

ANNE SKAJA ROBINSON

Chair of Chemical and Biomolecular Engineering
Catherine and Henry Boh Professor in Engineering
Tulane University, 300 Lindy Boggs Laboratory
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EDUCATION

- 1994 **Ph.D., Chemical Engineering** University of Illinois at Urbana-Champaign
Dissertation: *“Engineering the Yeast Secretory Pathway: The role of BiP and PDI in the secretion of foreign proteins in Saccharomyces cerevisiae.”*
Advisors: Prof. Douglas A. Lauffenburger, Prof. K. Dane Wittrup
- 1989 **M.S., Chemical Engineering** The Johns Hopkins University
Thesis: *“Isolation and Characterization of Proteolytic Enzymes from the Hyperthermophilic Archaeobacterium Pyrococcus furiosus.”*
Advisor: Prof. Robert M. Kelly
- 1988 **B.S., Chemical Engineering** The Johns Hopkins University
Departmental Honors GPA 3.9/4.0

PROFESSIONAL EXPERIENCE

- 2012-present **Chair, Chemical and Biomolecular Engineering and Catherine and Henry Boh Professor in Engineering**
Tulane University, School of Science and Engineering
- 2012-present **Adjunct Professor**
University of Delaware, Department of Chemical and Biomolecular Engineering
- 2008-2011 **Associate Chair for Biochemical Engineering**
University of Delaware, Department of Chemical Engineering
- 2008-2011 **Full Professor**
University of Delaware, Department of Chemical Engineering
- 2003-2008 **Associate Professor**
University of Delaware, Department of Chemical Engineering
- 1997-2003 **Assistant Professor**
University of Delaware, Department of Chemical Engineering
- 1994-1997 **Postdoctoral Fellow**
Massachusetts Institute of Technology, Department of Biology
Advised by Professor Jonathan King.

ACADEMIC HONORS

- 2015 Perlman Awardee, American Chemical Society (ACS) BIOT division
- 2013 American Institute of Chemical Engineers (AIChE) Women's Initiatives Committee's (WIC) Mentorship Excellence Award
- 2011 Fellow, American Institute for Medical and Biological Engineering

2002	Outstanding Junior Faculty Member, College of Engineering, University of Delaware
2000-2005	NSF Presidential Early Career Award in Science and Engineering (PECASE/Career)
2000-2003	DuPont Young Professor
2000	National Academy of Engineering, Sixth Annual Frontiers in Engineering
1996-97	NIH Postdoctoral Fellowship
1992-94	Clare Booth Luce Graduate Fellowship
1989-92	Department of Defense Fellowship (NDSEG)
1988-89	NSF Creativity Award for Scientists and Engineers
1988	National Science Foundation Fellowship awarded (declined)
1988-89	Tau Beta Pi Fellowship
1984-88	Beneficial Hodson Scholarship, Johns Hopkins University

PATENTS

- A) Kelly, R.M., A.K.S. Robinson, I.I. Blumentals, S.H. Brown, and C.B. Anfinsen. "Proteolytic Enzymes from Hyperthermophilic Bacteria and Processes for Their Production." Patent # 5,242,817. Filed 9/12/89. Accepted 9/7/93. Licensed to Takara Shuzo.
- B) Robinson, A.S. and K.D. Wittrup. "Methods for Increasing Secretion of Overexpressed Proteins." Patent # 5,773,245. Filed 10/92. Accepted 6/30/98.
- C) Robinson, A.S., D. Foguel, J.L. Silva, C.R. Robinson. "Use of Hydrostatic Pressure to Inhibit and Reverse Protein Aggregation and Facilitate Protein Refolding." Patent applied for, 60/161,035. Filed 10/99. Issued 11/10/09.

BOOK PUBLICATION

- 1) Russell, T.W.F., Robinson, A.S., and Wagner, N.J., (2008) Mass and Heat Transfer: Analysis of Mass Contactors and Heat Exchangers, Cambridge University Press, Cambridge, UK (www.cambridge.org/9780521886703).
- 2) Robinson, A.S., ed (2011) Production of Membrane Proteins – Strategies for Expression and Isolation, Wiley VCH (<http://onlinelibrary.wiley.com/book/10.1002/9783527634521>)

JOURNAL PUBLICATIONS (SINCE 2005)

- 32) Niebauer, R. T., and A.S. Robinson* (2006) "Exceptional total and functional yields of the human adenosine (A_{2a}) receptor expressed in the yeast *Saccharomyces cerevisiae*", *Prot. Exp. Purif.*, 46, p. 204-211.
- 33) Gage, M.J, Lefebvre, B.G., and A.S. Robinson* (2006) "Determinants of Protein Folding and Aggregation in P22 Tailspike," in Misbehaving Proteins, ACS Publications, eds. Regina Murphy and Amos Tsai.
- 34) Kim, J. and Robinson, A.S.* (2006) Dissociation of intermolecular disulfide bonds in P22 tailspike protein intermediates in the presence of SDS, *Protein Science*, 15 (7), p. 1791-3.
- 35) Wedekind, A.L.†, O'Malley, M., Niebauer, R.T., and Robinson, A.S.* (2006) Optimization of the Human Adenosine A_{2a} Receptor Yields in *Saccharomyces cerevisiae*, *Biotechnology Progress*, 22(5):1249-55. PMID:17022661; DOI: 10.1021/bp050431r
- 36) Powers, S.L., Robinson, C.R., and Robinson, A.S.* (2007) Denaturation of an Extremely Stable Hyperthermophilic Protein Occurs via a Dimeric Intermediate, *Extremophiles*, 11(1):179-89.
- 37) Forsten-Williams*, K.F., Cassino, T.R, Delo, L.J., Bellis, A.D., Robinson, A.S., and Ryan, T.E., (2007) Enhanced Insulin-like Growth Factor-I (IGF-I) Cell Association at Reduced pH is Dependent on IGF Binding Protein-3 (IGFBP-3) Interaction, *Journal of Cellular Physiology*, 210(2):298-308.

- 38) Famá, M.C., Raden, D., Zacchi, N., Lemos, D.R., Robinson, A.S., and Silberstein, S. * (2007) “The *Saccharomyces cerevisiae* YFR041C/ERJ5 gene encoding a type I membrane protein with a J domain is required to preserve the folding capacity of the endoplasmic reticulum” *Biochim Biophys Acta*, 1773(2):232-42.
- 39) Bane, S.E., Velasquez, J.E. †, and A.S. Robinson * (2007) “Expression and purification of milligram levels of inactive G-protein coupled receptors in *E. coli*”, *Protein Expression and Purification*, 52(2):348:355. PMID: 17166740; DOI: 10.1016/j.pep.2006.10.017
- 40) Powers, S.L. and A.S. Robinson * (2007) “PDI Improves Secretion of Redox-Inactive β -glucosidase”, *Biotech Prog.*, Mar-Apr;23(2):364-9. E-pub Feb 22, DOI: 10.1021/bp060287p
- 41) McCusker, E., Bane, S.E., O’Malley, M., and A.S. Robinson * (2007), “Heterologous GPCR expression: A bottleneck to obtaining crystal structures”, *Biotech Progress*, May-Jun;23(3):540-7. PMID:17397185
- 42) O’Malley, M., Lazarova, T., Britton, Z.T., and Robinson, A.S. * (2007) “High-level expression in *Saccharomyces cerevisiae* enables isolation and spectroscopic characterization of functional human adenosine A_{2a} receptor”, *J. Struct Biol.*, 159(2): 166-178. PMID: 17591446
- 43) McCusker, E., and Robinson, A.S.*, (2008) Refolding of G protein α subunits from inclusion bodies expressed in *Escherichia coli*, *Protein Exp. Purif.*, Apr;58(2): 342-55. PMID: 18249008
- 44) Hildebrandt, S., D. Raden, L. Petzold, A.S. Robinson, and F.J. Doyle III* (2008) “A top-down approach to mechanistic biological modeling: application to the single-chain antibody folding pathway”, *Biophysical Journal*, 95(8):3535-58. Epub 2008 Jul 18. PMID: 18641066
- 45) Webber T, Gurung S, Saul J, Baker T, Spataro M, Freyer M, Robinson AS, Gage MJ* (2009) “The C-terminus of the P22 tailspike protein acts as an independent oligomerization domain for monomeric proteins.”, *Biochem J*. May 1;419(3):595-602. PMID: 19196242
- 46) Spataro ML, Roberts CJ, Robinson AS* (2009) “Kinetic folding studies of the P22 tailspike beta-helix domain reveal multiple unfolded states.” *Biophys Chem*. 141(2-3):214-21. PMID: 19258192
- 47) Xu, P. and Robinson, A.S.* (2009) “Decreased secretion and unfolded protein response up-regulation are correlated with intracellular retention for single-chain antibody variants produced in yeast” *Biotech & Bioeng*, 104(1):20-9. PMID: 19415776
- 48) O’Malley MA, Mancini JD[†], Young CL, McCusker EC, Raden D, Robinson AS*. (2009) “Progress toward heterologous expression of active G-protein-coupled receptors in *Saccharomyces cerevisiae*: Linking cellular stress response with translocation and trafficking.” *Protein Sci*. 18(11):2356-2370. PMID: 19760666; DOI: 10.1002/pro.246
- 49) Spataro, ML and Robinson, AS* (2010) “Transgenic mouse and cell culture models demonstrate a lack of mechanistic connection between endoplasmic reticulum stress and tau dysfunction” *Journal of Neuroscience Research*, 88(9):1951-61. PMID: 20143409
- 50) O’Malley, MA, AN Naranjo, T Lazarova, AS Robinson * (2010) “Analysis of Adenosine A_{2a} Receptor Stability: Effects of Ligands and Disulfide Bonds”, *Biochemistry*, Nov 2;49(43):9181-9. PMID: 20853839
- 51) Yuraszcek, TM, Neveu, P, Rodriguez-Fernandez, M, Robinson, AS, Kosik, KS, and FJ Doyle III* (2010) Vulnerabilities in the Tau Network and The Role of Ultrasensitive Points in Tau Pathophysiology, *PLoS Computational Biology*, 6(11): e1000997. doi:10.1371/journal.pcbi.1000997 (with cover art)
- 52) O’Malley, MA, Helgeson, ME, Wagner, NJ and AS Robinson * (2011) Morphology and Composition of Cholesterol-rich Micellar Nanostructures Determine Transmembrane Protein (GPCR) Activity, *Biophysical Journal*, 100(2): L11-13. PMID: 3021673.
- 53) Sahin, E, JL Jordan, ML Spataro, AN Naranjo, WF Weiss IV, AS Robinson, EJ Fernandez *, CJ Roberts * (2011) “Computational Design and Biophysical Characterization of Aggregation-Resistant Point Mutations for γ D Crystallin Illustrate a Balance of Conformational Stability and Intrinsic Aggregation Propensity”, *Biochemistry*, Feb 8;50(5):628-39. PMID: 21184609

- 54) Young CL, Yuraszeck T, Robinson AS* (2011) “Decreased secretion and unfolded protein response upregulation,” *Methods Enzymol.* 491:235-60. PMID: 21329804
- 55) Robinson, A.* (2011) “New Tools for Breaking Barriers to GPCR Expression in *E. coli*” *Journal of Molecular Biology.* 408 (4): 597-598 PMID: 21420418
- 56) Britton, Z., Young, C., Can, Ö., McNeely, P., Naranjo, A. and Robinson, A. S. (2012) Membrane Protein Expression in *Saccharomyces cerevisiae*, in Production of Membrane Proteins: Strategies for Expression and Isolation (ed A. S. Robinson), Wiley-VCH Verlag GmbH & Co. KGaA, Weinheim, Germany. doi: 10.1002/9783527634521.ch2
- 57) O’Malley, Michelle A., Matthew E. Helgeson, Norman J. Wagner, Anne S. Robinson* (2011) “Toward Rational Design of Protein Detergent Complexes: Determinants of Mixed Micelles that are Critical for the *in vitro* Stabilization of a G-protein Coupled Receptor”, *Biophysical Journal*, 101 (8): 1938-1948. DOI: 10.1016/j.bpj.2011.09.018 PMID: 22004748
- 58) Griesemer, M, Young, CL, Doyle III, FJ, Robinson, AS and L Petzold* (2012) Spatial Localization of Chaperone Distribution in the Endoplasmic Reticulum of Yeast, *IET Systems Biology*, 6(2):54-63. <http://dx.doi.org/10.1049/iet-syb.2011.0006>
- 59) Young, CL, ZT Britton, AS Robinson,* (2012) “Recombinant Protein Expression and Purification: A Comprehensive Review of Affinity Tags and Microbial Applications” *Biotechnology Journal*, May;7(5):620-34. doi: 10.1002/biot.201100155. Epub 2012 Jan 10. PMID: 22442034
- 60) Young, CL, DL Raden, J Caplan, K Czymmek, AS Robinson,* (2012) “Cassette Series Designed for Live-Cell Imaging of Proteins and High Resolution Techniques in Yeast”, *Yeast*, Mar;29(3-4):119-36. doi: 10.1002/yea.2895. Epub 2012 Apr 4. PMID: 22473760
- 61) Britton, ZT, EI Hanle[†], and AS Robinson* (2012) An Expression and Purification System for the Biosynthesis of Adenosine Receptor Peptides for Biophysical and Structural Characterization, *Protein Exp. Purif.*, 84:224-235. PMID: 22722102; DOI: 10.1016/j.pep.2012.06.005
- 62) McNeely, P.M., Naranjo, A.N., and A.S. Robinson* (2012) Structure-function studies with G-protein coupled receptors as a paradigm for improving drug discovery and therapeutic development, *Biotechnology Journal*, 7(12): 1451-1461. DOI: 0.1002/biot.201200076 PMID: 23213015
- 63) Young, CL, DL Raden, AS Robinson,* (2013) Analysis of ER Resident Proteins in *S. cerevisiae*: Implementation of H/KDEL Retrieval Sequences, *Traffic*, 14(4):365-81. DOI: 10.1111/tra.12041. PMID: 23324027
- 64) Morozova, OA, ZM March[†], AS Robinson, and DW Colby* (2013) Conformational features of tau fibrils from Alzheimer's disease brain are faithfully propagated by unmodified recombinant protein, *Biochemistry*, 52(40):6960-7. <http://dx.doi.org/10.1021/bi400866w> PMID: 24033133
- 65) Maurer, Ronald W., Alan K Hunter, Xiangyang Wang, William K Wang, Anne S Robinson, and Christopher J. Roberts* (2013) Folding and aggregation of a multi-domain engineered immunotoxin, *Biochemical Engineering Journal*, 81: 8-14. <http://dx.doi.org/10.1016/j.bej.2013.09.015>.
- 66) Blanco, M, Sahin, E, Robinson, A, Roberts, C* (2013) Coarse-Grained Model for Colloidal Protein Interactions, B₂₂, and Protein Cluster Formation, *The Journal of Physical Chemistry, Part B*, Dec 19;117(50):16013-28. doi: 10.1021/jp409300j PMID: 24289039
- 67) Maurer, RW, AK Hunter, AS Robinson*, and CJ Roberts* (2014) Aggregates of alpha-chymotrypsinogen anneal to access more stable states, *Biotechnology & Bioengineering*, 111 (4): 782-791. doi: 10.1002/bit.25129 PMID: 24122552
- 68) St. Amand, MM, Ogunnaike, BA, and Robinson, AS* (2014) Development of At-Line Assay to Monitor Charge Variants of mAbs During Production, *Biotechnology Progress*, 30: 249–255. DOI: 10.1002/btpr.1848 PMID: 24382831

- 69) St. Amand, MM, K Tran[†], D Radhakrishnan, AS Robinson, BA Ogunnaike* (2014) Controllability Analysis of Protein Glycosylation in CHO cells, *PLoS One*, 9(2): e87973. DOI: 10.1371/journal.pone.0087973
- 70) Wu, H, R Kroe-Barrett, S Singh, AS Robinson, CJ Roberts* (2014) Competing aggregation pathways for monoclonal antibodies, *FEBS Letters* 588(6): 936-941. <http://dx.doi.org/10.1016/j.febslet.2014.01.051>
- 71) JA Costanzo, CJ O'Brien, K Tiller[†], E Tamargo[†], AS Robinson, CJ Roberts, and EJ Fernandez,* (2013) Computational Design to Control Protein Aggregation Rates Through Conformational Stability, *Protein Eng, Des, & Sel*, 27 (5): 157-167. 10.1093/protein/gzu008
- 72) St. Amand, MM, D Radhakrishnan, AS Robinson, BA Ogunnaike* (2014) Identification of Manipulated Variables for a Glycosylation Control Strategy, *Biotech Bioeng*, in press. Available online: 22 MAY 2014 DOI: 10.1002/bit.25251
- 73) Griesemer, M., Young, C., Robinson, A.S., Petzold, L. * (2014) BiP Clustering Facilitates Protein Folding in the Endoplasmic Reticulum, *PLOS Computational Biology*, 10(7):1-16, 10.1371/journal.pcbi.1003675
- 74) Young, CL and AS Robinson* (2014) Protein Folding and Secretion: Mechanistic Insights Advancing Recombinant Protein Production in *S. cerevisiae*, *Current Opinion in Biotechnology*, 30: 168-177. Available online 15 July 2014 DOI: 10.1016/j.copbio.2014.06.018
- 75) Naranjo, AN, A Chevalier[†], GD Cousins[†], E Ayettey[†], EC McCusker, C Wenk, AS Robinson (2015) Conserved disulfide bond is not essential for the adenosine A2A receptor: extracellular cysteines influence receptor distribution within the cell and ligand-binding recognition, *BBA Biomembranes*, 1848: 603-614 Available on-line Dec 5 2014 DOI: 10.1016/j.bbamem.2014.11.010
- 76) Blocker, KM, ZT Britton, AN Naranjo, PM McNeely, CL Young, AS Robinson*, (2015) Recombinant G protein-coupled receptor expression in *Saccharomyces cerevisiae* for protein characterization, in "Membrane Proteins – Production and Function Characterization", *Methods Enzymol.*, 556:165-83. doi: 10.1016/bs.mie.2014.12.025
- 77) Wu, H, K Truncali, J Ritchie, R Kroe-Barrett, S Singh, AS Robinson, and CJ Roberts* (2015) Weak protein interactions and pH- and temperature-dependent aggregation of human Fc1, *mAbs*, in press

PEER-REVIEWED CONFERENCE PROCEEDINGS

- 1) Hildebrandt, S., D. Raden, E. Bell[†], A.S. Robinson, and F.J. Doyle III* (2005) "Modeling the Unfolded Protein Response in *Saccharomyces Cerevisiae*", Proc. Int. Conf. Foundations of Systems Biology, Santa Barbara, California. [not indexed]
- 2) Griesemer, M., Young, C., Raden, D., Petzold, L., Robinson, A.S., Doyle, F.J. * (2007) "Computational Modeling of Chaperone Interactions in the Endoplasmic Reticulum of *Saccharomyces cerevisiae*." Proc. Int. Conf. Foundations of Systems Biology, Stuttgart, Germany.
- 3) Yuraszcek, T., Raden, D, Robinson, A.S., and Doyle, F.J. * (2007) "Microarray Analysis of the Unfolded Protein Response in *S. cerevisiae* Reveals Evidence of Down-regulation." Proc. Int. Conf. Foundations of Systems Biology, Stuttgart, Germany.
- 4) Yuraszcek, Theresa M., Pierre Neveu, Maria Rodriguez-Fernandez, Anne Robinson, Kenneth S. Kosik, Francis J. Doyle III (2009) Development of a mathematical model to investigate the pathophysiology of tau protein." Proc. Int. Conf. Foundations of Systems Biology, Denver, CO.

GRADUATE RESEARCH THESESPhD Students

- 1) Brian Lefebvre, May 2002; High Pressure Dissociates Tailspike Aggregates and Promotes Native Structure Formation; Assistant Professor, Rowan University, NJ (2004-2008); Current Position, Senior Research Engineer, DuPont & Co.
- 2) James Butz, June 2002; Characterizing and Optimizing GPCR Expression in Yeast; 2005-2006, Senior Scientist, Schering-Plough; 2006-2011, Associate Principal Scientist, Schering-Plough, NJ. Current position, Principal Scientist, Merck & Co.
- 3) Brenda Danek, March 2003; Characterization of the Role of Disulfides in Folding of Tailspike Protein; J.D., 2008, NYU; current position: Associate, Leydig, Voit & Mayer, LLP
- 4) Jessica Sinacola, August, 2003; Characterization and Reversal of the Aggregation of Single-Chain Antibodies; current position, Process Engineer, Sterile Process Technology & Engineering, Merck & Co, West Point, PA.
- 5) Jason Smith, July 2003; Folding and Expression of Extremophilic Enzymes; 2003-2006 Postdoctoral Fellow, Carnegie Mellon University; 2006-2007, Product Development Engineer, Cohera Medical, Pittsburgh, PA; current position, Manager of Research and Development at Carmell Therapeutics.
- 6) Ronald Niebauer, July 2005; Using GFP as a Sensor for Optimizing Expression of GPCRs; current position, Biotechnology Patent Examiner, US Patent Office, Washington, DC.
- 7) Junghwa Kim, June 2006; Roles of Folding Intermediate Conformation and Transient Disulfide Bonding on the Folding of P22 Tailspike Protein; current position, Scientist, MedImmune.
- 8) Sara Lawrence Powers, July 2006; Characterization and Expression of an Extremely Stable Hyperthermophilic Protein; 2006-2008, Postdoctoral Fellow, Wistar Institute; current position, Biomatrix.
- 9) Ping Xu, July 2006; Sensing and Analyzing the Unfolded Protein Response during Heterologous Protein Production; current position, Scientist, Glaxco Smith Kline.
- 10) Steven Bane, May 2007; Expression and Characterization of the Human Neurokinin 1 receptor from *E. coli*; Process Engineer, Sterile Process Technology & Engineering, Merck & Co, West Point, PA, 2007-2011; current position, Scientist, Amgen
- 11) Emily McCusker, December 2007; Overcoming Expression Obstacles in Producing Functional Components of the G-Protein Coupled Receptor Pathway; current position, Teva Pharmaceuticals.
- 12) Michelle Spataro, May 2009; Protein folding and aggregation *in vitro* and *in vivo*; current position, Investigator, GSK.
- 13) Michelle O'Malley, August 2009; Expression, Purification, and Biophysical Characterization of G-Protein Coupled Receptors Expressed from *Saccharomyces cerevisiae*; current position, Assistant Professor, UC Santa Barbara
- 14) Carissa Young, December 2012; Interrogation of Quality Control Mechanisms and Protein Trafficking in *Saccharomyces cerevisiae*; current position, post-doctoral fellow, MIT
- 15) Zachary Britton, December 2012; Novel Approaches to the Expression and Purification of G Protein-coupled Receptors; current position, Scientist, MedImmune
- 16) Melissa St. Amand, September 2013; Toward Online Quality Control during Biopharmaceutical Production; current position, post-doctoral fellow, Biotechnology Core Lab., ODIR, NIDDK, NIH
- 17) Andrea Naranjo, October 2014; Stability and Activity of a GPCR *in vivo* and in Membrane Mimetic Environments; post doc, NIH
- 18) Patrick McNeely, July 2015; Receptor-receptor, Ligand, and Membrane Interactions of the Adenosine A_{2A} Receptor

MChE Students

- 1) Sujata Bhatia, BChE/MChE, May 1999, Novel Role for Cysteines in *in vivo* folding of P22 Tailspike Protein; M.D./Ph.D, UPenn, 2003; Medical Research Scientist, DuPont Central Research & Development, Wilmington, DE, 2003-2011; current position, Assistant Director for Undergraduate Studies in Biomedical Engineering, Harvard University.

- Honors include: 1998 Barry M. Goldwater Award, 1999 AIChE Student Paper Award, 1999 NSF and NDSEG graduate fellowships
- 2) Nicole Sheatsley Richardson, March 2005; Optimizing Extremophile Expression in Yeast; Centocor, Malvern, PA, 2005-2008
- 3) Nathaniel Macapagal, May 2011; Characterization of the Folding and Assembly of Single-Chain Antibodies; current position, Associate Scientist I, Purification Process Sciences, MedImmune.
- 4) Nikki Ross, January 2012; The Role of Oxidative Stress on Tau Protein Homeostasis in Neurodegenerative Diseases