974. A third-stage mutant of *E. coli* capable of utilizing

- he polyhydric pentitols as sole source of carbon and energy. Abstr. Ann. Mtg. Am. Soc. Microbiol., p 156.
2. Stevens, F.J. and T.T. Wu. 1976. Acquisition by *E. coli* K-12 of the ability to utilize an unnatural pentose, D-lyxose. Federation Proc. **35**: 1660.
  3. Stevens, F.J. and R.L. Uffen. 1977. Acquisition of photosynthetic competence by anaerobic dark-grown cells of *R. rubrum* exposed to light. Presented at 4th Ann. Mtg. on the Molec. Biol. of Photosyn. Microorganisms; May 7.
  4. Stevens, F.J., H.S. Pankratz, and R.L. Uffen. 1977. Cytochemical and spectrophotometric examination of the effect of glutaraldehyde on the oxidation of 3,3'-diaminobenzidine in phototropic bacteria. Cell Biology **75**: 244a.
  5. Stevens, F.J., F. Westholm, A. Solomon, and M. Schiffer. 1979. Heterogeneity of κI light chains. Federation Proc. **38**: 940.
  6. Stevens, F.J., F.A. Westholm, A. Solomon, and M. Schiffer. 1979. Role of the third hypervariable region in the self-association of κI immunoglobulin light chains. Presented at Midwest Autumn Immunology Conference; November 5.
  7. Stevens, F.J., F.A. Westholm, A. Solomon, and M. Schiffer. 1980. Computer simulation of small zone gel filtration: heterologous association of immunoglobulin light chains. Federation Proc. **39**: 2210.
  8. Schiffer, M., F.J. Stevens, F. Westholm, D. Carlson, and B. Schoenborn. 1980. Small angle neutron scattering study of Bence Jones protein Mcg: comparison of structures in solution and in crystal. Presented at Gordon Conference on Diffraction Methods in Molecular Biology; June 21.
  9. Peraino, C., and F.J. Stevens. 1980. Characteristics of liver tumor promotion by phenobarbital. Presented at Symposium on Cocarcinogenesis and Biological Effects of Tumor Promoters; Oct. 13 (Krais, West Germany).
  10. Schiffer, M., F.J. Stevens, and F.A. Westholm. 1980. Structural properties of Bence Jones proteins in the crystal and in solution. 38th Annual Pittsburgh Diffraction Conference; Oct. 30.
  11. Wu, T.T., P.W. Stevens, J.G. Hovis, and F.J. Stevens. 1981. Production of an evolutionary remnant enzyme in *Escherichia coli* K-12. FEMS Symposium on Overproduction of Microbiol Products. Abstracts, p. 43.
  12. Short, M., F.J. Stevens, F. Westholm, A. Solomon, and M. Schiffer. 1980. Preliminary X-ray crystallographic studies of Bence Jones protein Loc. Presented at Midwest Autumn Immunology Conference; Oct. 18.
  13. Short, M., F.J. Stevens, F. Westholm, A. Solomon, and M. Schiffer. 1982. Crystallographic investigation of lambda Bence Jones protein Loc. Federation Proc. **41**: 287.
  14. Short, M.T., F.J. Stevens, and M. Schiffer. 1983. High pressure gel permeation chromatography

- in the study of self-associating immunoglobulin κI light chains and an idiotype-anti-idiotype complex. *Federation Proc.* **42**: 931.
15. Stevens, F.J., S. Anaokar, G. Korom, J. Tice, and P. Spillar. 1983. Solid phase enzyme immunoassay for serum leutinizing hormone. *Clin. Chem.* **29**: 1241.
  16. Schiffer, M., R.L. McMasters, C.-H. Chang, and F.J. Stevens. 1984. Interactions of immunoglobulin constant domains. Gordon Research Conference on Physics and Physical Chemistry of Biopolymers; June 27.
  17. Stevens, F.J. and C.F. Ainsworth. 1985. Antibody-antigen interactions: application of chromatography simulation and laboratory microcomputer. *Federation Proc.* **44**: 1077.
  18. Stevens, F.J. 1985. Application of microcomputer-controlled size-exclusion chromatography and computer simulation to analysis of macromolecular interactions. Presented at Amoco-sponsored University/Industry Poster Session; Oct. 23 (Naperville, IL).
  19. Schiffer, M., C.-H. Chang, F.J. Stevens, and A. Solomon. 1985. Implications of the protruding binding site of light-chain dimer Loc for the structure of the anti-idiotypic antibody. Presented at Symposium Idiotype Networks and Immune Regulation: Potential Uses in Vaccines and In Understanding Human Diseases; December 4 (San Antonio).
  20. Stevens, F.J. and M. Schiffer. 1985. Analysis of idiotype:anti-idiotype interactions by small-zone size-exclusion chromatography: consideration of equilibrium constant and rate constants. Presented at Symposium Idiotype Networks and Immune Regulation: Potential Uses in Vaccines and In Understanding Human Diseases; December 4 (San Antonio).
  21. Stevens, F.J. and C.F. Ainsworth. 1986. Microcomputer controlled size-exclusion chromatography. *Biophys. J.* **49**: 88a.
  22. Carperos, W., J. Jwo, and F.J. Stevens. 1986. Analysis of monoclonal antibodies by liquid chromatography. Presented at Abbott Laboratories, State of Illinois-Sponsored Poster Session; November 26 (North Chicago, IL).
  23. Boernke, W.E. and F.J. Stevens. 1987. HPLC-based size-exclusion chromatography for the detection of interactions between nucleic acids and proteins. *Biophys. J.* **51**: 442a.
  24. Stevens, F.J. and M. Schiffer. 1987. Modification of an ELISA-based method to determine affinity: correction for nonlinear relationship between occupancy of antibody binding sites and attachment of IgG to immobilized antigen. *Federation Proc.* **46**: 491.
  25. Stevens, F.J. 1987. Simulation of macromolecular interactions: kinetically controlled elution characteristics during size-exclusion high-performance liquid chromatography. Presented at Amoco-sponsored University-Industry Poster Session; Oct 8 (Naperville, IL).
  26. Carperos, W.E. and F.J. Stevens. 1987. Application of size-exclusion high performance liquid chromatography for the measurement of antibody-antigen dissociation rate constants. Presented at Amoco-sponsored University-Industry Poster Session; Oct 8 (Naperville, IL).

27. Stevens, F.J. 1987. Macromolecular interactions: kinetically controlled elution characteristics during size-exclusion HPLC. Presented at Conference on Biotechnology Research Directions: Biomolecules; Nov 6 (Argonne, IL)
28. Carperos, W.E. and F.J. Stevens. 1987. Application of size-exclusion high-performance liquid chromatography for the measurement of antibody-antigen dissociation rate constants. Presented at Conference on Biotechnology Research Directions: Biomolecules; Nov 6 (Argonne, IL)
29. Schiffer, M., C.-H. Chang and F.J. Stevens. 1987. Domain-domain interactions affect the expression of idiotope. Presented at Symposium on Idiotype Networks and Immune Regulation: II Autoimmunity, Cancer, Receptors and Vaccines; Nov 18 (San Antonio).
30. Stevens, F.J. 1988. Simulation of kinetically controlled elution characteristics exhibited by macromolecular interactions during size-exclusion HPLC. *Biophys. J.* **53**: 400a.
31. Stevens, F.J., and P. Kaumaya. 1988. Size-exclusion HPLC analyses of protein and peptide epitopes. Presented at Second Conference on Biotechnology Research Directions: Biomolecules; Nov 4 (Argonne, IL)
31. Schiffer, M., C.-H. Chang, Z.B. Xu and F.J. Stevens. 1989. Modification of antibody binding site through altered V-V domain interactions. *J. Cell. Biol.* **13A**:94.
33. Boernke, W.E. and F.J. Stevens. 1989. Study of interactions between myeloperoxidase and nucleic acids by size-exclusion HPLC. *Biophys. J.* **55**: 525a.
34. Stevens, F.J., and P. Kaumaya. 1989. Size-exclusion HPLC analyses of peptide epitopes. *FASEB J.* **3**: A1135.
35. Schiffer, M., C.-H. Chang, Z-B. Xu and F.J. Stevens. 1989. Conformational effects and diversity of antibodies. Presented at 7th International Congress of Immunology, (Berlin).
36. Spangler, B.D., F.J. Stevens, and E.M. Westbrook. 1989. Binding of monoclonal antibodies to cholera toxin: comparison of solid phase and solution phase techniques. Presented at Amoco-sponsored University-Industry Poster Session; Oct 12 (Naperville, IL).
37. Stevens, F.J. 1990. Simulation of macromolecular interactions during size-exclusion chromatography: positive cooperativity contribution by gel matrix. *Biophys. J.* **57**: 82a.
38. Stevens, F.J. 1990. Chromatographic analyses of protein:protein interactions: from the supercomputer to the kidney. Presented at the University of Illinois at Chicago, Dept. of Biochemistry; April 19.
39. Stevens, F.J. 1990. Chromatographic analyses of macromolecular interactions: from the supercomputer to the kidney. Presented at the Wayne State University, Dept. of Biochemistry; Sept 4.
40. Stevens, F.J. 1990. Biophysical aspects of myeloma pathology. University of Tennessee-Knoxville, School of Medicine; Sept 7.

41. Myatt, E.A. and F.J. Stevens. 1991. Induction of protein aggregation in nephromimetic solutions. *Biophys. J.* **59**: 119a.
42. Shen, B., F.J. Stevens, and U. Luthi. 1991. Elasticity of human erythrocyte spectrin inferred from conformational analysis. *FASEB J.* **5**: A685.
43. Myatt, E.A. and F.J. Stevens. 1991. Induction of protein aggregation in nephromimetic solutions. Presented at Gordon Research Conference; June 17--21.
44. Spangler, B.D., G.D. Armstrong, and F.J. Stevens. 1991. HPLC analysis of pertussis binding to glycoproteins. Presented at fifteh European Workshop on Bacterial Protein Toxins, Veldhoven, July.
45. Chen, S., S. Goldin, F.J. Stevens, and B.W. Shen. 1991. Elasticity of human erythrocyte spectrin inferred from electron microscopic and conformational analysis. Presented at Gordon Research Conference, Plymouth State College, New London, N.H.; August 5--9.
46. Stevens, F.J. 1991. Polymerization of immunoglobulin domains: a model system for the development of facilitated macromolecular assembly. (Presented at the Second Foresight Conference on Molecular Nanotechnology, Palo Alto, CA; (November 7--9)).
47. Stevens, F.J. 1991. Biophysical and computational aspects of immunoglobulin light chain pathology. Chicago Medical School, North Chicago, IL. Dept of Biochemistry; November 21.
48. Stevens, F.J., E.A. Myatt, A. Solomon, and M. Schiffer. 1992. Amyloidosis AL: Molecular model for a biophysical disease. *Biophys. J.* **61**: A211
49. Stevens, F.J. and R.L. Stevens. 1992. Protein-protein interaction kinetics. Presented at Concurrent Supercomputing Consortium Workshop, CalTech, February 11.
50. Harrison, H.H., T. Godsey, K. Bedford, A. Katta, K. Miller, E. Weisenberg, J.E. Bowman, E. Myatt, C.S. Giometti, and F.J. Stevens. 1992. Comparison of microheterogeneity patterns of purified monoclonal light chains (Bence Jones proteins) and polyclonal free light chains that produce the pseudo-oligoclonal ("ladder light chain") pattern in immunofixation studies of urine. *Clin. Chem.*
51. Stevens, F.J. 1992. Analysis of antibody light chain interactions: modeling of amyloid fibril formation. Presented at Northern Illinois University, De Kalb, IL, Oct. 2.
52. Gaasterland, T., R.L. Stevens, F.J. Stevens, and M. Schiffer. Critiquing protein molecule structures. Presented at The Third Keck Symposium on Computational Biology, Houston, TX, Nov. 1-3, 1992.
52. Myatt, E.A., A. Solomon, and F.J. Stevens. Aggregation of immunoglobulin light chains. *J. Immunol.* **150**: 861, 1993.
54. Li, D., F.J. Stevens, M. Schiffer, and L.E. Anderson. Cysteine clusters in the light activated

chloroplast enzymes. *Plant Physiol.* **102:** 37, 1993.

55. Li, D., F.J. Stevens, M. Schiffer, and L.E. Anderson. Identification of redox-sensitive cysteines distal to catalytic site in light-activated chloroplastic glyceraldehyde-3-phosphate dehydrogenase by modeling. Presented at The XIII Midwest Enzyme Conference, Loyola University, Oct 9, 1993.
56. Huang, D.-B., C.-H. Chang, C. Ainsworth, M. Schiffer, F.J. Stevens, A. Brunger, M. Eulitz, and A. Solomon. Novel immunoglobulin variable domain interaction is observed. ACA Meeting, Albuquerque, PK14, 1993.
57. Jiang, X., P. Lykos, M. Schiffer, and F.J. Stevens. The design and evaluation of the peptide inhibition of antibody light chain dimer formation by molecular modeling. Presented at The Second Annual Development of Small Molecule Mimetic Drugs, Philadelphia, April 11-12, 1994.
58. Stevens, F.J. A molecular model for amyloid fibril formation by antibody light chains. Presented at Loyola University, Chicago, April 21, 1994.
59. Li, D., F.J. Stevens, M. Schiffer, and L.E. Anderson. Mechanisms of light modulation: identification of potential redox-sensitive cysteines in chloroplast fructose-1,6-bisphosphatase and sedoheptulose-1,7-bisphosphatase. *FASEB J.* **8:** A1290, 1994.
60. Muslin, E.H., D. Li, M. Schiffer, F.J. Stevens, M.I. Donnelly, and L. Anderson. Construction of a redox-sensitive malate dehydrogenase. Presented at XIV Annual Midwest Autumn Enzymology Conference. 1994
61. Stevens, F.J. Physical chemical aspects of antibody interactions with itself, antigen, and solid surfaces. Presented at Surfaces in Biomaterials '94, Scottsdale, AZ., Sept 8--10, 1994.
62. Li, D., E. H. Muslin, L.E. Anderson, M.M. Pacold, M. Donnelly, M. Schiffer, and F.J. Stevens. Engineering a domain-locking disulfide into a bacterial malate dehydrogenase produces a redox-sensitive enzyme. *FASEB J.* **9:**A1284, 1995.
63. Paul, S., L. Li, P. Wilkins-Stevens, F.J. Stevens, and A.Solomon. Natural catalytic antibodies: peptide hydrolyzing activities of monoclonal human light chains and VL fragment. *FASEB J* **9:** 1265, 1995.
64. Kalaga, R., H. Huang, F.J. Stevens, A. Solomon, and S. Paul. gp120 hydrolysis by catalytic antibody light chain. Presented at The 9th International Congress of Immunology, San Fransisco, CA, July 23-29, 1995.
65. Jiang, X.-L., E.A. Myatt, R. Raffen, and F.J. Stevens. Study of interactions between amyloid forming protein and glycosaminoglycans. Presented at Protein Interactions. Pfizer/Beckman Institute Protein Symposium. Urbana, IL, June 1-4, 1995.
66. Anderson, L.E., H.C. Huppe, D. Li, and F.J. Stevens. Identification of a potential redox -sensitive inter-domain disulfide in the sedoheptulose bisphosphatase of *Chlamydomonas*

- reinhardtii*. Presented at Xth International Photosynthesis Congress, Montpellier, France, August 20-25, 1995.
67. Stevens, F.J. Molecular studies of light chain amyloidosis. Presented at FASEB Conference: Amyloid and other Abnormal Protein Assembly Processes, Copper Mountain, CO, August 20-25, 1995.
68. Mulsin, E.H., F.J. Stevens, and L.E. Anderson. A disulfide bridge stabilizes anaerobically induced lactate dehydrogenase in barley. Presented at Thioredoxins and Related Proteins, Witzenhausen, University of Kassel, August 27-31, 1995.
69. Anderson, L.E., D. Li, E.H. Muslin, M.E. Pacold, M.I. Donnelly, M. Schiffer, and F.J. Stevens. Mechanism of light modulation: Identification of redox-sensitive cysteines in malate dehydrogenase by modeling and by site-directed mutagenesis. Presented at Thioredoxins and Related Proteins, Witzenhausen, University of Kassel, August 27-31, 1995.
70. Muslin, E.H., F.J. Stevens, and L.E. Anderson. Anaerobically induced lactate dehydrogenase: The role of a disulfide bridge in enzyme stability. Presented at XV Annual Midwest Autumn Enzymology Conference. Chicago, IL. Oct 15, 1995
71. Li, A.D., H.C. Huppe, F.J. Stevens, and L.E. Anderson. *Chlamydomonas reinhardtii* NADP-linked glyceraldehyde-3-phosphate dehydrogenase is DTT-activatable. Presented at XV Annual Midwest Autumn Enzymology Conference. Chicago, IL. Oct 15, 1995
72. Anderson L.E., A. D. Li, E.H. Muslin, M.E. Pacold, M.I. Donnelly, M. Schiffer, and F.J. Stevens 1995. Mechanism of light modulation: identification of redox-sensitive cysteines in target enzymes by modeling and by site-directed mutagenesis. Presented at XV Annual Midwest Autumn Enzymology Conference. Chicago, IL. Oct 15.
73. Muslin E.H., A. D. Li, F.J. Stevens, M. Schiffer, M. Donnelly, and L.E. Anderson. 1995. The engineering of new domain-locking disulfides into a bacterial malate dehydrogenase strengthens evidence for original model of reductive activation of the chloroplast enzyme. Presented at Photosynthesis Symposium, Turkey Run, IN, Oct.
74. Raffen, R., P. Wilkins Stevens, D.K. Hanson, Y. Deng, M. Berrios-Hammond, F.A. Westholm, M. Schiffer, and F.J. Stevens. 1996. In vitro characterization of light chain amyloidosis using recombinant light chain variable domains. Biophysical J.
75. Levine, D. and F.J. Stevens. 1996. STALK: A Molecular Docking System. Presented at The Second Sandia National Laboratories Workshop on Computational Molecular Biology, March 4-6, 1996, Albuquerque, New Mexico
76. Huang, D.-B., C. Ainsworth, C.-H. Chang, F.J. Stevens, and M. Schiffer. 1996. Structure of the variable domain of human immunoglobulin κIV light chain Len Presented at the International Union of Crystallography XVII Congress and General Assembly, Seattle, WA., Aug 8-17.
77. Stevens, F.J. 1996. Biophysical pathology of human antibody light chains. Presented at NYU Medical School, New York, Nov 26.

78. Khurana, R., F.J. Stevens, S. Meredith, and Y. Argon. 1997. Interaction of BiP with partially unfolded variable domain of immunoglobulin light chain. Presented at Keystone Conference, March.
79. Schiffer, M., D.-B. Huang, R. Raffen, R. Pokkuluri, P. Wilkins-Stevens, and F.J. Stevens. 1997. Major unexpected effect of a Gln to Glu substitution on a dimer interface. 1997 Proteins Gordon Conference.
80. Raffen, R., L.J. Dieckman, P. Wilkins Stevens, P.R. Pokkuluri, M. Schiffer, and F.J. Stevens. 1997. Characterizing primary structural determinants of immunoglobulin light chain amyloidosis with recombinant light chain variable domains. FASEB Summer Research Conference on "Amyloid and other Abnormal Protein Assembly, July 13-18, Copper Mt. CO.
81. Pokkuluri, P.R., D.-B. Huang, R. Raffen, P. Wilkins-Stevens, F.J. Stevens, and M. Schiffer. 1997. How do metal ions help (promote) the formation of protein crystals? The Annual American Crystallographic Association Meeting, July 19-25, St. Louis, MO.
82. Stevens, F.J. 1997. Conformational diseases of immunoglobulins. Presented at University of Chicago, Dept of Pathology, Oct 13.
83. Stevens, F.J. 1997. Amino acid sequences, protein biophysics and the pathological expression of light chains. Presented at the 30th Annual Meeting of the American Society of Nephrology, San Antonio, Nov 2-5.
84. Deret, S., J. Chomilier, F. Stevens, and P Aucouturier. 1997. Particularites structurales des chaines legeres d'immunoglobulines responsables de la maladie des depots de chaines legeres (LCDD). Presented at the 10 Annual Meeting of the GGMM (Groupe du Graphisme et Modelisation Moleculaire), Dourdan, France.
85. Stevens, F.J. 1998. Conformational diseases of antibody light chains. Presented at University of Illinois, Chicago, Feb 12.
86. Pokkuluri, P.R., G. Johnson, X. Cai, L. Dieckman, F.J. Stevens, and M. Schiffer. 1998. Domain flips resulting from single amino acid substitutions. American Crystallographic Association Meeting.
87. Schiffer, M., P.R. Pokkuluri, X. Cai, G. Johnson, R. Raffen, and F.J. Stevens. 1998. A single amino acid substitutions changes the mode of dimerization. Presented at The Thrid Annual Structural Biology Symposium, University of Texas Medical Branch at Galveston, April 3-5.
88. Dogar, J. and F.J. Stevens. 1998. Use of computerized homology modeling and threading techniques to predict immunodominant epitopes of Fel d 1 on MHC Class-II molecules. Presented at Aspen Allergy Conference, July 29 - Aug1.
89. Raffen, R., P.R. Pokkuluri, P. Wilkins Stevens, M. Schiffer, and F.J. Stevens. 1998. Characterization of immunoglobulin light chain amyloidosis by site-directed mutagenses. Presented at Protein Society Symposium, San Diego, CA., July 25-29.

90. Stevens, F.J. 1998. Applications of protein engineering: From a killer protein to detection of actinides. Presented at Northwestern University, Dept. of Biomedical Engineering, Dec 3.
91. Stevens, F.J., R. Raffen, L. J. Dieckman, M. Szpunar, P.R. Pokkuluri, P.W. Stevens, and M. Schiffer. 1999. Characterization of immunoglobulin light chain pathoenic aggregation and amyloidosis by site-directed mutagenesis. Presented at 4<sup>th</sup> Annual Midwest Stress Response and Chaperone Meeting., Jan 16.
92. Davis, D.P., S. Vogen, C. Case, R. Raffen, F. Stevens, and Y. Argon. 1999. Unlocking the mechanisms of immunoglobulin aggregation disorders. Presented at 4<sup>th</sup> Annual Midwest Stress Response and Chaperone Meeting, Jan 16.
93. Schiffer, M., P.R. Pokkulur, R. Raffen, X. Cai, G. Johnson, and F.J. Stevens. 1999. Alteration of dimerization mode by single amino substitutions. Presented at 13<sup>th</sup> Symposium of the Protein Society, Boston, MA, July 24-28.
94. Li, A.D., L.E. Anderson, M. Schiffer, and F.J. Stevens. 1999. Identifying potential disulfides by homology modeling. Presented at XVIth International Botanical Congress, St. Louis, Aug 1-7.
95. Anderson, L.E., A.D. Li, and F.J. Stevens. 1999. Identification of potential redox-sensitive cysteine residues in the reductase domains of two plant dihydrofolate reductase-thymidylate synthases. Presented at XVIth International Botanical Congress, St. Louis, Aug 1-7.
96. Anderson, L.E., A.D. Li, and F.J. Stevens. 1999. Finding redox-regulated enzymes by homology modeling. Presented at Russian Society for Plant Physiology meeting, Moscow, Oct.
97. Dong, A., Y.-S. Kim, R. Raffen, T.W. Randolph, A. Solomon, P. Wilkins Stevens, F.J. Stevens, and J.F. Carpenter. 2000. Entrapping aggregation intermediates of  $\alpha$ -helix and  $\beta$ -domain proteins by thermochemical denaturation: an infrared spectroscopic study. Presented at FASEB Conference: Amyloid and other Abnormal Protein Assembly Processes, Copper Mountain, CO, June 11-16.
98. Kim, Y.-S., S.P., E. Chi, R. Raffen, P. Wilkins Stevens, F.J. Stevens, A. Solomon, M.C. Manning, T.W. and J.F. Carpenter. 2000. Counteracting effects of renal solutes on amyloid fibril formation by immunoglobulin light chains. Presented at FASEB Conference: Amyloid and other Abnormal Protein Assembly Processes, Copper Mountain, CO, June 11-16.
99. Stevens, F.J. and S. Filer. 2000. Novel strategy using synthetic nucleic acids and conventional immunoassays for biological agent detection. Third International Symposium; Chemical and Biological Medical Treatment Symposia (CBMTS). AC-Laboratorium, Spiez, Switzerland, May 7-12.
100. Manning, M.C., R. Meglan, and F.J. Stevens. 2000. Decoding the sequences of antibody light chains: the power of multivariate statistical analysis. Presented at the 2000 Colorado Protein Stability Conference, Breckenridge, Colorado, July 13-15.
101. Davis, D.P., K. Sciarretta, S.M. Vogen, J. L. Dul, F.J. Stevens, and Y. Argon. 2001. Both somatic mutation and environmental factors govern the aggregation pathway. Canadian Society of Biochemistry and Molecular & Cellular Biology. 11<sup>th</sup> Winternational Conference: "Dynamics of

Intracellular Organelles and Molecular Machines.” Mont Ste. Anne, Feb 8-11.

102. Dul, J.L., D.P. Davis, E.K. Williamson, F.J. Stevens, and Y. Argon. 2001. Hsp70 and anti-fibrillrogenic peptides promote degradation and inhibit intracellular aggregation of amyloidogenic light chains. Canadian Society of Biochemistry and Molecular & Cellular Biology. 11<sup>th</sup> International Conference: “Dynamics of Intracellular Organelles and Molecular Machines.” Mont Ste. Anne, Feb 8-11.
103. Stevens, F.J. 2001. Protein structure, biophysics, and disease. Presented at Northern Illinois University, April 20.
104. Wan, K., M. Duan, Y.-M. Lin, Y.-M., J. Lin, R. Raffen, and F.J. Stevens. 2001. The study of protein-protein interactions by H/D exchange and ESI MS. Presented at 49<sup>th</sup> Annual Conference on Mass Spectrometry and Allied Topics, Chicago, IL, May 27-May31.
105. Stevens, F.J. 2001. Protein structure, biophysics, and disease. Presented at MediChem Research, Inc. Woodridge, IL, June 5.
106. Silver, R.B., J.R. Hummel, J.H. Christiansen, S.R. Filer, and F.J. Stevens. 2001. A multi-modal biosensor and decision system. Presented at 5<sup>th</sup> Joint Conference on Standoff and Detection, Williamsburg, VA, Sept 24-27, 2001.
107. Stevens, F.J. 2002. The challenge of the megaproteins: Speculations on the structures of BRCA1 and BRCA2. Presented at the University of Illinois at Chicago, Bioengineering Dept, Sept 20.
108. Stevens, F.J. 2003. BRCA1 and BRCA2: Structure predictions without significant homology: soluble domains, functional validation. Presented at the University of Chicago, Feb 14.
109. Stevens, F.J. 2003. New directions for analysis of megaproteins and other inhabitants of the twilight and midnight zones. Presented at Advanced Life Sciences, Woodridge, Il, June 13.
110. Stevens, F.J. 2003. Structural relevance of germline gene to light chain amyloidosis. Presented at Mayo Clinic, Rochester, MN, Sept 24.
111. Stevens, F.J. 2003. Structural insights from sequence analysis. Presented at Children’s Hospital of Philadelphia, Oct 1.
112. Stevens, F.J. 2003. Protein self-assembly: a normal process, a pathological process, a nanotechnical process. Presented at Argonne National Laboratory, Nanolunch seminar series, Oct 15.
113. Stevens, F.J. 2003. Recognition of protein fold (and sometimes function) from sequence at very low levels of identity. Presented at Bioinformatics Workshop, Ft. Detrick, Md. Nov 5.
114. Wilton, R, P. Saxena, and F.J. Stevens. Expression, purification, and biochemical characterization of the human receptor for advanced glycation endproducts (RAGE). Presented 18<sup>th</sup> Symposium of the Protein Society, San Diego, Aug 14-18, 2004.

## PUBLICATIONS (book chapters):

1. Stevens, F.J. and C. Peraino. 1982. Development of a model to simulate the characteristics of rat-liver tumor promotion by phenobarbital. In *Proceedings of the Symposium on Cocarcinogenesis and Biological Effects of Tumor Promoters*, eds. E. Hecker, N.E. Fusenig, W. Kunz, F. Marks, and H.W. Thielmann, pp. 105-109, Raven Press, New York.
2. Wu, T.T., P.W. Stevens, J.G. Hovis, and F.J. Stevens. 1982. Production of an evolutionary remnant enzyme in *Escherichia coli* K-12. In *Overproduction of Microbial Products*, eds. V. Krumphanzl, B. Sikyta, Z. Vanek, pp. 417-423. Academic Press, London.
3. Peraino, C., W.L. Richards, and F.J. Stevens. 1983. Multistage hepatocarcinogenesis. In *Mechanisms of Tumor Promotion, Vol.I, Tumor Promotion in Internal Organs*, ed. T.J. Slaga, pp. 1-53. CRC Press.
4. Stevens, F.J. and C. Peraino. 1983. Liver as a model system for analyzing mechanisms of tumor initiation and promotion. In *Organ and Species Specificity in Chemical Carcinogenesis*, eds. R. Langenbach, S. Nesnow and J.M. Rice, Plenum Press, pp. 231-252.
5. Stevens, F.J. 1989. Size-exclusion high-performance liquid chromatography in analysis of protein and peptide epitopes. *Methods in Enzymology* **178**: 107-130.
6. Stevens, F.J. 1991. Considerations of the interpretation of the specificity of monoclonal antibodies determined in solid-phase immunoassays. In *The Immunochemistry of Solid-Phase Immunoassays*, pp 233-242, ed. J.E. Butler, CRC Press, New York.
7. Spangler, B.D., G.D. Armstrong, and F.J. Stevens. 1992. HPLC analysis of pertussis toxin binding to glycoproteins, in *Bacterial Protein Toxins (5)*, pp 300-301, Witholt et al., eds., Gustav Fischer Verlag, Stuttgart.
8. Stevens, F.J. and M. Schiffer. 1995. Structure and properties of human immunoglobulin light chain dimers, in *Methods in Molecular Biology, Vol 51: Antibody Engineering Protocols*, ed. S. Paul., pp 51-81. Humana Press Inc, Totowa, NJ.
9. Schiffer, M. and F.J. Stevens. 1995. Crystallographic and chromatographic methods for study of antibody light chains and other proteins, in *Methods in Molecular Biology, Vol 51: Antibody Engineering Protocols*, ed. S. Paul., pp 83-98. Humana Press Inc, Totowa, NJ.
10. Anderson, L.E., H.C. Huppe, A.D. Li, and F.J. Stevens. 1996. The sedoheptulose bisphosphatase of *Chlamydomonas reinhardtii* contains a potential inter-domain disulfide and is redox-sensitive. Proceedings of Xth International Photosynthesis Congress, Montpellier, France. In *Photosynthesis from Light to Biosphere, Vol. V*, pp 71-74. Ed, P. Mathis.
11. Stevens, F.J., D.T. Weiss, and A. Solomon. 1998. Structural bases of light chain-related pathology. In *The Antibodies, Vol 5*, pp 175-208. M. Zanetti and J.D. Capra, eds. Harwood

Academic Publishers, Australia

12. Raffen, R. and Stevens, F.J. 1999. Small-zone, high-speed gel filtration chromatography to detect protein aggregation associated with light chain pathologies. *Meth. Enzymol.* **309**: 318-332.
13. Stevens, F.J., M. Schiffer, and Y. Argon. 2002. Thermodynamic aspects of immunoglobulin light chain disease. In *Conformational Diseases – A Compendium. Based on the First International Workshop on Conformational Diseases*, pp 135-150, ed B. Solomon, A. Taraboulos, and E. Katchalski-Katzir. The Center for the Study of Emerging Diseases.
14. Wilton, R., E.A. Myatt, and F.J. Stevens. 2003. SCIMMS: Simulated chromatography of interactive macromolecular systems. In *Protein-Protein Interactions: Methods and Protocols*, ed. H. Fu, Humana Press, Totowa, NJ.
15. Kay, B.K., M. Scholle, and F.J. Stevens 2004. EH domains and their ligands. (in press)
16. Stevens, F.J. 2004. Amyloid proteins. Immunoglobulin (AL, AH). In *Amyloid Proteins: The Beta Pleated Sheet Conformation and Disease*, ed J. Sipe, Wiley.

#### PUBLICATIONS (peer reviewed):

1. Stevens, F.J. and T.T. Wu. 1976. Growth on D-lyxose of a mutant strain of *Escherichia coli* K-12 using a novel isomerase and enzymes related to D-xylose metabolism. *J. Gen. Microbiol.* **97**: 257-265.
2. Stevens, F.J., H.S. Pankratz, and R.L. Uffen. 1977. Demonstration of two 3,3'-diaminobenzidine reactions associated with photosynthetic membranes in anaerobic, light-grown *Rhodospirillum rubrum*. *J. Histochem. Cytochem.* **25**: 1264-1268.
3. Stevens, F.J., F. Westholm, A. Solomon, and M. Schiffer. 1980. Self-association of human immunoglobulin κI light chains: role of the third hypervariable region. *Proc. Natl. Acad. Sci USA* **77**: 1144-1148.
4. Peraino, C., E. Staffeldt, D.A. Haugen, L.S. Lombard, F.J. Stevens, and R.J.M. Fry. 1980. Effects of varying the dietary concentration of phenobarbital on its enhancement of 2-acetylaminofluorene-induced hepatic tumorigenesis. *Cancer Res.* **40**: 3268-3273.
5. Boernke, W.E., F.J. Stevens, and C. Peraino. 1981. Effects of self-association of ornithine aminotransferase on its physico-chemical characteristics. *Biochemistry* **20**: 115-121.
6. Stevens, F.J., P.W. Stevens, J.G. Hovis, and T.T. Wu. 1981. Some properties of D-mannose isomerase from *Escherichia coli* K-12. *J. Gen. Microbiol.* **124**: 219-223.
7. Stevens, F.J., F.A. Westholm, N. Panagiotopoulos, A. Solomon, and M. Schiffer. 1981.

- Preliminary crystallographic data on the human  $\lambda$  III Bence Jones dimer Cle. J. Molec. Biol. **147**: 179-183.
8. Stevens, F.J., F.A. Westholm, N. Panagiotopoulos, M. Schiffer, R.A. Popp, and A. Solomon. 1981. Characterization and preliminary crystallographic data from the V<sub>L</sub>-related fragment of the human  $\kappa$ I Bence Jones protein Wat. J. Molec. Biol. **147**: 185-193.
  9. Stevens, F.J. and M. Schiffer. 1981. Computer simulation of protein self-association during small-zone gel filtration: estimation of equilibrium constants. Biochem. J. **195**: 213-219.
  10. Schiffer, M., F.J. Stevens, F.A. Westholm, D. Carlson, and B. Schoenborn. 1982. Small angle neutron scattering study of Bence Jones protein Mcg: comparison of structures in solution and in crystal. Biochemistry **21**: 2874-2878.
  11. Gierczak, J.S., F.J. Stevens, H.S. Pankratz, and R.L. Uffen. 1982. Cytochemical localization and measurement of aerobic 3,3'-diaminobenzidine oxidation reactions in photosynthetically grown *Rhodospirillum rubrum*. J. Histochem. Cytochem. **30**: 901-907.
  12. Boernke, W.E., F.J. Stevens, J.E. Edwards, and C. Peraino. 1982. Differential changes in ornithine aminotransferase self-affinity produced by exposure to basic amino acids and increases in the intrinsic electronegativity of the enzyme monomer. Arch. Biochem. Biophys. **216**: 152-157.
  13. Short, M.T., W.J. Eisler, and F.J. Stevens. 1983. Automated failsafe control system for independent operation of two radiation shutters on a rotating anode X-ray source. J. Appl. Crystal. **16**: 359-360.
  14. Short, M.T., F.A. Westholm, M. Schiffer, and F.J. Stevens. 1985. Comparison of chromatographic characteristics of homologous Bence Jones proteins during size -exclusion chromatography by HPLC and by Sephadex. J. Chromatog. **323**: 418-423.
  15. Chang, C.-H., M.T. Short, F.A. Westholm, F.J. Stevens, B.-C. Wang, W. Furey, A. Solomon, and M. Schiffer. 1985. A novel arrangement of immunoglobulin variable domains: X-ray crystallographic analysis of the  $\lambda$  chain dimer, Bence Jones protein Loc. Biochemistry **24**: 4890-4897.
  16. Schiffer, M., C.-H. Chang, and F.J. Stevens. 1985. Formation of an infinite  $\beta$ -sheet arrangement dominates the crystallization behavior of  $\lambda$ -type antibody light chains. J. Mol. Biol. **186**: 475-478.
  17. Peraino, C., B.A. Carnes, and F.J. Stevens. 1986. Evidence for growth heterogeneity among different phenotypes in the population of histochemically detectable altered hepatocyte foci induced in a single neonatal treatment with carcinogen. Carcinogenesis **7**: 191-192.
  18. Stevens, F.J. 1986. Analysis of protein-protein interaction by simulation of small-zone size-exclusion chromatography: application to an antibody-antigen association. Biochemistry **25**: 981-993.
  19. Stevens, F.J., D.A. LeBuis, W.J. Eisler, and C.F. Ainsworth. 1986. Macromolecular interactions:

- application of microcomputer-controlled, high-speed, size-exclusion chromatography. *Liquid Chromatography Gas Chromatography* **4**: 340-348.
20. LeBuis, D.A., F.J. Stevens, and W.J. Eisler. 1986. An interface to provide microcomputer control of a motorized liquid chromatography sample injection valve. *Liquid Chromatography Gas Chromatography* **4**: 663-666.
  21. Stevens, F.J., J. Jwo, W. Carperos, H. Köhler, and M. Schiffer. 1986. Relationships between liquid- and solid-phase antibody association characteristics: implications for the use of competitive ELISA techniques to map the spatial locations of idiotopes. *J. Immunol* **137**: 1937-1944.
  22. Stevens, F.J. and M. Schiffer. 1987. Immunoglobulin structural diversities and idiotypic expression. *International Rev. Immunol.* **2**: 357-378.
  23. Stevens, F.J. 1987. Modification of an ELISA-based procedure for affinity determination: correction necessary for use with bivalent antibody. *Molec. Immunol.* **24**: 1055-1060.
  24. Murao, S-I., F.J. Stevens, A. Ito, and E. Huberman. 1988. Myeloperoxidase: a myeloid cell nuclear antigen with DNA-binding properties. *Proc. Natl. Acad. Sci. USA* **85**: 1232-1236.
  25. Stevens, F.J., W. Carperos, W.J. Monafo, and N.S. Greenspan. 1988. Size-exclusion HPLC analysis of epitopes. *J. Immunol. Methods* **108**: 271-278.
  26. Peraino, C., B.A. Carnes, F.J. Stevens, E.F. Staffeldt, J.J. Russell, A. Prapuolenis, J.A. Blomquist, S.D. Vessilinovitch, and R.R. Maronpot. 1988. Comparative developmental and phenotypic properties of altered hepatocyte foci and hepatic tumors. *Cancer Res.* **48**: 4171-4178.
  27. Stevens, F.J., C.-H. Chang, and M. Schiffer. 1988. Dual conformations of an immunoglobulin light chain dimer: heterogeneity of antigen specificity and idiotope profile may result from multiple variable domain interaction mechanisms. *Proc. Natl. Acad. Sci. USA* **85**: 6895-6899.
  28. Schiffer, M., C.-H. Chang, V.M. Naik, and F.J. Stevens. 1988. Analysis of immunoglobulin domain interactions: evidence for dominant role of salt bridges. *J. Mol. Biol.* **203**: 799-802.
  29. Schiffer, M., C. Ainsworth, Z.-B. Xu, W. Carperos, K. Olsen, A. Solomon, F.J. Stevens, and C.-H. Chang. 1989. The structure of a second crystal form of Bence Jones protein Loc: strikingly different domain associations in two crystal forms of a single protein. *Biochemistry* **28**: 4066-4072.
  30. Stevens, F.J. 1989. Analysis of protein-protein interaction by simulation of small-zone size-exclusion chromatography: stochastic formulation of kinetic rate contributions to observed elution characteristics. *Biophysical Journal* **55**: 1155-1167.
  31. Myatt, E.A., P.T.P. Kaumaya, and F.J. Stevens. 1989. Applications of size-exclusion HPLC in the analyses of peptide epitopes. *BioChromatography* **4**: 282-289.
  32. Stevens, F.J. 1990. Fractionation of macromolecules in an alternating transverse electric field: simulation of the method. *J. Biochem. Biophys. Meth.* **20**: 275-292.

33. Stevens, F.J., A. Solomon, and M. Schiffer. 1991. Bence Jones proteins: powerful tool for fundamental study of protein chemistry and pathophysiology. *Biochemistry* **30**: 6803-6805.
34. Myatt, E.A., F.J. Stevens, and P.B. Sigler. 1991. Self-association properties of two dimeric phospholipases A-2. *J. Biol. Chem.* **266**: 16331-16335.
35. M. Schiffer, C.-H. Chang, and F.J. Stevens. 1992. The functions of tryptophan residues in membrane proteins. *Protein Engineering* **5**: 213-214.
36. Stevens, F.J. and E.A. Myatt. 1992. Polymerization of immunoglobulin domains: a model system for the development of facilitated macromolecular assembly. *Nanotechnology* **2**: 206-213.
37. Kolmar, H., C. Frisch, G. Kleemann, K. Goetz, F.J. Stevens, and H.-J. Fritz. 1994. Dimerization of Bence Jones proteins: linking the rate of transcription from an *Escherichia coli* promoter to the association constant of REIV. *Biol. Chem. Hoppe Seyler's Z.* **375**: 61-70.
38. Myatt E.A., F.A. Westholm, D. T. Weiss, A. Solomon, M. Schiffer, and F.J. Stevens. 1994. Pathogenic potential of monoclonal human immunoglobulin light chains: relationship of *in vitro* aggregation to *in vivo* organ deposition. *Proc. Natl. Acad. Sci.* **91**: 3034-3038.
39. Li, D., F.J. Stevens, M. Schiffer, and L.E. Anderson. 1994. Identification of redox-sensitive cysteines distal to catalytic site in light-activated chloroplae enzymes. *Biophys. J.* **67**: 29-35.
40. Peak, M.J., F.J. Stevens, J.G. Peak, and M.W.W. Adams. 1994. The hyperthermophilic glycolytic enzyme enolase in the archaeon, *Pyrococcus furiosus*: comparison with mesophilic enolases. *Arch. Biochm. Biophys.* **313**: 280-286
41. Huang, D.-B., C.-H. Chang, C. Ainsworth, A.T. Brünger, M. Eulitz, A. Solomon, F.J. Stevens, and M. Schiffer. 1994. Comparison of crystal structures of two homologous proteins: structural origin of altered domain interactions in immunoglobulin light chain dimers. *Biochemistry* **33**: 14848-14857
42. Myatt, E.A., F.J. Stevens, and C. Benjamin. 1994. Solution-phase binding of monoclonal antibodies to bee venom phospholipase A2. *J. Immunological Methods* **177**: 35-42.
43. Wilkins-Stevens, P., R. Raffen, D.K. Hanson, Y.-L. Deng, M. Berrios-Hammond, F.A. Westholm, C. Murphy, M. Eulitz, R. Wetzel, A. Solomon, M. Schiffer, and F.J. Stevens. 1995. Recombinant immunoglobulin variable domains generated from synthetic genes provide a system for *in vitro* characterization of light-chain amyloid proteins. *Protein Science* **4**: 421-432.
44. Anderson, L.E., D. Li, N. Prakash, F.J. Stevens. 1995. Identification of potential redox-sensitive cysteines in cytosolic forms of fructosebiphosphatase and glyceraldehyde-3-phosphate dehydrogenase. *Planta* **196**: 118-124.
45. Pacold, M.E., L.E. Anderson, D. Li, and F.J. Stevens. 1995. Redox-sensitivity and light modulation of enzyme activity in the rhodophytes *Gracilaria tikvahiae* and *Chondrus crispus*. *J. Phycol.* **31**: 297-301.

46. Muslin, E.H., D. Li, F.J. Stevens, M. Donnelly, M. Schiffer, and L. Anderson. 1995. Engineering a domain-locking disulfide into a bacterial malate dehydrogenase produces a redox-sensitive enzyme. *Biophys. J.* **68**: 2218-2223.
47. Pacold, M.E., F.J. Stevens, D. Li, and L.E. Anderson. 1995. The NADP-linked glyceraldehyde-3-P dehydrogenases of *Anabena variabilis* and *Synechocystis* PCC 6803 which lack one of the cysteines found in the higher plant enzymes are not reductively activated. *Photosyn. Res.* **43**: 125-130.
48. Paul, S., L. Li, P. Wilkins-Stevens, F.J. Stevens, and A. Solomon. 1995. Natural catalytic antibodies: peptide hydrolyzing activities of Bence Jones proteins and a VL domain. *J. Biol. Chem.* **270**: 15257-15261.
49. Stevens, F.J., E.A. Myatt, C.-H. Chang, F.A. Westholm, M. Eulitz, D.T. Weiss, C. Murphy, A. Solomon, and M. Schiffer. 1995. A molecular model for the formation of amyloid: Immunoglobulin light chains. *Biochemistry* **34**: 10697-10702.
50. Boernke, W.E., C. Sanville Millard, S.N. Kakar, V. Jackiw, M. Schiffer, F.J. Stevens, and M.I. Donnelly. 1995. Stringency of substrate specificity of *Escherichia coli* malate dehydrogenase. *Arch. Bioch. Biophys* **322**: 43-52.
51. Shearer, M.H., F. J. Stevens, F.A. Westholm, H.B. Jenson, T.C. Chanh, K.D. Carey, G.L. White, and R.C. Kennedy. 1995. Serological crossreactions among primate immunoglobulins. *Devel. and Comp. Immunol.* **19**: 547-557.
52. Gallo, G. F. Goni, F. Boctor, R. Vidal, A. Kumar, F.J. Stevens, B. Frangione, and J. Ghiso. 1996. Light chain cardiomyopathy: Structural analysis of the light chain tissue deposits. *Am J. Pathol* **148**: 1397-1406.
53. Huang,D.-B., C.A. Ainsworth, F.J. Stevens, and M. Schiffer. 1996. Three quaternary structures for a single protein. *Proc. Natl. Acad. Sci.* **93**: 7017-7021.
54. Anderson, L., H.C. Huppe, L. Dong, F.J. Stevens. 1996. Identification of a potential redox-sensitive inter-domain disulfide in the sedoheptulosebisphosphatase of *Chlamydomonas reinhardtii*. *Plant J.* **10**: 553-560.
55. Levine, D., M. Facello, P. Hallstrom, G. Reeder, B. Walenz, and F.J. Stevens. 1997 STALK: An interactive virtual molecular docking system. *Comp. Sci. and Engineer.* **4**:55-65.
56. Li, A.D, F.J. Stevens, H.C. Huppe, R. Kersanach, and L.E. Anderson. 1997. *Chlamydomonas reinhardtii* NADP-linked glyceraldehyde-3-phosphated dehydrogenase contains the same Cys residues identified as potentially domain-locking in the higher plant enzyme and is light activated. *Photosyn. Res.* **51**: 167-177.
57. Anderson, L.E., A. Dong Li, S.C. Nehrlich, M.H. Hill, and F.J. Stevens. 1997. The cytosolic fructose bisphosphatase of *Brassica napus* contains a new potential regulatory disulfide and is redox-sensitive. *Plant Sci.* **128**: 23-30.
58. Jiang X, E. Myatt, P. Lykos, and F.J. Stevens. 1997. Interaction between glycosaminoglycans

- and immunoglobulin light chains. *Biochemistry* **36**: 13187-13194.
59. Stevens, F.J., Anderson, L.E., A. Dong Li, and S. Salman Lateef. 1997. Identification of potential inter-domain disulfides in three higher plant mitochondrial citrate synthases:paradoxical differences in redox-sensitivity as compared with the animal enzyme *Photosyn. Res.* **54**: 185-197
60. Anderson, L.E., D. Li, E.H. Muslin, F.J. Stevens, and M. Schiffer. 1997. Predicting redox-sensitive cysteines in plant enzymes by homology modeling. *Comp.Rendu Acad. Sci, Paris* **320**: 767-781.
61. Deret, S., J. Chomilier, D.-B. Huang, J.-L. Preud'homme, F.J. Stevens, and P. Aucouturier. 1997. Molecular modelling of immunoglobulin light chains implicates hydrophobic residues in non-amyloid light chain deposition disease. *Protein Engineering* **10**: 1191-1197.
62. Anderson, L.E., A.D. Li, and F.J. Stevens. 1998. The enolases of ice plant and arabidopsis contain a potential disulfide and are redox-sensitive. *Phytochem* **47**: 707-713.
63. Muslin, E.H., F.J. Stevens, and L.E. Anderson. 1998. New domain-locking disulfides into a bacterial malate dehydrogenase strengthens evidence for original model of reductive activation of the chloroplast enzyme. *Photosyn. Res.* **55**: 75-82
64. Huang, D.-B., C.-H. Chang, C. Ainsworth, G. Johnson, M. Eulitz, A. Solomon, F.J. Stevens, and M. Schiffer. 1998. Variable domain structure of the human κIV light chain Len: high homology to the murine light chain McPC603. *Molec. Immunol.* **34**: 1291-1301.
65. Raffen, R., P. Wilkins Stevens, M. Schiffer, and F.J. Stevens. 1998. Reengineering immunoglobulin domain interactions by introduction of charged residues. *Protein Engineering* **11**: 303-309.
66. Schiffer, M., D.-B. Huang, P.R. Pokkuluri, R. Raffen, G. Johnson, P. Wilkins Stevens, and F.J. Stevens. 1998. A domain flip as a result of a single amino acid substitution. *Structure* **6**: 1067-1073.
67. Hellman, R., M. Vanhove, A. Leujeune, F.J. Stevens, and L.M. Hendershot. 1998. The *in vivo* association of BiP with newly synthesized proteins is dependent on the rate and stability of folding and not simply on the presence of sequences that can bind to BiP. *J. Cell. Biol.* **144**: 21-30.
68. Raffen, R., C. Wunsch, P.R. Pokkuluri, P. Dave, P. Wilkins Stevens, M. Schiffer, and F.J. Stevens. 1999. Physicochemical consequences of amino acid variations that contribute to fibril formation by immunoglobulin light chains. *Protein Science* **8**: 509-517.
69. Deret, S., L. Denoroy, M. Lamarine, R. Vidal, B. Mougenot, B. Frangione, F.J. Stevens, P.M. Ronco and P. Aucouturier. 1999. Myeloma-associated Fanconi's syndrome: Molecular analyses of monoclonal immunoglobulin light chains from patients with and without intracellular crystals. *Protein Engineering* **12**: 363-369.
70. Stevens, F.J. and Y. Argon. 1999. Pathogenic light chains and the B cell repertoire

Immunology Today **20**: 451-457.

71. Davis, D.P., Khurana, R., S. Meredith, F.J. Stevens, and Y. Argon. 1999. Mapping the major interaction between binding protein and Ig light chains to sites within the variable domains. *J. Immunol.* **163**: 3842-3850.
72. Stevens, F.J. and Y. Argon. 1999. Protein folding in the ER. *Semin. Cellular. Devel. Biol.* **10**: 443-454.
73. Pokkuluri, P.R., M. Klebig, A. Solomon, F.J. Stevens, and M. Schiffer. 1999. Tertiary structure of human  $\lambda$ 6 light chains. *Amyloid: Int. J. Exp. Clin. Invest.* **6**: 165-171.
74. Wall, J., M. Schel, C. Murphy, R. Hrncic, M. Klebig, R. Wetzel, F.J. Stevens, and A. Solomon. 1999. The thermodynamic stability of two  $\lambda$ 6 V<sub>L</sub> proteins correlates with fibrillogenicity. *Biochemistry* **38**: 14101-14108
75. Vidal, R., F. Goni, F.J. Stevens, P. Aucouturier, A. Kumar, B. Frangione, J. Ghiso, and G. Gallo. 1999. Somatic mutation of the L12a gene in V $\kappa$ I light chain deposition disease: potential effects on aberrant protein conformation and deposition. *Am. J. Pathol.* **155**: 2009-2017.
76. Stevens, F.J. and R. Kisilevsky. 2000. Immunoglobulin light chains, glycosaminoglycans and amyloid. *Cell. Molec. Life Sci.* **57**: 441-449.
77. Stevens, F.J. 2000. Four structural risk factors identify most fibril-forming kappa light chains. *Amyloid* **7**: 200-211.
78. Pokkuluri, P.R., X. Cai, F.J. Stevens, and M. Schiffer. 2000. Change in dimerization mode by removal of a single unsatisfied polar residue. *Protein Sci.* **9**: 1852-1855
79. Setterdahl, A., M. Hirasawa, L.M. Bucher, C.A. Dholakia, P. Jacquot, H. Yards, F. Miller, F.J. Stevens, D.B. Knaff, and L.E. Anderson. 2000. Oxidation-reduction properties of two engineered redox-sensitive mutant Escherichia coli malate dehydrogenases. *Arch. Bioch. Biophys.* **382**: 15-21
80. Davis, D.P., R. Raffen, J.L. Dul, S. Vogen, E.K. Williamson, F.J. Stevens, and Y. Argon. 2000. Inhibition of amyloid fiber assembly by both BiP and its target peptide. *Immunity* **13**: 433-442.
81. Stevens, F.J., P.R. Pokkuluri, and M. Schiffer. 2000. Protein conformation and disease: Pathological consequences of analogous changes in homologous proteins. *Biochemistry* **39**: 15291-15296.
82. Kim, Y.-S., S.P. Cape, E. Chi, R. Raffen, P. Wilkins-Stevens, F.J. Stevens, M.C. Manning, T.W. Randolph, A. Solomon, and J.F. Carpenter. 2000. Counteracting effects of renal solutes on amyloid formation by immunoglobulin light chains. *J. Biol. Chem.* **276**: 1626-1633.
83. Louise E. Anderson, L.E., A. Dong Li, E. H. Muslin, M. Schiffer, and F.J. Stevens. 2000. Identifying redox-sensitive extra-chloroplastic enzymes by homology modeling. *Physiol Plantarum* **110**: 296-303.

84. Dul, J.L., D.P. Davis, F.J. Stevens, and Y. Argon. 2001. Hsp70 and anti-fibrillogenic peptides promote degradation and inhibit intracellular aggregation of amyloidogenic light chains. *J. Cell. Biol.* **152**: 705-716.
85. Lin, Y-M., R. Raffen, Y. Zhou, M.T. Flavin, and F.J. Stevens. 2001. Amyloid fibril formation in microwell plates for screening of inhibitors. *Amyloid* **8**: 182-193.
86. Stevens, F.J. 2001. Caveat receptor: proteomes on display. *Comb. Chem. & High Throughput Screening* **4**: 599-602.
87. Davis, D., S. G. Gallo, S.M. Vogen, J.L. Dul, K.L. Sciarretta, A. Kumar, R. Raffen, F.J. Stevens, and Y. Argon. 2001. Both the environment and somatic mutations govern the aggregation pathway of pathogenic immunoglobulin light chains. *J. Mol. Biol.* **313**: 1043-1056.
88. Pokkuluri, P.R., R. Raffen, L. Dieckman, C. Boogaard, F.J. Stevens, and M. Schiffer. 2002. Increasing protein stability by polar surface residues: domain-wide consequences of interactions within a loop. *Biophysical J.* **82**: 391-398.
89. Zavaljevski, N., F.J. Stevens, and J. Reifman. 2002. Support vector machines for protein classification and identification of key amino acid positions. *Bioinformatics* **18**: 689-696.
90. Pokkuluri, P.R., M. Gu, X. Cai, R. Raffen, F.J. Stevens, and M. Schiffer. 2002. Factors contributing to decreased protein stability when aspartic acid residues are in  $\beta$ -sheet regions. *Protein Sci.* **11**:1687-1694
91. Kim, Y.-S., T.W. Randolph, F.J. Stevens, and J.F. Carpenter. 2002. Kinetics and energetics of assembly, nucleation, and growth of aggregates and fibrils for an amyloidogenic protein. *J. Biol. Chem.* **277**: 27240-27246.
92. Kim, Y-S., T.W. Randolph, M.C. Manning, F.J. Stevens, and J.F. Carpenter. 2003. Congo red populates partially unfolded states of an amyloidogenic protein to enhance aggregation and amyloid fibril formation. *J. Biol. Chem.* **278**: 10842-10850.
93. Vilim, R.B., R.M. Cunningham, B. Lu, P. Kheradpour, and F.J. Stevens. 2004. Fold-specific protein classifier for protein classification. *Bioinformatics* **20**: 839-846.
94. Coriu, D., A. Dispenzieri, F.J. Stevens, C.L. Murphy, S. Wang, D.T. Weiss and A. Solomon. 2004. Hepatic amyloidosis resulting from deposition of the apolipoprotein A-I variant Leu75Pro. *Amyloid* **10**: 215-223
95. Wu, W., J.T. Watson, F.J. Stevens, R.Yousefzai, and L.E. Anderson. 2004. Mass spectrometric evidence for an alternated disulfide bond in chloroplast fructose bisphosphatase. *Photosyn. Res.* **79**: 189-200.
96. Gidalevitz, T. C. Biswas, H. Ding, D. Duhovny, H.J. Wolfson, F. Stevens, S. Radford, and Y. Argon. (2004) Identification of the N-terminal peptide binding site of GRP94. *J. Biol. Chem.* **279**: 16543-16552
97. Stevens, F.J. 2004. Hypothetical structure of SAA: Helical bundle may be

- destabilized by proteolytic processing resulting in fibril formation. *Amyloid* 11: 71-80.
98. Stevens, F.J. 2004. Efficient recognition of protein fold in the twilight and midnight zones by conservative application of Psi-BLAST: Validation. *J. Molecular Recognition*. 18: 139-149.
  99. Stevens, F.J., C. Kuemmel, G. Babnigg, and F. Collart. 2004. Efficient recognition of protein fold in the twilight and midnight zones by conservative application of Psi-BLAST: Application. *J. Molecular Recognition* 18: 150-157.
  100. Stevens, F.J. 2004. Amyloid formation: An emulation of matrix protein assembly? *Amyloid* 11: 232-244.
  101. Wall, J. , V. Gupta, M. Wilkerson, R. Loris, P. Adams, A. Solomon, F.J. Stevens, and C. Dealwis. (2004) Structural basis of light chain amyloidogenicity: Comparison of the thermodynamic properties, fibrillogenic potential, and tertiary structural features of four V( $\lambda$ )6 proteins. *J. Molec. Recognition* 17: 323-331.
  102. Pirun, M., G. Babnigg, and F.J. Stevens. 2005. Template-based recognition of protein fold within the midnight and twilight zones of protein sequence similarity. *J. Molec. Recogn.* 18: 203-212.
  103. Bobrovnik, S.A. and F.J. Stevens. 2004. Evaluation of the intramolecular interaction of complementary domains, connected with a flexible chain. (in Russian) *Ukr. Biochem. J.* 76: 148-150.
  104. Yu, C., N. Zavaljevski, F.J. Stevens, K. Yackovich, and J. Reifman. 2005. Classifying noisy protein sequence data: a case study of immunoglobulin light chains. *Bioinformatics* (in press).
  105. Ghadge, G.D., L. Wang, K. Sharma, A.L. Monti, V. Bindokas, F.J. Stevens, and R.P. Roos. 2005. Electroporation of truncated and mutant SOD2 into the chick embryo spinal cord: A critical role for protein misfolding in neuronal death. *J. Neurology* (in press)
  106. Stevens, F.J. 2005. Modeling multiple p53 binding sites in BRCA1. (in preparation)
  107. Gu, M., R. Wilton, M. Szpunar, and F.J. Stevens. 2005. Fibril restriction by disulfide bonds: aggressive fibrillogenesis by RNaseA. (in preparation)
  108. Gu, M., P. Pokkuluri, M. Schiffer, and F.J. Stevens. 2005. Binding of ThioflavinT by non-fibrillar antibody variable domains. (in preparation)
  109. Stevens, F.J. and A. Solomon. 2005. Amyloidogenic mutations in ApolipoproteinA-1 may be predictable and segregate into three classes of energetic consequence. (in preparation)
  110. Stevens, F.J. and A. Solomon. 2005. The database of light chain variable domain sequences from patients with plasma cell diseases: A resource for study of folding disorders of immunoglobulins. (in preparation).
  111. F. Collart, S. Moy, L. Dieckman, M. Schiffer, J. Fackenthal, G. Green, A. Venkataraman, O.I. Olopade, and F.J. Stevens. 2005. Generation of functional subunits of BRCA1: Demonstration of p53 and estrogen receptor binding activities. (in preparation)