

Charles H. Reynolds

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Experience

2011-present Founder, Gfree Bio, LLC

- Gfree Bio develops and delivers modeling, informatics, and structure-based design capabilities for emerging and established biotech companies. Gfree Bio has helped advance drug discovery programs for more than 60 companies since its founding.
- Critical expert witness for a successful patent settlement in excess of \$1 billion.
- Principal Investigator for NIH STTR program (2016-2017) at Gfree Bio to develop new methodology for refining protein-ligand cocrystal structures.
- Acting CSO, PPI Mimics, LLC (2013-2015), and PI for two NIH grants.
- Other collaborations include: NIH SBIR The Wistar Institute; multiple NIH SBIRs Phelix Therapeutics; NIH STTR Fox Chase Chemical Diversity and Drexel College of Medicine, NIH SBIR for CerSci Therapeutics (acquired by Acadia).
- Co-inventor of phase III clinical candidate Resmetirom (Madriral Pharmaceuticals).

2010-2011 Senior Director and Head of Discovery Technologies, Ansaris

- Responsible for scientific computing, proprietary discovery technology, structural biology, and IT. Member of senior management team with significant business development responsibilities.
- Initiated a major update of core computational fragment-based drug design platform. Refocused technology to target small molecule ligands for protein-protein interactions. Updated and improved software (speed, efficiency, user-interface).
- Leadership role in identification and prosecution of an internal target that led to a significant pharma partnering opportunity.
- Executive management of major research collaboration with Novartis.

2000-2010 Research Fellow, Johnson & Johnson Pharmaceutical R&D

- Responsibilities included managing modeling and informatics research teams at the Spring House, PA and La Jolla, CA sites. Managed significant hardware and software budgets across multiple US sites.
- Teams under my direction provided critical support for progression of tens of compounds to early development, multiple compounds to clinical development. Contributed to carisbamate NDA and European registration for Dacogen®.
- Led development of new discovery methodology at J&J including: informatics tools for fragment-based design, novel protocols for improved crystal-structure refinement, linear interaction energy approaches for ligand optimization, a similarity-based tool for scoring reagents for “drug-likeness,” application of linear-scaling quantum calculations to analysis of protein structures, and structure-based ADME-Tox. The latter effort

includes a published multi-state homology model for the hERG channel that was employed successfully in multiple discovery programs and was honored in 2008 as a most cited article by Elsevier.

- Migrated J&J Computer-Aided Drug Discovery (CADD) groups to linux-based clusters and workstations, and consolidated CADD software licenses across sites. This effort resulted in a significant improvement in computational capabilities while reducing the overall cost of high-performance computing.
- Member of the New Targets Evaluation Committee for drug discovery. Organized a series of global J&J CADD and structural-biology meetings. Initiated external research collaborations at Penn State, Yale, and the University of Florida. Coordinated J&J's first structural-biology collaboration with an outside crystallography CRO.

1995-2000 Research Fellow, **1984-1994** Senior Scientist, Rohm and Haas (DowDuPont)

- Co-inventor of a new class of ergosterol biosynthesis inhibitor antifungals. Awarded the highest scientific award in the company (Otto Haas Award).
- Led an interdisciplinary team of a dozen scientists tasked with developing new drug therapies and diagnostic aids. This project resulted in a novel nanoparticle-based targeted contrast agent for MR imaging. Roles included developing the concept and selling it to R&D management, leading the team, establishing a collaboration with the University of Pennsylvania Medical School, and developing a market analysis in collaboration with Argonex (a biotechnology company acquired by Millipore).

2019-present Editorial Advisory Board, Journal Chemical Information and Modeling

2014-present Editorial Advisory Board, ACS Medicinal Chemistry Letters

2009-2013 Director (Industry Advisory Board) QuantumBio Inc.

2013 J.T. Oden Visiting Faculty Fellow, ICES, The University of Texas at Austin

2010-2017 Adjunct Professor, Pennsylvania Drug Discovery Institute

2008-2018 Member, American Chemical Society Publications Committee

2007-2016 NIH study sections, BDMA, BCMB, IMST-10, IMST-11, NCI ZCA1

2006-2011 Scientific Advisory Board, Keystone Symposia

2006-2009 Councilor, American Chemical Society

1998-2006 Editorial Advisory Board, Journal of Molecular Graphics and Modeling

1998 Joint postdoctoral research supervisor, University of North Carolina at Chapel Hill

1996-1998 Joint graduate research supervisor, The Pennsylvania State University

1990 Adjunct Professor, The University of Pennsylvania

Additional Experience and Leadership Activities

2015 National Academy of Sciences STTR program workshop; **2008** Co-organizer, first Keystone Symposium on computer-aided drug discovery; **2006-2009** Advisory Board for NSF sponsored science and math education initiative at Temple University (TU-SMART); **2007** Co-organizer, ACS Prospectives Meeting, Advances in Structure-Based Drug Discovery; **2007** Inaugural Keck Foundation Lecture, St. Edward's University; **2007** Lecturer, ACS computational chemistry short course; **2005** Organizer, ACS Prospectives Meeting, Advances in Structure-Based Drug Discovery; **1999-2000** Rohm and Haas Technology Center Coordinator for Life Sciences; **1998-2000** Membership Committee, ACS Computers in Chemistry Division; **1999** Co-taught a computational chemistry short course (Jan Term) at Austin College; **1997** Chair, ACS Computers in Chemistry Division; **1991-1995** Treasurer, ACS Computers in Chemistry Division; **1989** Faculty, QCPE course.

Education

1984 Ph.D. in Theoretical Organic Chemistry, The University of Texas at Austin, Austin, Texas. Supervisor: Michael J.S. Dewar.
1979 B.A. in Chemistry, Austin College, Sherman, Texas. Honors in Chemistry.

Management Training

2013 Kauffman Institute Entrepreneur Training Program
2002 Johnson & Johnson Management Fundamentals
2001 Johnson & Johnson Manager and the Law
1993 First-Line Leadership
1992 The Leadership Challenge

Awards

2015 Gfree Bio selected as a Kauffman Foundation FastTrac success story
2012 Distinguished Alumni Award, Austin College
2010 Fellow of the American Chemical Society
2007 Standards of Leadership Award, Johnson & Johnson
2007 Platinum Encore Award, Johnson & Johnson (Dacogen registration in Europe)
2000 Johnson & Johnson Excellence in Science Award (four awarded company wide)
1987 Otto Haas Award, Rohm and Haas (highest award for scientific achievement).
1993 Marion Cox Award, Austin College; **1984** University of Texas Career Development Award; **1983** Robert A. Welch Foundation Pre-doctoral Fellowship, The University of Texas at Austin; **1978** P.S. Wharton Fellowship in chemistry, Austin College; **1978** National Science Foundation Undergraduate Research Scholarship, The University of Texas at Arlington; **1975-1979** Robert A. Welch Undergraduate Chemistry Scholarship.

Publications

Author of over 100 papers, chapters, books, and patents

Publications

Bratkowski, M.; Burdett, T.C.; Danao, J.; Wang, X.; Mathur, P.; Gu, W.; Beckstead, J.; Talreja, S.; Yang, Y.-S.; Danko, G.; Park, J.H.; Walton, M.; Brown, S.P.; Tegley, C.M.; Joseph, P.R.B.; Reynolds, C.H.; Sambashivan, S. **Uncompetitive, adduct-forming SARM1 inhibitors are neuroprotective in Preclinical models of nerve injury and disease**, *Neuron*, 2022, <https://doi.org/10.1016/j.neuron.2022.08.017>

Carta, L.; Hutcheson, R.; Davis, S.A.; Rudolph, M.J.; Reynolds, C.H.; Quick, M.; Williams, T.M.; Schmertzler, M.; Hadari, Y.R. **Developing small molecules that inhibit K-Ras/GTP binding based on new affinity measurements**, *bioRxiv*, 2020, doi: <https://doi.org/10.1101/2020.07.27.218248>

Agragiotis, D.K.; Holloway, M.K.; Johnson, S.A.; Reynolds, C.H.; Stouch, T.R.; Tropsha, A.; Waller, C.L. **Chemistry, information, and Frank: a tribute to Frank Brown**, *J. Computer-Aided Molecular Design*, 2018, 32, 723-729. <https://doi.org/10.1007/s10822-018-0135-9>

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Reynolds, C.H.; Reynolds, R.C. **Group additivity in ligand binding affinity: An alternative approach to ligand efficiency** *J. Chem. Inf. Model*, 2017, 57, 3086-3093 DOI: 10.1021/acs.jcim.7b00381

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Murray, Christopher W., Daniel A. Erlanson, Andrew L. Hopkins, György M. Keserü, Paul D. Leeson, David C. Rees, Charles H. Reynolds, and Nicola J. Richmond. **Validity of Ligand Efficiency Metrics**. *ACS Medicinal Chemistry Letters* (2014) 5, 2-5. (One of the most read papers in *ACS Medicinal Chemistry Letters* 2014)

Martha J. Kelly, Charles H. Reynolds, et al. **Discovery of 2-[3,5-Dichloro-4-(5-isopropyl-6-oxo-1,6-dihydropyridazin-3-yloxy)phenyl]-3,5-dioxo-2,3,4,5-tetrahydro[1,2,4]triazine-6-carbonitrile (MGL-3196), a Highly Selective Thyroid Hormone Receptor β Agonist in Clinical Trials for the Treatment of Dyslipidemia**, *Journal of Medicinal Chemistry*, 2014, 57, 3912-3923. DOI: 10.1021/jm4019299

Hopkins, A.L.; Keseru, G.M.; Leeson, P.D.; Rees, D.C.; Reynolds, C.H. **The role of ligand efficiency measures in drug discovery**, *Nature Reviews Drug Discovery*, 2014, 13, 105-121.

Reynolds, Charles, H., **Impact of computational structure-based methods on drug discovery**, *Current Pharmaceutical Design*, 2014, 20, 3380-3386. DOI: 10.2174/138161282020140528105532

Reynolds, Charles H.; Holloway, M. Katharine, **Thermodynamics of ligand binding and efficiency**, *ACS Medicinal Chemistry Letters*, (2011) 2, 433-437. *One of the top ten accessed papers in ACS Medicinal Chemistry Letters and the Journal of Medicinal Chemistry for 2011 and 2012.*

Céline Schalk-Hihi, Carsten Schubert, Richard Alexander, Shariff Bayoumy, Jose C. Clemente, Ingrid Deckman, Renee L. DesJarlais, Keli C. Dzordzorme, Christopher M. Flores, Bruce Grasberger, James K. Kranz, Frank Lewandowski, Li Liu, Hongchang Ma, Diane Maguire, Mark J. Macielag, Mark E. McDonnell, Tara Mezzasalma Haarlander, Robyn Miller, Cindy Milligan, Charles Reynolds, Lawrence C. Kuo, **Crystal structure of a soluble form of human monoglyceride lipase in complex with an inhibitor at 1.35 Å resolution** *Protein Science* (April 2011) 20, 670-683.

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Zhang, Xiaohua; Gibbs, Alan C.; Reynolds, Charles H.; Peters, Martin B.; Westerhoff, Lance M. **Quantum Mechanical Pairwise Decomposition Analysis of Protein Kinase B Inhibitors: Validating a New Tool for Guiding Drug Discovery**. J. Chem. Inf. Model. (2010), 50, 651-661.

Lawson, Edward C.; Luci, Diane K.; Ghosh, Shyamali; Kinney, William A.; Reynolds, Charles H.; Qi, Jenson; Smith, Charles E.; Wang, Yuanping; Minor, Lisa K.; Haertlein, Barbara J.; Parry, Tom J.; Damiano, Bruce P.; Maryanoff, Bruce E. **Nonpeptide Urotensin-II Receptor Antagonists: A New Ligand Class Based on Piperazino-Phthalimide and Piperazino-Isoindolinone Subunits**. Journal of Medicinal Chemistry (2009), 52, 7432-7445.

Li, Jian; Reynolds, C. H. **A quantum mechanical approach to ligand binding: Calculation of ligand-protein binding affinities for stromelysin-1 (MMP-3) inhibitors**, Canadian Journal of Chemistry, (2009), 87, 1480-1484.

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Baxter, Ellen W.; Conway, Kelly A.; Kennis, Ludo; Bischoff, Francois; Mercken, Marc H.; De Winter, Hans L.; Reynolds, Charles H.; Tounge, Brett A.; Luo, Chi; Scott, Malcolm K.; Huang, Yifang; Braeken, Mirielle; Pieters, Serge M. A.; Berthelot, Didier J. C.; Masure, Stefan; Bruinzeel, Wouter D.; Jordan, Alfonso D.; Parker, Michael H.; Boyd, Robert E.; Qu, Junya; Alexander, Richard S.; Brenneman, Douglas E.; Reitz, Allen B. **2-Amino-3,4-dihydroquinazolines as inhibitors of BACE-1 (β -site APP cleaving enzyme): use of structure based design to convert a micromolar hit into a nanomolar lead**. Journal of Medicinal Chemistry (2007), 50(18), 4261-4264. *One of the most accessed articles in JMC for 2007.*

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Reynolds, Charles H.; Tropsha, Alexander; Pfahler, Lori B.; Druker, Ross; Chakravorty, Subhas; Ethiraj, G.; Zheng, Weifan. **Diversity and coverage of structural sublibraries selected using the SAGE and SCA algorithms.** Journal of Chemical Information and Computer Sciences (2001), 41(6), 1470-1477.

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Reynolds, Charles H. **Semiempirical MO methods: the middle ground in molecular modeling.** THEOCHEM (1997), 401(3), 267-277.

Best, Scott A.; Merz, Kenneth M., Jr.; Reynolds, Charles H.. **GB/SA-Based Continuum Solvation Model for Octanol.** Journal of Physical Chemistry B (1997), 101(49), 10479-10487.

Reynolds, Charles H.. **Molecular modeling in agrochemicals.** Chemistry & Industry (London) (1997), (15), 592-595.

Reynolds, Charles H.; Hormann, Robert E. **Theoretical Study of the Structure and Rotational Flexibility of Diacylhydrazines: Implications for the Structure of Nonsteroidal Ecdysone Agonists**

and Azapeptides. Journal of the American Chemical Society (1996), 118(39), 9395-9401.

Reynolds, Charles H.. **Estimating Lipophilicity Using the GB/SA Continuum Solvation Model: A Direct Method for Computing Partition Coefficients.** Journal of Chemical Information and Computer Sciences (1995), 35(4), 738-42.

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Reynolds, Charles H.. **Structure and relative stability of halogenated carbocations: the C₂H₄X⁺ and C₄H₈X⁺ (X = fluoro, chloro, bromo) cations.** Journal of the American Chemical Society (1992), 114(22), 8676-82.

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Dewar, Michael J. S.; Ford, George P.; Reynolds, Charles H.. **Ground states of molecules. 61. Relative stabilities of o-, m-, and p-benzyne.** Journal of the American Chemical Society (1983), 105(10), 3162-7.

Dewar, Michael J. S.; Reynolds, Charles H. **Tritium migration in tritiated anisole.** Journal of the American Chemical Society (1982), 104(11), 3244-6.

Patents

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