

Curriculum Vitae

DOUGLAS H. TURNER

Department of Chemistry
University of Rochester
Rochester, NY 14627-0216

E-mail: Douglas.Turner@rochester.edu

Personal Data:

Year of Birth: 1946
Citizenship: U.S.
Marital Status: Married to Joanna B. Olmsted; one child, Richard
Military Status: Captain, U. S. Army Reserve, inactive

Education:

1967 Harvard College, A.B. cum laude in Chemistry
1972 Columbia University, Ph.D., Physical Chemistry,
Advisors, G. W. Flynn and N. Sutin (Brookhaven National Labs)
1973-74 University of California, Berkeley, Postdoctoral Fellow,
Biophysical Chemistry, with I. Tinoco, Jr.

Professional Experience:

1975-81 Assistant Professor of Chemistry, University of Rochester
1981-85 Associate Professor of Chemistry, University of Rochester
1986- Professor of Chemistry, University of Rochester
1999-09 Professor of Pediatrics, University of Rochester,
School of Medicine and Dentistry

Honors:

National Institutes of Health Predoctoral Fellowship	1968-1972
Alfred P. Sloan Fellow	1979-1983
NIH Senior Fellowship, University of Colorado, Boulder	1984-1985
Member, Biomedical Sciences Study Section, NIH	1984-1988
Member, Editorial Board of <i>Biophysical Journal</i>	1992-1995, 1997-2003
Guggenheim Fellow and American Cancer Society Scholar	1993-1994
Co-chair, Nucleic Acids Gordon Conference	1995
Member, BBKA Study Section, NIH	1995-1999
Member, Editorial Board of <i>Folding and Design</i>	1996-1998
Associate-director, University of Rochester M.D./Ph.D. Program	1998-2014
Bridging Fellowship to Department of Pediatrics, University of Rochester	1999
Fellow, American Association for the Advancement of Science	1999
Member, International Advisory Board of the Institute of Bioorganic Chemistry, Polish Academy of Sciences, Poznan	2002-2013
Member, Biomedical Research and Research Training Subcommittee A, NIH	2004-2008
Gordon Hammes Lectureship sponsored by the journal, <i>Biochemistry</i> , and the Biological Chemistry Section of the American Chemical Society	2011

University of Rochester Doctoral Commencement Award for Lifetime Achievement in Graduate Education	2014
American Association for the Advancement of Science Poland – US Science Award (shared with Ryszard Kierzek)	2016
Honorary member, Phi Beta Kappa, Harvard College	2017
Total publications exceed 250	2022
Total citations exceed 20,000 (H-index > 70)	2022

Publications

1. "The Stimulated Raman Effect. A New Source of Laser Temperature-Jump Heating," J. V. Beitz, G. W. Flynn, D. H. Turner, and N. Sutin, *J. Am. Chem. Soc.*, 92, 4130-4132 (1970).
2. "Laser Raman Temperature-Jump Study of the Kinetics of the Triiodide Equilibrium. Relaxation Times in the 10^{-7} - 10^{-8} Second Range," D. H. Turner, G. W. Flynn, N. Sutin, and J. V. Beitz, *J. Am. Chem. Soc.*, 94, 1554-1559 (1972).
3. "Dimerization of Proflavin by the Laser Raman Temperature-Jump Method," D. H. Turner, G. W. Flynn, S. K. Lundberg, L. D. Faller, and N. Sutin, *Nature*, 239, 215-217 (1972).
4. "Raman Laser Temperature-Jump Studies," Ph.D. Thesis, Columbia University (1972).
5. "Rate of Intersystem Crossing between 1A and 5A States of an Iron(II) Complex in Solution," J. K. Beattie, N. Sutin, D. H. Turner and G. W. Flynn, *J. Am. Chem. Soc.*, 95, 2052-2054 (1973).
6. "Fluorescence Detected Circular Dichroism," D. H. Turner, I. Tinoco, Jr., and M. Maestre, *J. Am. Chem. Soc.*, 96, 4340-4342 (1974).
7. "Kinetics of the Stacking of Ethidium Bromide by the Raman Laser Temperature-Jump Method," D. H. Turner, R. Yuan, G. W. Flynn and N. Sutin, *Biophysical Chemistry*, 2, 385-389 (1974).
8. "The Kinetics of Codon-Anticodon Interaction in Yeast Phenylalanine Transfer RNA," K. Yoon, D. H. Turner, and I. Tinoco, Jr., *J. Mol. Biol.*, 99, 507-518 (1975).
9. "Fluorescence Detected Circular Dichroism Study of the Anticodon Loop of Yeast tRNA^{Phe}," D. H. Turner, I. Tinoco, Jr., and M. Maestre, *Biochemistry*, 14, 3794-3799 (1975).
10. "Some Very Rapid Reactions of Porphyrins in Aqueous Solution," R. F. Pasternack, N. Sutin, and D. H. Turner, *J. Am. Chem. Soc.*, 98, 1908-1913 (1976).
11. "Fluorescence Detected Circular Dichroism. Theory," I. Tinoco, Jr. and D. H. Turner, *J. Am. Chem. Soc.*, 98, 6453-6456 (1976).
12. "The Kinetics of Binding of U-U-C-A to a Dodecanucleotide Anticodon Fragment from Yeast tRNA^{Phe}," K. Yoon, D. H. Turner, I. Tinoco, Jr., F. von der Haar, and F. Cramer, *Nucleic Acids Research*, 3, 2233-2241 (1976).
13. "Solvent Effects on Stacking: A Kinetic and Spectroscopic Study of Thionine Association in Aqueous Alcohol Solutions," T. G. Dewey, P. S. Wilson, and D. H. Turner, *J. Am. Chem. Soc.*, 100, 4550-4554 (1978).
14. "Fluorescence-Detected Circular Dichroism," D. H. Turner, *Methods in Enzymology*, Vol. 49G, ed. by C. H. W. Hirs and S. N. Timasheff, Chapter 8, Academic Press, 1978.
15. "Raman Laser Temperature-Jump Kinetics," T. G. Dewey and D. H. Turner, *Adv. Molec. Relaxation*, 13, 331-350 (1978).
16. "Comparative Biophysical Analysis of Monoclonal Cryo and Noncryoprecipitable IgG1, IgG2 and IgG3 Globulins," C. D. Scoville and D. H. Turner, *Fed. Proc.*, 37, 1853 (1978).

17. "Photoselected Fluorescence Detected Circular Dichroism," E. W. Lobenstine and D. H. Turner, *J. Am. Chem. Soc.*, 101, 2205-2207 (1979).
18. "Laser Temperature Jump Study of Solvent Effects on Proflavin Stacking," T. G. Dewey, D. A. Raymond, and D. H. Turner, *J. Am. Chem. Soc.*, 101, 5822-5826 (1979).
19. "Spectroscopic and Kinetic Analysis of a Monoclonal IgG Cryoglobulin. Effect of Mild Reduction on Cryoprecipitation," C. D. Scoville, G. N. Abraham, and D. H. Turner, *Biochemistry*, 18, 2610-2615 (1979).
20. "Raman Laser Temperature Jump Study of Solvent Effects on Simple Stacking Systems," T. G. Dewey and D. H. Turner, "NATO Adv. Study Inst. Ser., Techniques and Applications of Fast Reactions in Solution," ed. by E. Wyn-Jones, D. Reidel, Pub., p. 235-238 (1979).
21. "Laser Temperature-Jump Study of Stacking in Adenylic Acid Polymers," T. G. Dewey and D. H. Turner, *Biochemistry*, 18, 5757-5762 (1979).
22. "Laser Temperature Jump Study of Solvent Effects on Poly (adenylic acid) Stacking," T. G. Dewey and D. H. Turner, *Biochemistry*, 19, 1681-1685 (1980).
23. "Study of the Kinetic and Structural Properties of a Monoclonal Immunoglobulin G Cryoglobulin," C. D. Scoville, D. H. Turner, J. L. Lippert, and G. N. Abraham, *J. Biol. Chem.*, 255, 5847-5852 (1980).
24. "Intersystem-Crossing Dynamics and Coordination Geometry Changes Observed by Ultrasonic and Laser Temperature-Jump Relaxation of the 2T_6A Spin Equilibrium of Hexadentate Iron (III) Complexes in Solution," R. A. Binstead, J. K. Beattie, T. G. Dewey and D. H. Turner, *J. Am. Chem. Soc.*, 102, 6442-6451 (1980).
25. "Further Verification of Fluorescence-Detected Circular Dichroism," E. W. Lobenstine and D. H. Turner, *J. Am. Chem. Soc.*, 102, 7786-7787 (1980).
26. "Thermodynamics of (dG-dC)₃ Double-Helix Formation in H₂O and D₂O," D. D. Albergo, L. A. Marky, K. J. Breslauer, and D. H. Turner, *Biochemistry*, 20, 1409-1413 (1981).
27. "Solvent Effects on Thermodynamics of Double-Helix Formation in (dG-dC)₃," D. D. Albergo and D. H. Turner, *Biochemistry*, 20, 1413-1418 (1981).
28. "Solvent Effects on the Kinetics and Thermodynamics of Stacking in Poly (cytidylic acid)," S. M. Freier, K. O. Hill, T. G. Dewey, L. A. Marky, K. J. Breslauer, and D. H. Turner, *Biochemistry*, 20, 1419-1426 (1981).
29. "Fluctuation in Optical Activity: A Probe of Fast Reactions Using Light Scattering," T. G. Dewey, D. H. Turner, and M. G. Sceats, *J. Chem. Phys.*, 74, 6592-6602 (1981).
30. "Fluorescence Detected Circular Dichroism of Proteins with Single Fluorescent Tryptophans," E. W. Lobenstine, W. C. Schaefer, and D. H. Turner, *J. Am. Chem. Soc.*, 103, 4936-4940 (1981).
31. "Why Do Nucleic Acids Form Helices?" D. H. Turner, M. Petersheim, D. D. Albergo, T. G. Dewey, and S. M. Freier, in "Bimolecular Stereodynamics," ed. by R. H. Sarma, pp. 429-438 (1981).
32. "Laser Crosslinking of *E. coli* RNA Polymerase and T7 DNA," C. A. Harrison, D. H. Turner,

and D. C. Hinkle, *Nucleic Acids Res.*, 10, 2399-2414 (1982).

33. "Base-Stacking and Pairing Contributions to Helix Stability: Thermodynamics of Double Helix Formation with CCGG, CCGGp, ACCGGp, CCGGAp CCGGUp, and ACCGGUp," M. Petersheim and D. H. Turner, *Biochemistry*, 22, 256-263 (1983).
34. "Nuclear Overhauser Studies of CCGGAp, ACCGGp, and ACCGGUp," M. Petersheim and D. H. Turner, *Biochemistry*, 22, 264-268 (1983).
35. "Proton Magnetic Resonance Melting Studies of CCGGp, CCGGAp, CCGGUp, ACCGGp, and ACCGGUp," M. Petersheim and D. H. Turner, *Biochemistry*, 22, 269-277 (1983).
36. "Solvent Effects on the Dynamics of (dG-dC)₃," S. M. Freier, D. D. Albergo, and D. H. Turner, *Biopolymers*, 22, 1107-1131 (1983).
37. "Effects of 3' Dangling End Stacking on the Stability of GGCC and CCGG Double Helices," S. M. Freier, B. J. Burger, D. Alkema, T. Neilson, and D. H. Turner, *Biochemistry*, 22, 6198-6206 (1983).
38. "Thermodynamic Studies of RNA Stability," S. M. Freier, M. Petersheim, D. R. Hickey, and D. H. Turner, *J. Biomolecular Structure and Dynamics*, 1, 1229-1242 (1984).
39. "Solvent Effects on the Stability of A₇U₇," D. R. Hickey and D. H. Turner, *Biochemistry*, 24, 2086-2094 (1985).
40. "Fluorescence-Detected Circular Dichroism of Ethidium in Vivo and Bound to Deoxyribonucleic Acid in Vitro," M. L. Lamos and D. H. Turner, *Biochemistry*, 24, 2819-2822 (1985).
41. "Effects of Terminal Mismatches on RNA Stability: Thermodynamics of Duplex Formation for ACCGGGp, ACCGGAp, and ACCGGCp," D. R. Hickey and D. H. Turner, *Biochemistry*, 24, 3987-3991 (1985).
42. "Contributions of Dangling End Stacking and Terminal Base Pair Formation to the Stabilities of XGGCCp, XCCGGp, XGGCCYp and XCCGGYp Helices," S. M. Freier, D. Alkema, A. Sinclair, T. Neilson, and D. H. Turner, *Biochemistry*, 24, 4533-4539 (1985).
43. "Improved Free Energies for G•C Base-Pairs," S. M. Freier, A. Sinclair, T. Neilson, and D. H. Turner, *J. Mol. Biol.*, 185, 645-647 (1985).
44. "Molecular Complexes of Nucleosides and Nucleotides with a Monomer Cationic Porphyrin and Some of Its Metal Derivatives," R. F. Pasternack, E. J. Gibbs, A. Gaudemer, A. Antebi, S. Bassner, L. De Poy, D. H. Turner, A. Williams, F. Laplace, M. H. Lansard, C. Merienne and M. Peree-Fauvet, *J. Am. Chem. Soc.*, 107, 8179-8186 (1985).
45. "Improved Parameters for Prediction of RNA Secondary Structure and Insights into Why RNA Forms Double Helices," D. H. Turner, S. M. Freier, N. Sugimoto, D. R. Hickey, J. A. Jaeger, A. Sinclair, D. Alkema, T. Neilson, M. H. Caruthers, and R. Kierzek, in "Structure and Dynamics of RNA," ed. by P. H. van Knippenberg, C. W. Hilbers, and L. Bosch, Plenum Press, pp. 1-13 (1986).
46. "Temperature-Jump Methods," D. H. Turner, in "Investigation of Rates and Mechanisms of Reactions, vol. VI, Part II," 4th edition, ed. by C. F. Bernasconi, Chapter 3, Techniques of Chemistry series, Wiley-Interscience, (1986).

47. "Fluorescence-Detected Circular Dichroism of Ethidium Bound to Poly(dG-dC) and Poly(dG-m⁵dC) under B- and Z-Form Conditions," M. L. Lamos, G. T. Walker, T. R. Krugh, and D. H. Turner, *Biochemistry*, 25, 687-691 (1986).
48. "Free Energy Contributions of G•U and Other Terminal Mismatches to Helix Stability," S. M. Freier, R. Kierzek, M. H. Caruthers, T. Neilson, and D. H. Turner, *Biochemistry*, 25, 3209-3213 (1986).
49. "Stability of XGCGCp, GCGCYp, and XGCGCYp Helices: An Empirical Estimate of the Energetics of Hydrogen Bonds in Nucleic Acids," S. M. Freier, N. Sugimoto, A. Sinclair, D. Alkema, T. Neilson, R. Kierzek, M. H. Caruthers, and D. H. Turner, *Biochemistry*, 25, 3214-3219 (1986).
50. "Fluorescence Detected Circular Dichroism of Ethidium Bound to Nucleic Acids," M. L. Lamos, E. W. Lobenstine, and D. H. Turner, *J. Am. Chem. Soc.*, 108, 4278-4284 (1986).
51. "Energetics of Internal GU Mismatches in Ribooligonucleotide Helices," N. Sugimoto, R. Kierzek, S. M. Freier, and D. H. Turner, *Biochemistry*, 25, 5755-5759 (1986).
52. "Polymer Supported RNA Synthesis and Its Application to Test the Nearest Neighbor Model," R. Kierzek, M. H. Caruthers, C. E. Longfellow, D. Swinton, D. H. Turner, and S. M. Freier, *Biochemistry*, 25, 7840-7846 (1986).
53. "Improved Free Energy Parameters for Prediction of RNA Duplex Stability," S. M. Freier, R. Kierzek, J. A. Jaeger, N. Sugimoto, M. H. Caruthers, T. Neilson, and D. H. Turner, *Proc. Natl. Acad. Sci. U.S.A.*, 83, 9373-9377 (1986). [by 9/2021 cited over 1900 times]
54. "Free Energy Increments for Hydrogen Bonds in Nucleic Acid Base Pairs," D. H. Turner, N. Sugimoto, R. Kierzek and S. D. Dreiker, *J. Am. Chem. Soc.*, 109, 3783-3785 (1987).
55. "Improved Parameters for Prediction of RNA Structure," D. H. Turner, N. Sugimoto, J. A. Jaeger, C. E. Longfellow, S. M. Freier, and R. Kierzek, *Cold Spring Harbor Symposium, Quant. Biol.*, 52, 123-133 (1987).
56. "Sequence Dependence for the Energetics of Dangling Ends and Terminal Base Pairs in Ribonucleic Acid," N. Sugimoto, R. Kierzek, and D. H. Turner, *Biochemistry*, 26, 4554-4558 (1987).
57. "Sequence Dependence for the Energetics of Terminal Mismatches in Ribooligonucleotides," N. Sugimoto, R. Kierzek, and D. H. Turner, *Biochemistry*, 26, 4559-4562 (1987).
58. "RNA Structure Prediction," D. H. Turner, N. Sugimoto and S. M. Freier, *Ann. Rev. Biophysics and Biophysical Chem.*, 17, 167-192 (1988). [7/2022 cited over 850 times]
59. "Kinetics for Reaction of a Circularized Intervening Sequence with CU, UCU, CUCU, and CUCUCU: Mechanistic Implications from the Dependence on Temperature, and on Oligomer and Mg²⁺ Concentrations," N. Sugimoto, R. Kierzek, and D. H. Turner, *Biochemistry*, 6384-6392 (1988).
60. "Hydrogen Bonding and Stacking Contributions to Nucleic Acid Stability," D. H. Turner, N. Sugimoto, S. D. Dreiker, S. M. Freier, and R. Kierzek, in *Structure and Expression, Vol. 1: From Proteins to Ribosomes*, ed. by R. H. Sarma and M. H. Sarma, p. 249-259 (1988).

61. "Effects of Substrate Structure on the Kinetics of Circle Opening Reactions of the Self-Splicing Intervening Sequence from *Tetrahymena thermophila*: Evidence for Substrate and Mg^{2+} Binding Interactions," N. Sugimoto, M. Tomka, R. Kierzek, P. Bevilacqua, and D. H. Turner, *Nucleic Acids Research*, 17, 355-371 (1989).
62. "Laser Temperature-Jump, Spectroscopic and Thermodynamic Study of Salt Effects on Duplex Formation by dGCATGC," A. P. Williams, C. E. Longfellow, S. M. Freier, R. Kierzek, and D. H. Turner, *Biochemistry*, 28, 4283-4291 (1989).
63. "Binding of a Fluorescent Oligonucleotide to a Circularized Intervening Sequence from *Tetrahymena thermophila*," N. Sugimoto, M. Sasaki, R. Kierzek, and D. H. Turner, *Chemistry Letters*, 2223-2226 (1989).
64. "Improved Predictions of Secondary Structures for RNA," J. A. Jaeger, D. H. Turner, and M. Zuker, *Proc. Natl. Acad. Sci. U.S.A.*, 86, 7706-7710 (1989).
65. "Predicting Optimal and Suboptimal Secondary Structure for RNA," J. A. Jaeger, D. H. Turner, and M. Zuker, *Methods in Enzymology*, Vol. 183, 280-305 (1990).
66. "Thermodynamic and Spectroscopic Study of Bulge Loops in Oligoribonucleotides," C. E. Longfellow, R. Kierzek, and D. H. Turner, *Biochemistry*, 29, 278-285 (1990).
67. "Thermodynamics and Kinetics of Base-Pairing and of DNA and RNA Self-Assembly and Helix-Coil Transition," D. H. Turner, N. Sugimoto, and S. M. Freier, in "Nucleic Acids," Vol. 1c by W. Saenger, Landolt-Bornstein series, Springer-Verlag, Berlin, p. 201-227 (1990).
68. "Effects of GA Mismatches on the Structure and Thermodynamics of RNA Internal Loops," J. SantaLucia, Jr., R. Kierzek, and D. H. Turner, *Biochemistry*, 29, 8813-8819 (1990).
69. "Melting and Chemical Modification of a Cyclized Self-Splicing Group I Intron: Similarity of Structures in 1 M Na^+ , in 10 mM Mg^{2+} , and in the Presence of Substrate," J. A. Jaeger, M. Zuker, and D. H. Turner, *Biochemistry*, 29, 10147-10158 (1990).
70. "Functional Group Substitutions as Probes of Hydrogen Bonding Between GA Mismatches in RNA Internal Loops," J. SantaLucia, Jr., R. Kierzek, and D. H. Turner, *J. Am. Chem. Soc.*, 113, 4313-4322 (1991).
71. "Thermodynamic Study of Internal Loops in Oligoribonucleotides: Symmetric Loops are More Stable than Asymmetric Loops," A. E. Peritz, R. Kierzek, N. Sugimoto, and D. H. Turner, *Biochemistry*, 30, 6428-6436 (1991).
72. "A Comparison of Optimal and Suboptimal RNA Secondary Structures Predicted by Free Energy Minimization with Structures Determined by Phylogenetic Comparison," M. Zuker, J. A. Jaeger, and D. H. Turner, *Nucleic Acids Res.*, 19, 2707-2714 (1991).
73. "Stabilities of Consecutive A•C, C•C, G•G, U•C, and U•U Mismatches in RNA Internal Loops: Evidence for Stable Hydrogen Bonded U•U and C•C+ Pairs," J. SantaLucia, Jr., R. Kierzek, and D. H. Turner, *Biochemistry*, 30, 8242-8251 (1991).
74. "Comparison of Binding of Mixed Ribose-Deoxyribose Analogues of CUCU to a Ribozyme and to GGAGAA by Equilibrium Dialysis: Evidence for Ribozyme Specific Interactions with 2' OH Groups," P. C. Bevilacqua and D. H. Turner, *Biochemistry*, 30, 10632-10640 (1991).

75. "Nearest Neighbor Parameters for GU Mismatches: 5'GU/3'UG is Destabilizing in the Contexts 5'CGUG/3'GUGC, 5'UGUA/3'AUGU, and 5'AGUU/3'UUGA, but Stabilizing in 5'GGUC/3'CUGG," L. He, R. Kierzek, J. SantaLucia, Jr., A. Walter, and D. H. Turner, *Biochemistry*, **30**, 11124-11132 (1991).
76. "Inhibition of Deoxyribo Oligonucleotides on the Circle Opening Reaction of the Intervening Sequence from *Tetrahymena thermophila*," N. Sugimoto, M. Sasaki, R. Kierzek, and D. H. Turner, *Chemistry Letters*, 747-748 (1990).
77. "Association of 2'-5' Oligoribonucleotides," R. Kierzek, L. He, and D. H. Turner, *Nucleic Acids Res.*, **20**, 1685-1690 (1992).
78. "Bulges in Nucleic Acids," D. H. Turner, invited review for *Current Opinion in Structural Biology*, **2**, 334-337 (1992).
79. "Context Dependence of Hydrogen Bond Free Energy Revealed by Substitutions in an RNA Hairpin," J. SantaLucia, Jr., R. Kierzek, and D. H. Turner, *Science*, **256**, 217-219 (1992).
80. "Dynamics of Ribozyme Binding of Substrate Revealed by Fluorescence Detected Stopped-Flow," P. C. Bevilacqua, R. Kierzek, K. A. Johnson, and D. H. Turner, *Science*, **258**, 1355-1358 (1992).
81. "Thermodynamic Considerations for Evolution by RNA," D. H. Turner and P. C. Bevilacqua, invited chapter in *The RNA World*, ed. by R. F. Gesteland and J. F. Atkins, Cold Spring Harbor Press, p. 447-464 (1993).
82. "Thermal Unfolding of a Group I Ribozyme: The Low Temperature Transition is Primarily Disruption of Tertiary Structure," A. R. Banerjee, J. A. Jaeger, and D. H. Turner, *Biochemistry*, **32**, 153-163 (1993).
83. "5'-Amino Pyrene Provides a Sensitive, Non-Perturbing Fluorescent Probe of RNA Secondary and Tertiary Structure Formation," R. Kierzek, D. H. Turner, Y. Li, and P. C. Bevilacqua, *J. Am. Chem. Soc.*, **115**, 4985-4992 (1993).
84. "Binding of Guanosine and 3' Splice Site Analogues to a Group I Ribozyme: Interactions with Functional Groups of Guanosine and with Additional Nucleotides," S. Moran, R. Kierzek, and D. H. Turner, *Biochemistry*, **32**, 5247-5256 (1993).
85. "Effect of a 5'-Phosphate on the Stability of Triple Helix," K. Yoon, C. A. Hobbs, A. E. Walter, and D. H. Turner, *Nucleic Acids Research*, **21**, 601-606 (1993).
86. "RNA Hairpin Loop Stability Depends on Closing Base Pair," M. J. Serra, M. H. Lyttle, T. J. Axenson, C. A. Schadt, and D. H. Turner, *Nucleic Acids Res.*, **21**, 3845-3849 (1993).
87. "Cooperative and Anticooperative Binding to a Ribozyme," P. C. Bevilacqua, K. A. Johnson, and D. H. Turner, *Proc. Natl. Acad. Sci. U.S.A.*, **90**, 8357-8361 (1993).
88. "A Novel NAD Derivative Produced During tRNA Splicing: ADP-ribose 1"-2" Cyclic Phosphate," G. M. Culver, S. M. McCraith, M. Zillman, R. Kierzek, N. Michaud, R. D. LaReau, D. H. Turner, and E. M. Phizicky, *Science*, **261**, 206-208 (1993).
89. "Structure of (rGGCGAGCC)₂ in Solution from NMR and Restrained Molecular Dynamics," J. SantaLucia, Jr. and D. H. Turner, *Biochemistry*, **32**, 12612-12623 (1993).

90. "Coaxial Stacking of Helices Enhances Binding of Oligoribonucleotides and Improves Predictions of RNA Folding," A. E. Walter, D. H. Turner, J. Kim, M. H. Lyttle, P. Müller, D. H. Mathews, and M. Zuker, *Proc. Natl. Acad. Sci. U.S.A.*, 91, 9218-9222 (1994).
91. "Fluorescence-Detected Stopped Flow with a Pyrene Labeled Substrate Reveals that Guanosine Facilitates Docking of the 5' Cleavage Site into a High Free Energy Binding Mode in the *Tetrahymena* Ribozyme," P. C. Bevilacqua, Y. Li, and D. H. Turner, *Biochemistry*, 33, 11340-11348 (1994).
92. "The Stability and Structure of Tandem GA Mismatches in RNA Depend on Closing Base Pairs," A. E. Walter, M. Wu, and D. H. Turner, *Biochemistry*, 33, 11349-11354 (1994).
93. "Sequence Dependence of Stability for Coaxial Stacking of RNA Helices with Watson-Crick Base Paired Interfaces," A. E. Walter and D. H. Turner, *Biochemistry*, 33, 12715-12719 (1994).
94. "A Model for the Stabilities of RNA Hairpins Based on a Study of the Sequence Dependence of Stability for Hairpins of Six Nucleotides," M. J. Serra, T. J. Axenson, and D. H. Turner, *Biochemistry*, 33, 14289-14296 (1994).
95. "Replacement of the Conserved G-U with a G-C Pair at the Cleavage Site of the *Tetrahymena* Ribozyme Decreases Binding, Reactivity, and Fidelity," A. M. Pyle, S. Moran, S. A. Strobel, T. Chapman, D. H. Turner, and T. R. Cech, *Biochemistry*, 33, 13856-13863 (1994).
96. "A Periodic Table of Symmetric Tandem Mismatches in RNA," M. Wu, J. A. McDowell, and D. H. Turner, *Biochemistry*, 34, 3204-3211 (1995).
97. "The Time Dependence of Chemical Modification Reveals Slow Steps in the Folding of a Group I Ribozyme," A. R. Banerjee and D. H. Turner, *Biochemistry*, 34, 6504-6512 (1995).
98. "Thermodynamic and Activation Parameters for Binding of a Pyrene-Labeled Substrate by the *Tetrahymena* Ribozyme: Docking is Not Diffusion Controlled and is Driven by a Favorable Entropy Change," Y. Li, P. C. Bevilacqua, D. H. Mathews, and D. H. Turner, *Biochemistry*, 34, 14394-14399 (1995).
99. "Predicting Thermodynamic Properties of RNA," M. J. Serra and D. H. Turner, *Methods in Enzymology*, 259, 242-261 (1995).
100. "A Mechanistic Framework for the Second Step of Splicing Catalyzed by the *Tetrahymena* Ribozyme," P. C. Bevilacqua, N. Sugimoto, and D. H. Turner, *Biochemistry*, 35, 648-658 (1996).
101. "Structural Features of a Six-Nucleotide RNA Hairpin Loop Found in Ribosomal RNA," M. A. Fountain, M. J. Serra, T. R. Krugh, and D. H. Turner, *Biochemistry*, 35, 6539-6548 (1996).
102. "Solution Structure of (rGCGGACGC)₂ by Two-Dimensional NMR and the Iterative Relaxation Matrix Approach," M. Wu and D. H. Turner, *Biochemistry*, 35, 9677-9689 (1996).
103. "Thermodynamics of Base Pairing," D. H. Turner, invited review, *Current Opinion in Structural Biology*, 6, 299-304 (1996).
104. "Dynamics of a Group I Ribozyme Detected by Spectroscopic Methods," D. H. Turner, Y. Li, M. Fountain, L. Profenno, and P. C. Bevilacqua, *Nucleic Acids and Molecular Biology*, 10, 19-

32 (1996).

105. "Thermodynamics of Coaxially Stacked Helices with GA and CC Mismatches," J. Kim, A. E. Walter, and D. H. Turner, *Biochemistry*, 35, 13753-13761 (1996).
106. "An Investigation of the Structural Basis for Thermodynamic Stabilities of Tandem GU Mismatches: Solution Structure of (rGAGGUCUC)₂ by Two-Dimensional NMR and Simulated Annealing," J. A. McDowell and D. H. Turner, *Biochemistry*, 35, 14077-14089 (1996).
107. "G·A and U·U Mismatches Can Stabilize RNA Internal Loops of Three Nucleotides," S. Schroeder, J. Kim, and D. H. Turner, *Biochemistry*, 35, 16105-16109 (1996).
108. "Secondary Structure Model of the RNA Recognized by the Reverse Transcriptase from the R2 Retrotransposable Element," D. H. Mathews, A. R. Banerjee, D. D. Luan, T. H. Eickbush, and D. H. Turner, *RNA*, 3, 1-16 (1997).
109. "Solution Structure of (rGGCAGGCC)₂ by Two-Dimensional NMR and the Iterative Relaxation Matrix Approach," M. Wu, J. SantaLucia, Jr., and D. H. Turner *Biochemistry*, 36, 4449-4460 (1997).
110. "Investigation of the Structural Basis for Thermodynamic Stabilities of Tandem GU Wobble Pairs: NMR Structures of (rGGAGUUC)₂ and (rGGAUGUCC)₂," J. A. McDowell, L. He, X. Chen, and D. H. Turner, *Biochemistry*, 36, 8030-8038 (1997).
111. "Effects of Mg²⁺ and the 2' OH of Guanosine on Steps Required for Substrate Binding and Reactivity with the *Tetrahymena* Ribozyme Reveal Several Local Folding Transitions," Y. Li and D. H. Turner, *Biochemistry*, 36, 11131-11139 (1997).
112. "Guanosine Binds to the *Tetrahymena* Ribozyme in More than One Step, and its 2'-OH and the Nonbridging *pro*-S_p Phosphoryl Oxygen at the Cleavage Site Are Required for Productive Docking," L. A. Profenno, R. Kierzek, S. M. Testa, and D. H. Turner, *Biochemistry*, 36, 12477-12485 (1997).
113. "Thermodynamics of Nonsymmetric Tandem Mismatches Adjacent to G·C Base Pairs in RNA," T. Xia, J. A. McDowell, and D. H. Turner, *Biochemistry*, 36, 12486-12497 (1997).
114. "The Crystal Structure of an RNA Oligomer Incorporating Tandem Adenosine-Inosine Mismatches," R. J. Carter, K. J. Baeyens, J. SantaLucia, D. H. Turner and S. R. Holbrook, *Nucleic Acids Res.*, 25, 4117-4122 (1997).
115. "A *Pneumocystis carinii* Group I Intron Ribozyme that Does Not Require 2' OH Groups on its 5' Exon Mimic for Binding to the Catalytic Core," S. M. Testa, C. G. Haidaris, F. Gigliotti, and D. H. Turner, *Biochemistry*, 36, 15303-15314 (1997).
116. "Measuring the Thermodynamics of RNA Secondary Structure Formation," J. SantaLucia, Jr. and D. H. Turner, *Biopolymers*, 44, 309-319 (1997).
117. "An Updated Recursive Algorithm for RNA Secondary Structure Prediction with Improved Thermodynamic Parameters," D. H. Mathews, T. C. Andre, J. Kim, D. H. Turner, and M. Zuker in *Molecular Modeling of Nucleic Acids* (Leontis, N. B., and SantaLucia, J., Jr., eds.), Am. Chem. Soc. Pub., New York p. 246-257 (1998).
118. "The Application of Thermodynamics to the Modeling of RNA Secondary Structure," D. H.

Mathews, J. M. Diamond, and D. H. Turner, in *Thermodynamics in Biology*, ed. by E. Di Cera, Oxford University Press, pp. 177-201 (2000).

119. "Antisense Binding Enhanced by Tertiary Interactions: Binding of Phosphorothioate and N3'→P5' Phosphoramidate Hexanucleotides to the Catalytic Core of a Group I Ribozyme from the Mammalian Pathogen *Pneumocystis carinii*," S. M. Testa, S. M. Gryaznov, and D. H. Turner, *Biochemistry*, 37, 9379-9385 (1998).
120. "Thermodynamic Parameters for an Expanded Nearest-Neighbor Model for Formation of RNA Duplexes with Watson-Crick Base Pairs," T. Xia, J. SantaLucia, Jr., M. E. Burkard, R. Kierzek, S. J. Schroeder, X. Jiao, C. Cox, and D. H. Turner, *Biochemistry*, 37, 14719-14735 (1998).
121. "The Interactions that Shape RNA," M. E. Burkard, D. H. Turner, and I. Tinoco, Jr., in *The RNA World*, 2nd edition, ed. by R. F. Gesteland, T. R. Cech and J. F. Atkins, Cold Spring Harbor Press, p. 233-264 (1999).
122. "Structures of Base Pairs Involving at Least Two Hydrogen Bonds," M. E. Burkard, D. H. Turner, and I. Tinoco, Jr., in *The RNA World*, Appendix 1, 2nd edition, ed. by R. F. Gesteland, T. R. Cech, and J. F. Atkins, Cold Spring Harbor Press, 675-680 (1999).
123. "Schematic Diagrams of Secondary and Tertiary Structure Elements," M. E. Burkard, D. H. Turner, and I. Tinoco, Jr., in *The RNA World*, Appendix 2, 2nd edition, ed. by R. F. Gesteland, T. R. Cech, and J. F. Atkins, Cold Spring Harbor Press, 681-685 (1999).
124. "Transient ADP-ribosylation of a 2'-phosphate Implicated in its Removal from Ligated tRNA during Splicing in Yeast," S. L. Spinelli, R. Kierzek, D. H. Turner, and E. M. Phizicky, *J. Biol. Chem.*, 274, 2637-2644 (1999).
125. "Thermodynamics of RNA Secondary Structure Formation," T. Xia, D. H. Mathews, and D. H. Turner, in *Prebiotic Chemistry, Molecular Fossils, Nucleosides, and RNA*, ed. by D. G. Söll, S. Nishimura, and P. B. Moore, Elsevier, p. 21-48 (1999) (reprinted in paperback as *RNA* (2001)).
126. "*In vitro* Suicide Inhibition of Self-Splicing of a Group I Intron from *Pneumocystis carinii* by an N3'→P5' Phosphoramidate Hexanucleotide," S. M. Testa, S. M. Gryaznov, and D. H. Turner, *Proc. Natl. Acad. Sci. U.S.A.*, 96, 2734-2739 (1999).
127. "Expanded Sequence Dependence of Thermodynamic Parameters Improves Prediction of RNA Secondary Structure," D. H. Mathews, J. Sabina, M. Zuker, and D. H. Turner, *J. Mol. Biol.*, 288, 911-940 (1999). [by 7/2022, cited over 3000 times]
128. "Thermodynamics of Unpaired Terminal Nucleotides on Short RNA Helices Correlates with Stacking at Helix Termini in Larger RNAs," M. E. Burkard, R. Kierzek, and D. H. Turner, *J. Mol. Biol.*, 290, 967-982 (1999).
129. "Thermodynamics of Single Mismatches in RNA Duplexes," R. Kierzek, M. E. Burkard, and D. H. Turner, *Biochemistry*, 38, 14214-14223 (1999).
130. "Thermodynamics of RNA-RNA Duplexes with 2- or 4-Thiouridines: Implications for Antisense Design and Targeting a Group I Intron," S. M. Testa, M. D. Disney, D. H. Turner, and R. Kierzek, *Biochemistry*, 38, 16655-16662 (1999).
131. "Predicting Oligonucleotide Affinity to Nucleic Acid Targets," D. H. Mathews, M. E. Burkard, S. M. Freier, J. R. Wyatt, and D. H. Turner, *RNA*, 5, 1458-1469 (1999).

132. "Algorithms and Thermodynamics for RNA Secondary Structure Prediction: A Practical Guide," M. Zuker, D. H. Mathews, and D. H. Turner, in RNA Biochemistry and Biotechnology, ed. by J. Barciszewski and B. F. C. Clark, Kluwer Academic Publishers, p. 11-43 (1999).
133. "RNA Secondary Structure Prediction," D. H. Mathews, D. H. Turner, and M. Zuker, in Current Protocols in Nucleic Acid Chemistry, ed. by S. Beaucage, D. E. Bergstrom, G. D. Glick, and R. A. Jones, John Wiley & Sons, p. 11.2.1-11.2.10 (2000).
134. "Expanded CUG Repeat RNAs form Hairpins that Activate the Double-Stranded-RNA-Dependent Protein Kinase PKR," B. Tian, R. J. White, T. Xia, S. Welle, D. H. Turner, M. B. Mathews, and C. A. Thornton, RNA, 6, 79-87 (2000).
135. "Targeting a *Pneumocystis carinii* Group I Intron with Methylphosphonate Oligonucleotides: Backbone Charge is Not Required for Binding or Reactivity," M. D. Disney, S. M. Testa, and D. H. Turner, Biochemistry, 39, 6991-7000 (2000).
136. "Nuclear Magnetic Resonance Spectroscopy and Molecular Modeling Reveal that Different Hydrogen Bonding Patterns are Possible for G·U Pairs: One Hydrogen Bond for each G·U Pair in r(GGCGUGCC)₂ and Two for each G·U Pair in r(GAGUGCUC)₂," X. Chen, J. A. McDowell, R. Kierzek, T. R. Krugh, and D. H. Turner, Biochemistry, 39, 8970-8982 (2000).
137. "Factors Affecting the Thermodynamic Stability of Small Asymmetric Internal Loops in RNA," S. J. Schroeder and D. H. Turner, Biochemistry, 39, 9257-9274 (2000).
138. "Conformational Changes," D. H. Turner, Chapter 8 in Nucleic Acids: Structures, Properties and Functions ed. by V. A. Bloomfield, D. M. Crothers and I. Tinoco, Jr. with chapters by J. Hearst, P. Kollman, D. Wemmer and D. H. Turner, University Science Press (2000).
139. "The Chemical Synthesis of Oligoribonucleotides with Selectively Placed 2'-O-Phosphates," R. Kierzek, M. A. Steiger, S. L. Spinelli, D. H. Turner, and E. M. Phizicky, Nucleosides, Nucleotides & Nucleic Acids 19, 917-933 (2000).
140. "NMR Structures of r(GCAGGCGUGC)₂ and Determinants of Stability for Single Guanosine-Guanosine Base Pairs," M. E. Burkard and D. H. Turner, Biochemistry, 39, 11748-11762 (2000).
141. "Contributions of Individual Nucleotides to Tertiary Binding of Substrate by a *Pneumocystis carinii* Group I Intron," M. D. Disney, S. M. Gryaznov, and D. H. Turner, Biochemistry, 39, 14269-14278 (2000).
142. "Stability and Structure of RNA Duplexes Containing Isoguanosine and Isocytidine," X. Chen, R. Kierzek, and D. H. Turner, J. Am. Chem. Soc., 123, 1267-1274 (2001).
143. "Thermodynamics of RNA Internal Loops with a Guanosine-Guanosine Pair Adjacent to Another Noncanonical Pair," M. E. Burkard, T. Xia, and D. H. Turner, Biochemistry, 40, 2478-2483 (2001).
144. "Use of Fluorescence Spectroscopy to Elucidate RNA Folding Pathways," P. C. Bevilacqua and D. H. Turner, in Current Protocols in Nucleic Acid Chemistry, ed. by S. Beaucage, D. E. Bergstrom, G. D. Glick, and R. A. Jones, John Wiley & Sons, p. 11.8.1-11.8.6.
145. "Use of Chemical Modification to Elucidate RNA Folding Pathways," D. H. Mathews and D. H. Turner, in Current Protocols in Nucleic Acid Chemistry, ed. by S. Beaucage, D. E. Bergstrom, G. D. Glick, and R. A. Jones, John Wiley & Sons, p. 11.9.1-11.9.4.

146. "Recognition Elements for 5' Exon Substrate Binding to the *Candida albicans* Group I Intron," M. D. Disney, C. G. Haidaris, and D. H. Turner, *Biochemistry*, 40, 6507-6519 (2001).
147. "Binding Enhancement by Tertiary Interactions and Suicide Inhibition of a *Candida albicans* Group I Intron by Phosphoramidate and 2'-O-Methyl Hexanucleotides," M. D. Disney, T. Matray, S. M. Gryaznov, and D. H. Turner, *Biochemistry*, 40, 6520-6526 (2001).
148. "Thermodynamics of Three-Way Multibranch Loops in RNA," J. M. Diamond, D. H. Turner, and D. H. Mathews, *Biochemistry*, 40, 6971-6981 (2001).
149. "The Energetics of Small Internal Loops in RNA," S. J. Schroeder, M. E. Burkard, and D. H. Turner, *Biopolymers Nucleic Acid Sci.*, 52, 157-167 (2001).
150. "Thermodynamic Stabilities of Internal Loops with GU Closing Pairs in RNA," S. J. Schroeder and D. H. Turner, *Biochemistry*, 40, 11509-11517 (2001).
151. "Long-Range Cooperativity in Molecular Recognition of RNA by Oligodeoxynucleotides with Multiple C5-(1-Propynyl) Pyrimidines," T. W. Barnes III, and D. H. Turner, *J. Am. Chem. Soc.*, 123, 4107-4118 (2001a).
152. "Long-Range Cooperativity due to C5-Propynylation of Oligopyrimidines Enhances Specific Recognition by Uridine of ribo-Adenosine Over ribo-Guanosine," T. W. Barnes III and D. H. Turner, *J. Am. Chem. Soc.*, 123, 9186-9187 (2001b).
153. "C5-(1-Propynyl)-2'-deoxy-Pyrimidines Enhance Mismatch Penalties of DNA:RNA Duplex Formation," T. W. Barnes III and D. H. Turner, *Biochemistry*, 40, 12738-12745 (2001c).
154. "Substrate Recognition by a Yeast 2'-Phosphotransferase Involved in tRNA Splicing and by its *E. coli* Homolog," M. A. Steiger, R. Kierzek, D. H. Turner, and E. M. Phizicky, *Biochemistry*, 40, 14098-14105 (2001).
155. "Experimentally Derived Nearest Neighbor Parameters for the Stability of RNA Three- and Four-Way Multibranch Loops," D. H. Mathews and D. H. Turner, *Biochemistry*, 41, 869-880 (2002).
156. "Dyalign: An Algorithm for Finding the Secondary Structure Common to Two RNA Sequences," D. H. Mathews and D. H. Turner, *J. Mol. Biol.*, 317, 191-203 (2002).
157. "Molecular Recognition by the *Candida albicans* Group I Intron: Tertiary Interactions with an Imino G•A Pair Facilitate Binding of the 5' Exon and Lower the K_M for Guanosine," M. D. Disney and D. H. Turner, *Biochemistry*, 41, 8113-8119 (2002).
158. "Oligonucleotide Directed Misfolding of RNA Inhibits *Candida albicans* Group I Intron Splicing," J. L. Childs, M. D. Disney, and D. H. Turner, *Proc. Natl. Acad. Sci. USA*, 99, 11091-11096 (2002).
159. "Sheared A_{anti}•A_{anti} Base Pairs in a Destabilizing Internal Loop: The NMR Structure of 5'(rGGCAAGCCU)₂," B. M. Znosko, M. E. Burkard, S. J. Schroeder, T. R. Krugh, and D. H. Turner, *Biochemistry*, 41, 14969-14977 (2002).
160. "Molecular Recognition in Purine-Rich Internal Loops: Thermodynamics, Structural, and Dynamic Consequences of Purine for Adenine Substitutions in 5'(rGGCAAGCCU)₂," B. M. Znosko, M. E. Burkard, T. R. Krugh, and D. H. Turner, *Biochemistry*, 41, 14978-14987 (2002).

161. "Uptake and Antifungal Activity of Oligonucleotides in *Candida albicans*," M. D. Disney, C. G. Haidaris, and D. H. Turner, *Proc. Natl. Acad. Sci. USA*, 100, 1530-1534 (2003).
162. "NMR Studies of DNA Single Strands and DNA:RNA Hybrids with and without 1-Propynylation at C5 of Oligopyrimidines," B. M. Znosko, T. W. Barnes, III, T. R. Krugh, and D. H. Turner, *J. Am. Chem. Soc.*, 125, 6090-6097 (2003).
163. "Inhibition of *Escherichia coli* RNase P by Oligonucleotide Directed Misfolding of RNA," J. L. Childs, A. W. Poole, and D. H. Turner, *RNA*, 9, 1437-1445 (2003).
164. "Thermodynamic Stability and Structural Features of the J4/5 Loop in a *Pneumocystis carinii* Group I Intron," S. J. Schroeder, M. A. Fountain, S. D. Kennedy, P. J. Lukavsky, J. D. Puglisi, T. R. Krugh, and D. H. Turner, *Biochemistry*, 42, 14184-14196 (2003).
165. "New Approaches to Targeting RNA with Oligonucleotides: Inhibition of Group I Intron Self-Splicing," M. D. Disney, J. L. Childs, and D. H. Turner, *Biopolymers*, 73, 151-161 (2004).
166. "Incorporating Chemical Modification Constraints into a Dynamic Programming Algorithm for Prediction of RNA Secondary Structure," D. H. Mathews, M. D. Disney, J. L. Childs, S. J. Schroeder, M. Zuker, and D. H. Turner, *Proc. Natl. Acad. Sci. USA*, 101, 7287-7292 (2004). [Cited over 1000 times]
167. "Structures of Two RNA Octamers Containing Tandem GA Base Pairs," S. B. Jang, K. Baeyens, M. S. Jeong, J. SantaLucia, Jr., D. Turner, and S. R. Holbrook, *Acta Cryst.* D60, 829-835 (2004).
168. "Secondary Structure Models of the 3' Untranslated Regions of Diverse R2 RNAs," A. M. Ruschak, D.H. Mathews, A. Bibillo, S. L. Spinelli, J. A. Childs, T. H. Eickbush, and D. H. Turner, *RNA*, 10, 978-987 (2004).
169. "Factors Affecting Thermodynamic Stabilities of RNA 3X3 Internal Loops," G. Chen, B. M. Znosko, X. Jiao, and D. H. Turner, *Biochemistry*, 43, 12865-12876 (2004).
170. "Structural Features and Thermodynamics of the J4/5 Loop from the *Candida albicans* and *Candida dubliniensis* Group I Introns," B. M. Znosko, S. D. Kennedy, P. C. Wille, T. R. Krugh, and D. H. Turner, *Biochemistry*, 43, 15822-15837 (2004).
171. "Hoechst 33258 Selectively Inhibits Group I Intron Self-Splicing by Affecting RNA Folding," M. D. Disney, J. L. Childs, and D. H. Turner, *ChemBioChem* 5, 1647-1652 (2004).
172. "Activity of Hoechst 33258 Against *Pneumocystis carinii* f. sp. muris, *Candida albicans*, and *Candida dubliniensis*," M. D. Disney, R. Stephenson, T. W. Wright, C. G. Haidaris, D. H. Turner, and F. Gigliotti, *Antimicrobial Agents and Chemotherapy* 49, 1326-1330 (2005).
173. "Solution Structure of an RNA Internal Loop with Three Consecutive Sheared GA Pairs," G. Chen, B. M. Znosko, S. D. Kennedy, T. R. Krugh, and D. H. Turner, *Biochemistry*, 44, 2845-2856 (2005).
174. "The influence of locked nucleic acid residues on the thermodynamic properties of 2'-O-methyl RNA/RNA heteroduplexes," E. Kierzek, A. Ciesielska, K. Pasternak, D. H. Mathews, D. H. Turner, and R. Kierzek, *Nucleic Acids Res.*, 33, 5082-5093 (2005).
175. "RNA Challenges for Computational Chemists," I. Yildirim and D. H. Turner, *Biochemistry* 44, 13225-13234 (2005).

176. "Predicting RNA Secondary Structure," D. H. Mathews, S. J. Schroeder, D. H. Turner, and M. Zuker, in *The RNA World*, 3rd edition, ed. By R. F. Gesteland, T. R. Cech, and J. F. Atkins, Cold Spring Harbor Press, pp. 631-657(2006).
177. "Facilitating RNA Structure Prediction with Microarrays," E. Kierzek, R. Kierzek, D. H. Turner, and I. E. Catrina, *Biochemistry*, 45, 581-593 (2006).
178. "Consecutive GA Pairs Stabilize Medium Size RNA Internal Loops," G. Chen and D. H. Turner, *Biochemistry* 45, 4025-4043 (2006).
179. "An Alternating Sheared AA Pair and Elements of Stability for a Single Sheared Purine-Purine Pair Flanked by Sheared GA Pairs," G. Chen, S. D. Kennedy, T. R. Krugh, and D. H. Turner, *Biochemistry* 45, 6889-6903 (2006).
180. "The Crystal Structure at 1.5 Å Resolution of an RNA Octamer Duplex Containing Tandem G-U Base Pairs," S. B. Jang, L-W. Hung, M. S. Jeong, E. L. Holbrook, X. Chen, D. H. Turner, and S. R. Holbrook, *Biophysical J.*, 90, 4530-4537 (2006).
181. "Prediction of RNA Secondary Structure by Free Energy Minimization," D. H. Mathews and D. H. Turner, *Current Opinions in Structural Biology* 16, 270-278 (2006).
182. "Nearest Neighbor Parameters for Watson-Crick Complementary Heteroduplexes Formed between 2'-O-methyl RNA and RNA Oligonucleotides," E. Kierzek, D. H. Mathews, A. Ciesielska, D. H. Turner, and R. Kierzek, *Nucleic Acids Res.*, 34, 3609-3614 (2006).
183. "Interpreting Oligonucleotide Microarray Data to Determine RNA Secondary Structure: Application to the 3' End of *Bombyx mori* R2 RNA," S. Duan, D. H. Mathews, and D. H. Turner, *Biochemistry*, 45, 9819-9832 (2006).
184. "The NMR Structure of an Internal Loop from 23S Ribosomal RNA Differs from its Structure in Crystals of 50S Ribosomal Subunits," N. Shankar, S. D. Kennedy, G. Chen, T. R. Krugh, and D. H. Turner, *Biochemistry*, 45, 11776-11789 (2006).
185. "A Set of Nearest Neighbor Parameters for Predicting the Enthalpy Change of RNA Secondary Structure Formation," Z. J. Lu, D. H. Turner, and D. H. Mathews, *Nucleic Acids Res.*, 34, 4912-4924 (2006).
186. "NMR Structures of (rGCUGAGGCU)₂ and (rGCGGAUGCU)₂: Probing the Structural Features that Shape the Thermodynamic Stability of GA Pairs," B. S. Tolbert, S. D. Kennedy, S. J. Schroeder, T. R. Krugh, and D. H. Turner, *Biochemistry*, 46, 1511-1522 (2007).
187. "RNA Secondary Structure Prediction," D. H. Mathews, D. H. Turner, and M. Zuker, *Current Protocols in Nucleic Acid Chemistry*, Supplement 28, 11.2.1-11.2.17 (2007).
188. "Stacking Effects on Local Structure in RNA: The Structure of Tandem GA Pairs Changes When Flanking GC Pairs are Replaced by isoG-isoC Pairs," G. Chen, R. Kierzek, I. Yildirim, T. R. Krugh, D. H. Turner, and S. D. Kennedy, *J. Phys. Chem. B*, 111, 6718-6727 (2007).
189. "The Chemical Synthesis of 2'-O-methyl-2,6-diaminopurine Riboside and LNA-2,6-diaminopurine Riboside and their Influence on the Thermodynamic Properties of 2'-O-methyl RNA/RNA Heteroduplexes." A. Pasternak, E. Kierzek, K. Pasternak, D.H. Turner, and R. Kierzek, *Nucleic Acids Res.*, 35, 4055-4063 (2007).

190. "NMR Reveals Absence of Hydrogen Bonding in Adjacent UU and AG Mismatches in an Isolated Internal Loop from Ribosomal RNA," N. Shankar, T. Xia, S.D. Kennedy, T.R. Krugh, D.H. Mathews, and D.H. Turner, *Biochemistry*, **46**, 12665-12678 (2007).
191. "Selective Nanoparticle Quenching of Fluorescence from Unbound Oligonucleotides as a Probe of RNA Structure," H. Li, R. Liang, D.H. Turner, L.J. Rothberg, and S. Duan, *RNA*, **13**, 2034-2041 (2007).
192. "Isoenergetic Penta- and Hexanucleotide Microarray Probing and Chemical Mapping Provide a Secondary Structure Model for an RNA Element Orchestrating R2 Retrotransposon Protein Function," E. Kierzek, R. Kierzek, W.N. Moss, S.M. Christensen, T.H. Eickbush, and D.H. Turner, *Nucleic Acids Res.*, **36**, 1770-1782 (2008).
193. "The thermodynamics of 3'-terminal pyrene and guanosine for the design of isoenergetic 2'-O-methyl-RNA-LNA chimeric oligonucleotide probes of RNA structure," A. Pasternak, E. Kierzek, K. Pasternak, A. Fraczak, D.H. Turner, and R. Kierzek, *Biochemistry*, **47**, 1249-1258 (2008).
194. "NMR-Assisted Prediction of RNA Secondary Structure: Identification of a Probable Pseudoknot in the Coding Region of an R2 Retrotransposon," J.M. Hart, S.D. Kennedy, D.H. Mathews, and D.H. Turner, *J. Am. Chem. Soc.* **130**, 10233-10239 (2008).
195. "Contributions of Stacking, Preorganization, and Hydrogen Bonding to the Thermodynamic Stability of Duplexes between RNA and 2'-O-methyl RNA with Locked Nucleic Acids (LNA)," E. Kierzek, A. Pasternak, K. Pasternak, Z. Gdaniec, I. Yildirim, D. Turner, and R. Kierzek, *Biochemistry*, **48**, 4377-4387 (2009).
196. "A CA⁺ Pair Adjacent to a Sheared GA or AA Pair Stabilizes Size-Symmetric RNA Internal Loops," G. Chen, S.D. Kennedy, and D.H. Turner, *Biochemistry*, **48**, 5738-5752 (2009).
197. "Secondary Structures for 5' Regions of R2 Retrotransposon RNAs Reveal a Novel Conserved Pseudoknot and Regions that Evolve under Different Constraints," E. Kierzek, S.M. Christensen, T.H. Eickbush, R. Kierzek, D.H. Turner, and W.N. Moss, *J. Mol. Biol.*, **390**, 428-442 (2009).
198. "Effects of Restrained Sampling Space and Non-planar Amino Groups on Free Energy Predictions for RNA with Imino and Sheared Tandem GA Base Pairs Flanked by GC, CG, iGiC or iCiG Base Pairs," I. Yildirim, H. Stern, J. Sponer, N. Spackova, and D.H. Turner, *J. Chem. Theory and Computation*, **5**, 2088-2100 (2009).
199. "Optical Melting Measurements of Nucleic Acid Thermodynamics," S.J. Schroeder and D.H. Turner, in *Methods in Enzymology*, **468**, 371-387, ed. by D. Herschlag, Academic Press, Burlington, (2009).
200. "The Chemical Synthesis of LNA-2-thiouridine and its Influence on Stability and Selectivity of Oligonucleotide Binding to RNA," M. Carlucci, E. Kierzek, A. Olejnik, D.H. Turner, and R. Kierzek, *Biochemistry* **48**, 10882-10893 (2009).
201. "Folding and Finding RNA Secondary Structure," D.H. Mathews, W.N. Moss, and D.H. Turner, in *The RNA Worlds*, ed. by J. F. Atkins, R. F. Gesteland, and T. R. Cech, Cold Spring Harbor Press, p. 293-307 (2010) cited as: Cold Spring Harb Perspect Biol doi: 10.1101/cshperspect.a003665.
202. "RNA Pseudoknots: Folding and Finding," B. Liu, D.H. Mathews, and D.H. Turner, *fl000 Biology Reports*, **2**, 8 (2010).

203. "Fluorescence Competition Assay Measurements of Free Energy Changes for RNA Pseudoknots," B. Liu, N. Shankar, and D.H. Turner, *Biochemistry*, 49, 623-634 (2010).
204. "NNDB: The Nearest Neighbor Parameter Database for Predicting Stability of Nucleic Acid Secondary Structure," D.H. Turner and D.H. Mathews, *Nucleic Acids Res.* 38, D280-D282 (2010).
205. "An RNA Molecular Switch: Intrinsic Flexibility of 23S rRNA Helices 40 and 68 5'-UAA/5'-GAN Internal Loops Studied by Molecular Dynamics Methods," K. Reblova, Z. Strelcova, P. Kulhanek, I. Besseova, D.H. Mathews, K. Van Nostrand, I. Yildirim, D.H. Turner, and J. Sponer, *J. Chem. Theory and Computation*, 6, 910-929 (2010).
206. "Re-parameterization of RNA χ Torsion Parameters for the AMBER Force Field and Comparison to NMR Spectra for Cytidine and Uridine," I. Yildirim, H.A. Stern, S.D. Kennedy, J.D. Tubbs, and D.H. Turner, *J. Chem. Theory and Computation*, 6, 1520-1531 (2010).
207. "RNA Internal Loops with Tandem AG Pairs: The Structure of the 5'GAGU/3'UGAG Loop can be Dramatically Different from Others, Including 5'AAGU/3'UGAA," N. B. Hammond, B. S. Tolbert, R. Kierzek, D. H. Turner, and S. D. Kennedy, *Biochemistry*, 49, 5817-5827 (2010).
208. "Comparisons between Chemical Mapping and Binding to Isoenergetic Oligonucleotide Microarrays Reveal Unexpected Patterns of Binding to the *Bacillus subtilis* RNase P RNA Specificity Domain," R. Liang, E. Kierzek, R. Kierzek, and D.H. Turner, *Biochemistry* 49, 8155-8168 (2010).
209. "Fluorescence Competition and Optical Melting Measurements of RNA Three-Way Multibranch Loops Provide a Revised Model for Thermodynamic Parameters," B. Liu, J.M. Diamond, D.H. Mathews, and D.H. Turner, *Biochemistry*, 50, 640-653 (2011).
210. "Identification of Potential Conserved RNA Secondary Structure throughout Influenza A Coding Regions," W. N. Moss, S. F. Priore, and D. H. Turner, *RNA*, 17, 991-1011 (2011).
211. "NMR Structure of a 4X4 Nucleotide RNA Internal Loop from an R2 Retrotransposon: Identification of a Three Purine-Purine Sheared Pair Motif and Comparison to MC-SYM Predictions," Y. Lerman, S. D. Kennedy, N. Shankar, M. Parisien, F. Major, and D. H. Turner, *RNA*, 17, 1664-1677 (2011).
212. "Biophysical analysis of influenza A virus RNA promoter at physiological temperatures," E. Noble, D. H. Mathews, J. L. Chen, D. H. Turner, T. Takamoto, and B. Kim, *J. Biol. Chem.*, 286, 24872-24881 (2011).
213. "Benchmarking AMBER Force Fields for RNA: Comparisons to NMR Spectra for Single-stranded r(GACC) are Improved by Revised χ Torsions," I. Yildirim, H. A. Stern, J. D. Tubbs, S. D. Kennedy, and D. H. Turner, *J. Phys. Chem. B*, 115, 9261-9270 (2011).
214. "The R2 Retrotransposon RNA Families," W. N. Moss, D. G. Eickbush, M. J. Lopez, T. H. Eickbush, and D. H. Turner, *RNA Biology*, 8, 714 - 718 (2011).
215. "Molecular Mechanics Investigation of an Adenine-Adenine Non-Canonical Pair Conformational Change," K. Van Nostrand, S. D. Kennedy, D. H. Turner, and D. H. Mathews, *J. Chem. Theory Computation*, 7, 3779 - 3792 (2011).

216. "Revision of AMBER Torsional Parameters for RNA Improves Free Energy Predictions of Tetramer Duplexes with GC and iGiC Base Pairs," I. Yildirim, S. D. Kennedy, H. A. Stern, J. M. Hart, R. Kierzek, and D. H. Turner, *J. Chem. Theory Computation*, 8, 172-181 (2012).
217. "Testing the Nearest Neighbor Model for Canonical RNA Base Pairs: Revision of GU Parameters," J. L. Chen, A. L. Dishler, S. D. Kennedy, I. Yildirim, B. Liu, D. H. Turner, and M. J. Serra, *Biochemistry* 51, 3508-3522 (2012).
218. "Influenza A Virus Coding Regions Exhibit Host-specific Global Ordered RNA Structure," S. F. Priore, W. N. Moss, and D. H. Turner, *PLoS ONE*, 7(4): e35989, doi:10.1371/journal.pone.0035989 (2012).
219. "The 3' Splice Site of Influenza A Segment 7 mRNA Can Exist in Two Conformations: A Pseudoknot and a Hairpin," W. N. Moss, L. I. Dela-Moss, E. Kierzek, R. Kierzek, S. F. Priore, and D. H. Turner, *PLoS ONE*, 7(6): e38323. Doi:10.1371/journal.pone.0038323. (2012).
220. "The Influenza A Segment 7 mRNA 3' Splice Site Pseudoknot/Hairpin Family," W. N. Moss, L. I. Dela-Moss, S. F. Priore, and D. H. Turner, *RNA Biology* 9, 1305-1310 (2012).
221. "Understanding the Role of Base Stacking in Nucleic Acids. MD and QM Analysis of Tandem GA Pairs in RNA Duplexes," C. A. Morgado, D. Svozil, D. H. Turner, and J. Sponer, *Physical Chemistry Chemical Physics*, 14, 12580-12591 (2012). Doi: 10.1039/c2cp40556c.
222. "Novel Conformation of an RNA Structural Switch," S. D. Kennedy, R. Kierzek, and D. H. Turner, *Biochemistry* 51, 9257-9259 (2012).
223. "The Nuclear Magnetic Resonance of CCCC RNA Reveals a Right-Handed Helix, and Revised Parameters for AMBER Force Field Torsions Improve Predictions from Molecular Dynamics," J. D. Tubbs, D. E. Condon, S. D. Kennedy, M. Hauser, P. C. Bevilacqua, and D. H. Turner, *Biochemistry* 52, 996-1010 (2013).
224. "Secondary Structure of a Conserved Domain in the Intron of Influenza A NS1 mRNA," S. F. Priore, E. Kierzek, R. Kierzek, J. R. Baman, W. N. Moss, L. I. Dela-Moss, and D. H. Turner, *PLoS ONE*, 8(9): e70615. Doi: 10.1371/journal.pone.0070615 (2013).
225. "Influenza B Virus has Global Ordered RNA Structure in (+) and (-) Strands but Relatively Less Stable Predicted RNA Folding Free Energy than Allowed by the Encoded Protein Sequence," S. F. Priore, W. N. Moss, and D. H. Turner, *BMC Research Notes* 6:330. Doi: 10.1186/1756-0500-6-330. (2013).
226. "Fundamental Interactions in RNA: Questions Answered and Remaining," D. H. Turner, *Biopolymers* 99, 1097-1104 (2013).
227. "The Contribution of Pseudouridine to Stabilities and Structure of RNAs," E. Kierzek, M. Malgowska, J. Lisowiec, D. H. Turner, Z. Gdaniec, and R. Kierzek, *Nucleic Acids Res.* 42, 3492-3501 (2014).
228. "Identification of Conserved RNA Secondary Structures at Influenza B and C Splice Sites Reveals Similarities and Differences between Influenza A, B, and C," L. I. Dela-Moss, W. N. Moss, and D. H. Turner, *BMC Research Notes*, 7:22. Doi:10.1186/1756-0500-7-22 (2014).
229. "Optimization of an AMBER Force Field for the Artificial Nucleic Acid, LNA, and Benchmarking with NMR of L(CAAU)," D. E. Condon, I. Yildirim, S. D. Kennedy, B. C. Mort, R. Kierzek, and D. H. Turner. *J. Phys. Chem. B.* 118. 1216-1228 (2014).

230. "Structure Determination of Noncanonical RNA Motifs Guided by NMR Chemical Shifts," P. Sripakdeevong, M. Cevec, A. T. Chang, M. C. Erat, M. Ziegler, Q. Zhao, G. E. Fox, X. Gao, S. D. Kennedy, R. Kierzek, E. P. Nikonowicz, H. Schwalbe, R. K. Sigel, D. H. Turner, and R. Das, *Nature Methods* 11, 413-416 (2014).
231. "Secondary Structure of a Conserved Domain in an Intron of Influenza A mRNA," T. Jiang, S. D. Kennedy, W. N. Moss, E. Kierzek, and D. H. Turner, *Biochemistry* 53, 5236-5248 (2014).
232. "Microarrays for Identifying Binding Sites and Probing Structure of RNAs," R. Kierzek, D. H. Turner, and E. Kierzek, *Nucleic Acids Res.* 43, 1-12 (2015).
233. "Stacking in RNA: NMR of Four Tetramers Benchmark Molecular Dynamics," D. E. Condon, S. D. Kennedy, B. C. Mort, R. Kierzek, I. Yildirim, and D. H. Turner, *J. Chem. Theory Comput.* 11, 2729-2742 (2015).
234. "Structural Features of a 3' Splice Site in Influenza A," J. L. Chen, S. D. Kennedy, and D. H. Turner, *Biochemistry* 54, 3269-3285 (2015).
235. "The Influenza A PB1-F2 and N40 Start Codons Are Contained within an RNA Pseudoknot," S. F. Priore, A. D. Kauffmann, J. R. Baman, and D. H. Turner, *Biochemistry* 54, 3413-3415 (2015).
236. "Nuclear Magnetic Resonance-Assisted Prediction of Secondary Structure for RNA: Incorporation of Direction-Dependent Chemical Shift Constraints," J. L. Chen, S. Bellaousov, J. D. Tubbs, S. D. Kennedy, M. J. Lopez, D. H. Mathews, and D. H. Turner, *Biochemistry* 54, 6769-6782 (2015).
237. "Self-folding of Naked Segment 8 Genomic RNA of Influenza A Virus," E. Lenartowicz, J. Keszy, A. Ruzkowska, M. Soszynska-Jozwiak, P. Michalak, W. N. Moss, D. H. Turner, R. Kierzek, and E. Kierzek, *PLoS ONE* 11:e0148281 (2016).
238. "Mutations Designed by Ensemble Defect to Misfold Conserved RNA Structures of Influenza A Segments 7 and 8 Affect Splicing and Attenuate Viral Replication in Cell Culture," T. Jiang, A. Nogales, S. F. Baker, L. Martinez-Sobrido, and D. H. Turner, *PLoS ONE* 11: e0156906. (2016).
239. "Antisense Oligonucleotides Targeting Influenza A Segment 8 Genomic RNA Inhibit Viral Replication," E. Lenartowicz, A. Nogales, E. Kierzek, R. Kierzek, L. Martinez-Sobrido, and D. H. Turner, *Nucleic Acid Therapeutics* 26, 277-285 (2016) (with cover illustration).
240. "Crystal Structure of a Poly(rA) Staggered Zipper at Acidic pH: Evidence that Adenine N1 Protonation Mediates Parallel Double Helix Formation," M. L. Gleghorn, J. Zhao, D. H. Turner, and L. E. Maquat, *Nucleic Acids Res.* 44, 8417-8424 (2016).
241. "RNA Structure Determination", D.H. Turner and D.H. Mathews, eds., Springer Humana Press (2016).
242. "RNA Secondary Structure Determination by NMR," J. L. Chen, S. Bellaousov, and D.H. Turner, *Meth. Molec. Biol.* 1490, 177- 186 (2016) [chap. 11 in *RNA Structure Determination*].
243. "RNA Secondary Structure Prediction," D. H. Mathews, D. H. Turner, and R. M. Watson, *Curr. Protoc. Nucleic Acid Chem.* 67, 11.2.1 – 11.2.19 (2016), doi: 10.1002/cpnc.19

244. "Predicting the Kinetics of RNA Oligonucleotides Using Markov State Models," G. Pinamonti, J. Zhao, D. E. Condon, F. Paul, F. Noe, D. H. Turner, and G. Bussi, *J. Chem. Theory Comput.* 13, 926-934 (2017).
245. "Nuclear Magnetic Resonance Structure of an 8X8 Nucleotide RNA Internal Loop Flanked on Each Side by Three Watson-Crick Pairs and Comparison to Three-Dimensional Predictions," A. D. Kauffmann, S. D. Kennedy, J. Zhao, and D. H. Turner, *Biochemistry*, 56, 3733-3744 (2017).
246. "Physics-based all-atom modeling of RNA energetics and structure," L.G. Smith, J. Zhao, D. H. Mathews, and D. H. Turner, *WIREs RNA*, 8e1422 (2017).
247. "Surprising Sequence Effects on GU Closure of Symmetric 2X2 Nucleotide Internal Loops," K. D. Berger, S. D. Kennedy, S. J. Schroeder, B. M. Znosko, H. Y. Sun, D. H. Mathews, and D. H. Turner, *Biochemistry* 57, 2121-2131 (2018).
248. "Conformational Ensembles of RNA Oligonucleotides from Integrating NMR and Molecular Simulations," S. Bottaro, G. Bussi, S. D. Kennedy, D. H. Turner, and K. Lindorff-Larsen, *Science Advances*, 4, eaar8521 (2018).
249. "Improving RNA Nearest Neighbor Parameters for Helices by Going Beyond the Two-State Model," A. Spasic, K. D. Berger, J. L. Chen, M. G. Seetin, D. H. Turner, and D. H. Mathews, *Nucleic Acids Res.* 46, 4883-4892 (2018).
250. "Molecular Dynamics Correctly Models the Unusual Major Conformation of the GAGU RNA Internal Loop and with NMR Reveals an Unusual Minor Conformation," A. Spasic, S. D. Kennedy, L. Needham, M. Manoharan, R. Kierzek, D. H. Turner and D. H. Mathews, *RNA* 24, 656-672 (2018).
251. "NMR Reveals that GU Base Pairs Flanking Internal Loops can Adopt Diverse Structures," K. D. Berger, S. D. Kennedy, and D. H. Turner, *Biochemistry* 58, 1094-1108 (2019).
252. "Accurate Geometrical Restraints for Watson-Crick Base Pairs," M. Gilski, J. Zhao, M. Kowiel, D. Brzezinski, D.H. Turner, and M. Jaskolski, *Acta Crystallographica Section B: Structural Science, Crystal Engineering and Materials* B75, 235-245 (2019).
253. "*In Vivo* Analysis of Influenza A mRNA Secondary Structures Identifies Critical Regulatory Motifs," L. M. Simon, E. Morandi, A. Luganini, G. Gribaudo, L. Martinez-Sobrido, D. H. Turner, S. Oliviero, and D. Incarnato, *Nucleic Acids Res.* 47, 7003-7017 (2019).
254. "Nuclear Magnetic Resonance of Single Stranded RNAs and DNAs of CAAU and UCAAUC as Benchmarks for Molecular Dynamics Simulations," J. Zhao, S.D. Kennedy, K.D. Berger, and D.H. Turner, *J. Chem. Theory Comput.* 16, 1968-1984 (2020).
255. "Nuclear Magnetic Resonance Spectra and AMBER OL3 and ROC-RNA Simulations of UCUCGU Reveal Force Field Strengths and Weaknesses for Single Stranded RNA," J. Zhao, S.D. Kennedy, and D.H. Turner, *J. Chem. Theory Comput.* 18, 1241-1254 (2022).
256. "Nuclear Magnetic Resonance Reveals a Two Hairpin Equilibrium near the 3'-Splice Site of Influenza A Segment 7 mRNA that can be Shifted by Oligonucleotides," A.D. Kauffmann, S.D. Kennedy, W.N. Moss, E. Kierzek, R. Kierzek, and D.H. Turner, *RNA* 28, 508-522. (2022).

257. "Nearest Neighbor Rules for RNA Helix Folding Thermodynamics: Improved End Effects," J. Zuber, S.J. Schroeder, H. Sun, D.H. Turner, and D.H. Mathews, *Nucleic Acids Res.* 50, 5251-5262 (2022).
258. "Norman Sutin, founding editor of *Comments in Inorganic Chemistry*: A Remembrance and Tribute," B.S. Brunschwig and D.H. Turner, *Comments Inorg. Chem.* (2022).

Research Support

American Chemical Society - Petroleum Research Fund, Type G Grant, "The Kinetics and Thermodynamics of Stacking Reactions", \$9,000, 1975-1977.

Research Corporation, "Fluorescence Detected Circular Dichroism Studies", \$16,790, 1975-1976.

National Science Foundation Department Equipment Grant, "Circular Dichroism Instrumentation", coauthored with T. R. Krugh, \$37,200, 1975-1976.

American Cancer Society Institutional Research Grant, "Sequence Specificity of Drug-Nucleic Acid Complexes", \$1,966, 1975-1976.

Alfred P. Sloan Fellowship, \$20,000, 1979-1983.

Biomedical Research Support Grant, "Laser Cross Linking of Nucleic Acids and Proteins, coauthored with D. C. Hinkle, \$5,000, 1979-1980.

National Institutes of Health, "Laser Cross Linking of Proteins and Nucleic Acids", \$79,765 (direct cost), 1980-1983.

National Institutes of Health, "Fluorescence Detected CD Studies of Nucleic Acids", \$81,006 (direct cost), 1981-1984.

Biomedical Research Support Grant, "Temperature Control for Absorption Spectroscopy", \$4,040, 1984-1985.

National Institutes of Health (National Research Service Award for Senior Fellowship), "Studies of Self Splicing RNA", \$30,140, 1984-1985.

National Institutes of Health, "Effects of Drugs on Nucleic Acids", \$94,346 (direct cost), 1984-1987.

National Institutes of Health, "500 MHz NMR Spectrometer" (DRR BRS Shared Instrumentation), \$300,000 (equipment only), Co-P.I. with T. R. Krugh, 1987.

Office of Naval Research, "Molecular Basis of RNA Catalysis," \$155,337 (direct cost), 1988-1991.

National Institutes of Health, "Computer Graphics Equipment" (DRR BRS Shared Instrumentation), \$172,000 (equipment only).

National Institutes of Health, GM22939, "Kinetic and Spectroscopic Studies of Nucleic Acids",

\$128,415 (direct cost), 1976-1979.

\$423,255 (direct cost), 1982-1987.

\$154,759 (direct cost), 1979-1982.

\$691,574 (direct cost), 1987-1992.

National Institutes of Health, GM22939, "Folding RNA: from AIDS to the Human Genome".

\$753,829 (direct cost), 1992-1996 (funded with priority score in top 2%)

\$876,545 (direct cost), 1996-2000, (funded with priority score in top 1%)

\$1,059,037 (direct cost), 2000-2004 (funded with priority score in top 6%)

\$1,211,970 (direct cost), 2004-2008 (funded with priority score in top 7%)

\$1,252,046 (direct cost), 2008-2013 (funded with priority score in top 10%)

\$ 110,000 (direct cost), 2012-2013 (funded supplement)

\$2,356,955 (total cost). 2013-2017 (funded with priority score in top 16%)

National Institutes of Health, 1 R03 TW 01068-01, "Folding RNA with Modified Oligonucleotides," (Fogarty International Research Collaboration Award to fund Dr. Ryszard Kierzek's synthesis program at the Polish Academy of Sciences, Institute of Bioorganic Chemistry in Poznan, Poland) \$82,285 (direct cost), 1999-2002, \$96,000 (direct costs), 2002-2005.

National Institutes of Health, 1 R03 AI 45398-01, "RNA Targeting with Tertiary Interactions," (Innovative Drug Discovery Research in AIDS Opportunistic Infections) \$100,000, 1999-2001.

National Institutes of Health, 1 R03 TW008739, "Folding RNA: Influenza," (Fogarty International Research Collaboration Award to fund Dr. Elzbieta Kierzek (Institute of Bioorganic Chemistry in Poznan, Poland) to work on determining structures of domains in influenza RNA and to discover oligonucleotides that could potentially serve as therapeutics. \$174, 509 (direct cost), 2011-2014 (priority score in top 2%).

Research Students

[“Position” is either last prior to retirement or current. Subsequent Positions are last known.]

Research Associates

Susan M. Freier

Ph.D., University of California, Berkeley, 1976

Position: Director of Molecular and Structural Biology at IONIS Pharmaceuticals, Carlsbad, CA.
Retired.

Naoki Sugimoto

Ph.D., Kyoto University, 1985

Positions: Professor of Chemistry, Konan University, Kobe, JAPAN

Director, Frontier Institute of Biomolecular Engineering Research (FIBER)

Sean Moran

Ph.D., University of Colorado, Boulder, 1987

Position: Director, Structural Biology Core Facility, Uniform Services University of the Health Sciences

Matthew Fountain

Ph.D., University of Rochester, 1994

Position: Professor and Chair of Chemistry, State University College at Fredonia NY

Stephen M. Testa (NIH Postdoctoral Fellow)

Ph.D., Purdue University, 1994

Position: Associate Professor of Chemistry, University of Kentucky

Sherry Spinelli

Ph.D., University of Rochester, 1999

Position: Research Associate Professor of Pathology and Laboratory Medicine, University of Rochester Medical Center

Irina Catrina

Ph.D., Utah State University, 2001

Position: Clinical Assistant Professor of Chemistry, Yeshiva University

Elzbieta Kierzek

Ph.D., Institute for Bioorganic Chemistry, Poznan, Polish Academy of Sciences

Position: Professor, Institute of Bioorganic Chemistry, Poznan

Elzbieta Lenartowicz

Ph.D., Institute for Bioorganic Chemistry, Poznan, Polish Academy of Sciences, 2016

Position: Head of Laboratory of Molecular Diagnostics, Institute for Bioorganic Chemistry, Poznan, Polish Academy of Sciences

Ph.D. Students

1979 T. Gregory Dewey

“Laser Temperature Jump Kinetics of Solvent Effects on Stacking Reactions in Dyes and Nucleic Acids”

Position: President, Albany College of Pharmacy and Health Sciences

- 1980 Craig D. Scoville
“Studies on a Monoclonal IgG Cryoglobulin”, joint student with G. N. Abraham
(Department of Microbiology)
Position: Practicing Physician
- 1981 Eric W. Lobenstine
“Fluorescence Detected Circular Dichroism: Verification and Applications to Proteins”
Position: Manager of Computers and Network, Chemistry Department, University of Rochester
- 1981 Diane DePrisco Albergo
“Solvent Effects on the Thermodynamics and Kinetics of Double Helix Formation in Nucleic Acids”
Position: Scientist at Abbott Laboratories, Irving, Texas
- 1982 Matthew Petersheim (deceased)
“Stabilization of Ribonucleic Acid Double Helices by 'Dangling' Bases: Optical and NMR Studies”
Position: Associate Professor and Chair of Chemistry, Seton Hall University, South Orange, NJ
- 1984 Catherine A. Harrison
“Photochemistry of Nucleic Acids Including Crosslinking to Proteins”
Subsequent position: Scientist at Eastman Kodak Company, Rochester, NY
- David R. Hickey
“Thermodynamic Effects of Solvent and Terminal Mismatches on Oligoribonucleotide Stability”
Position: Instructor, Department of Chemistry, University of Rochester
- 1985 Michael Lamos
“Fluorescence Detected Circular Dichroism of Ethidium-Nucleic Acid Complexes”
Subsequent Position: Scientist at Sienna Biotech, Stillwater, MN
- 1988 Alison Williams
“Raman Laser Temperature-Jump, Spectroscopic and Thermodynamic Studies of Salt Effects on Duplex Formation by dGCATGC”
Position: Vice President for Equity and Inclusion, Wesleyan University
- 1989 Carl Longfellow
“Thermodynamic and Spectroscopic Studies of Bulge Loop Structures Formed by Oligoribonucleotides”
Subsequent Position: Scientist at Wyeth-Ayerst Labs, Rouses Point, NY
- 1989 John Jaeger
“RNA Secondary Structure: Prediction, Melting, and Chemical Modification”
Subsequent Position: Information Technology Business Partner for Research, Bristol Myers Squibb
- 1991 John SantaLucia, Jr.
“The Role of Hydrogen Bonding in the Thermodynamics and Structure of Mismatches in RNA Oligonucleotides”

Positions:

Professor of Chemistry, Wayne State University, Detroit, MI
CEO, DNA Software, Ann Arbor, MI

1993 Adam E. Peritz

“Studies of Symmetric and Asymmetric Internal Loops in RNA”

Subsequent position: Staff member, Nucleic Acid Facility, University of Pennsylvania

Philip C. Bevilacqua

“Dynamics of Substrate-Ribozyme Interaction: Binding, Conformational Changes, and Catalysis”

Position: Professor of Chemistry (Department Chairman), Pennsylvania State University, University Park, PA

Liyan He

“Thermodynamic and Structural Studies of G•U Mismatches in RNA”

Subsequent position: Sr. Technology Commercialization and Liaison Officer at Weill Cornell Medical College, New York City

1994 Amy E. Walter (Morsch)

“Thermodynamic Studies of the Sequence Dependence of Stability for Interfaces of Coaxially Stacked Helices and Tandem GA Mismatches in RNA”

Subsequent position: Programmer, A-LIFE Medical, Inc., San Diego, CA

Aloke Raj Banerjee

“Investigation of the Folding Pathway of a Group I Ribozyme: Melting, Chemical Modification and Kinetics Studies”

Subsequent position: Manager, Assay Research, Nanosphere

1996 Ming Wu

“Structure and Thermodynamic Stability of Symmetric Tandem Mismatches in RNA”

Subsequent position: Manager, Gilead Sciences Inc., Foster City, CA

Jeffrey A. McDowell

“Investigation of the Structural Basis of the Thermodynamic Stability of Symmetric, Tandem GU Mismatches in RNA”

Position: Senior Manager – The Janssen Pharmaceutical Companies of Johnson & Johnson, Greater Philadelphia Area

1997 James Kim

“The Thermodynamics of Coaxial Stacking and Its Effect on RNA Secondary Structure”

Position: Associate Professor of Internal Medicine, University of Texas Southwestern Medical School

Yi Li

“Effects of Temperature, Mg²⁺ and the 2' OH of Guanosine on Substrate Binding and Reactivity with the Tetrahymena Ribozyme”

Subsequent Position: Scientist, Alza Pharmaceuticals, Mountain View, CA

1998 Louis Profenno

“Conformational Rearrangements in the Catalytic Mechanism of the *Tetrahymena* Ribozyme”

Position: Psychiatrist, Syracuse, NY

- 1999 Tianbing Xia
“Sequence Dependence of Stabilities and Structures of Tandem Mismatches and Watson-Crick Base Pairs in RNA”
Position: Systems Engineer, Abbott Labs, Dallas
- 2000 Xiaoying Chen
“Stability and Structure of Guanosine-Uridine and Isoguanosine-Isocytidine Pairs”
Subsequent position: Manager, Bioinformatics, Roche Molecular Systems, Inc., Pleasanton, CA
- Mark E. Burkard
“Base Stacking Interactions and the Thermodynamics and Structure of Guanine-Guanine Pairs in RNA”
Position: Professor of Medicine and Oncology, University of Wisconsin, Madison
- 2001 David H. Mathews
“Prediction of RNA Secondary Structure”
Position: Maquat Distinguished Professor of RNA Biology, Department of Biochemistry and Biophysics, University of Rochester
- Thomas W. Barnes III
“The Impact of Long-Range Cooperativity, Base Composition and Structure on the Molecular Recognition of RNA by C5-(1-Propynyl) Pyrimidine-Containing Oligodeoxynucleotides”
Subsequent position: Patent Attorney, International Paper Co.
- 2002 Susan J. Schroeder
“Exploring the Thermodynamic Stabilities and Structures of Asymmetric Internal Loops in RNA”
Positions: Associate Professor, Departments of Chemistry and of Microbiology and Plant Biology, University of Oklahoma
- Matthew D. Disney
“*In vitro* and *in vivo* Targeting of RNA in Fungal Pathogens with Oligonucleotides and Small-Molecules”
Position: Professor of Chemistry, Scripps Research Institute - Florida
- 2003 Jessica Childs (Disney)
“Oligonucleotide Directed Misfolding of RNAs”
Position: Senior Staff Scientist, Chemistry Dept., Scripps Research Institute - Florida
- 2004 Brent Znosko
“Thermodynamic and Structural Properties of RNA Oligomers Containing Purine Rich Internal Loops and DNA:RNA Hybrids with C5-Propynyl Substitutions”
Position: Professor of Chemistry, St. Louis University
- 2005 Gang Chen
“Expanded Sequence Dependence Improves Prediction of RNA Internal Loop Stability and NMR Reveals Molecular Recognition Interactions Accounting for Stability, Structure and Dynamics”
Position: Associate Professor, School of Life and Health Sciences, The Chinese University of Hong Kong, Shenzhen

- 2006 Shenghua Duan
“Applying Oligonucleotide Microarrays in Determining RNA Secondary Structure”
Subsequent position: Bioinformatics Analyst II, Center for Cancer Computational Biology, Dana Farber Cancer Institute
- Blanton Tolbert
“The Solution NMR Structures of Model RNA Duplexes Containing Non-Canonical GA Pairs: Insights into the Factors Affecting Thermodynamic Stability of RNA 2X2 Nucleotide Internal Loops”
Position: Professor of Chemistry, Case Western Reserve University
- 2007 Neelaabh Shankar
“NMR Studies of Two Conserved RNA Internal Loops Found in Ribosomes”
Position: Patent Attorney
- 2008 Ilyas Yildirim
“Free Energy Calculations of RNA Duplexes with Tandem GA Base Pairs Using the Thermodynamic Integration Approach”
Position: Assistant Professor of Chemistry, Florida Atlantic University
- 2009 James M. Hart
“NMR-Assisted Prediction of RNA Secondary Structure”
Position: Medical Director at Abbott Core Diagnostics
- 2010 Ruiting Liang
“RNA Secondary Structure Determination: Isoenergetic Oligonucleotide Microarray Compared with Chemical Mapping and MALDI MS Detection of Chemical Mapping”
Position: Director, Analytical Research and Development, Beigene
- Biao Liu
“Fluorescence Competition Assay Measurements of Thermodynamics for RNA Pseudoknots and Multibranch Loops”
Position: Computational Scientist, Intellia Therapeutics, Cambridge MA
- 2011 Walter N. Moss
“RNA Secondary Structure Discovery and Characterization”
Position: Associate Professor of Molecular Biology and Biophysics, Iowa State University
- Nicolas Hammond
“Investigating the mechanism of Hoechst 33258 inhibition of *Candida* spp. growth and II. RNA internal loops with tandem AG pairs: the structure of the 5’GAGU/3’UGAG loop can be dramatically different from others, including 5’AAGU/3’UGAA”
Position: Assistant Director of University of Rochester Workshop Program
- 2013 Jason Tubbs
“Computational and Experimental Advances in the RNA Therapeutic Pipeline”
Position: Associate Director of Quality Operations, CoreRX, Clearwater, FL
- Salvatore Priore
“Discovery and Characterization of Influenza Virus RNA Secondary Structures”
Position: Assistant Professor of Clinical Pathology and Laboratory Medicine, Perelman School of Medicine, University of Pennsylvania

- Indee Dela-Moss
“Analysis of RNA Secondary Structure: Common Themes in Spliced Segments of Influenza A, B, and C”
Subsequent Position: Postdoctoral Fellow, Iowa State University
- 2015 David Condon
“Nucleic Acid Force Fields in Prediction of Ensemble NMR Properties”
Position: Assistant Professor of Internal Medicine, University of South Dakota, Lead Computational Bioinformatics Analyst at Sanford Health, Sioux Falls, South Dakota
- Jonathan Chen
“Two- and Three-Dimensional Modeling of RNA Structures with NMR and Thermodynamics Methods”
Position: Staff Scientist, Center for RNA Biology, University of Rochester Medical Center
- Tian Jiang
“RNA Structure and Function of Influenza Virus”
Position: Systems Engineer, Abbott Labs, Dallas
- 2018 Kyle Berger
“Thermodynamic and Structural Studies of RNA Internal Loops Closed by GU Pairs”
Position: Postdoctoral Fellow in laboratory of David MacLean, University of Rochester School of Medicine and Dentistry
- 2019 Andrew Kauffmann
“Structures of RNA Sequences from Influenza”
Position: Assistant Professor of Chemistry, Truman State University, Missouri
- 2020 Jianbo Zhao
“Molecular Dynamics and Quantum Mechanics Studies of RNA Structures”
Position: Senior Scientist, Nested Therapeutics, Boston, MA

Visiting Scientists

- Dr. Ryszard Kierzek, Professor, Institute of Bioorganic Chemistry, Poznan, Poland
- Dr. Elzbieta Kierzek, Professor, Institute of Bioorganic Chemistry, Poznan, Poland
- Dr. Peter Müller, Scientist, Dr. Karl Thomae, GmbH, Bieberach, Germany
Head, Department of Chemical Research, Dr. Karl Thomae, GmbH, Bieberach, Germany
Senior Vice President, Research and Development, Boehringer Ingelheim Pharmaceuticals, Ridgefield, CT
Chief Scientific Officer & Senior Vice President, Vertex Pharmaceuticals
Retired
- Dr. Martin J. Serra, Paul E. and Mildred L. Hill Professor of Chemistry, Allegheny College
Retired
- Dr. Janet Morrow, Professor of Chemistry, SUNY at Buffalo
- Dr. Neena Grover, Professor of Chemistry & Biochemistry, Colorado College