Shu-Bing Qian

Web Bio

Information

Biography

Biographical Statement

Professor Shu-Bing Qian received MSc and PhD degrees in Molecular Biology & Biochemistry with honors in 1997 and 2000, respectively, from Shanghai Jiaotong University Medical School (formerly Shanghai Second Medical University). He then conducted two postdoctoral fellowships at the National Institutes of Health (Bethesda, MD) and University of North Carolina (Chapel Hill, NC). Dr. Qian joined the Division of Nutritional Sciences at Cornell University in July 2008. In 2009, he received Young Investigator Award from Ellison Medical Foundation, and NIH Director's New Innovator Award. In 2010, Dr. Qian received DOD Development Award. In 2013, Dr. Qian received Peter Reeds Young Investigator Award. In 2014, Dr. Qian received DOD Idea Award.

Most of the research work in Dr. Qian's laboratory is broadly interdisciplinary, with a primary emphasis on protein synthesis, nutrient signaling pathway, and stress response. Using biochemical, genetic, and cell biological approach, the Qian laboratory investigates translational control of gene expression, molecular mechanisms of adaptive stress response, and the implications in human health and diseases. Specific disease aspects include but are not limited to, diabetes, cancer, aging and neurodegenerative disorders.

Teaching

Teaching and Advising Statement

Teaching is an exciting, enriching, and an integral component of an academic career, and I am firmly committed to excellence in teaching. My primary goal is to instill interest and to expose students to the course topics, which in turn enables me to achieve a better understanding of the material taught, which then provides all parties an opportunity to learn new material. Overall, I enjoy teaching and respect this responsibility as a core value and of critical importance to society.

Professional

Current Professional Activities

Graduate Field Membership: Nutrition; Genetics & Development; Biochemistry, Molecular & Cellular Biology, Biological and Biomedical Sciences

Faculty Member: Center for Vertebrate Genomics

Faculty Member: Chemical Biology Interface (CBI) Program Faculty Member: Leadership Program for Veterinary Students

Research

Current Research Activities

How is mRNA translation controlled by nutrient signaling? How does protein folding and degradation occur during protein synthesis? How do cells get rid of misfolded proteins? These are a few of the problems we would like to understand. Elucidation of the molecular mechanisms underlying protein quality and quantity control will ultimately define new therapeutic strategies to human diseases such as cancer, diabetes, and neurodegenerative disorders.

Specifically, we use biochemistry, cell biological, and genetic approaches to study translational control of gene expression and protein triage (folding, degradation, and aggregation) using mammalian system. We established high resolutionribosomal profiling analysis to monitor mRNA translation, which allows us to investigate ribosome dynamics as well as co-translational events. By focusing on chaperone network and the translation machinery, we are dedicated to elucidate fundamental principles of protein homeostasis.

Extension

Education

Education

PostDoc, 2004 ~ 2006 University of North Carolina, Chapel Hill, NC PostDoc, 2000 ~ 2004 National Institutes of Health, Bethesda, MD Ph.D., 2000 Shanghai Jiaotong University Medical School, Biochemistry M.Sc., 1997 Shanghai Jiaotong University Medical School, Biochemistry

Courses

Courses Taught

NS3200 - Human Biochemistry NS7030 - Seminar in Nutritional Sciences BIOG4990 - Independent Research in Biology II NS4010 - Empirical Research

Websites

Related Websites

Lab: http://qian.human.cornell.edu/

Administration

Administrative Responsibilities

Member of DNS Seminar Committee Member of BMCB Admission Committee

Publications

Selected Publications

Gao X, Wan J, Liu B, Ma M, Shen B, and Qian SB. Quantitative profiling of initiating ribosomes in vivo. Nat Methods 2015; 12(2):147-53. PMCID: in process

Han Y, Gao X, Liu B, Wan J, Zhang X, and <u>Qian SB</u>. Ribosome profiling reveals sequence-independent post-initiation pausing as a signature of translation. **Cell Res** 2014; 24(7):842-51. PMCID: PMC4085768

Liu B and <u>Qian SB</u>. Invited review: Mechanisms of translational regulation during stress. **Wiley Interdiscip Rev RNA** 2014; 5(3):301-5. PMCID: PMC3991730

Wan J and <u>Qian SB</u>. TISdb: a database for alternative translation initiation in mammalian cells. **Nucleic Acids Res** 2014; 42(1):D845-50. PMID: 24203712

Liu B and <u>Qian SB</u>. Translational reprogrammin in cellular stress response. **WIREs RNA** 2013 (in press). PMID: 24375939

Sherman MY and Qian SB. Less is more: Improving proteostasis by translation slow-down. **Trends Biochem Sci** 2013; 38(12):585-91. PMID: 24126073

Conn CS and Qian SB. mTORC1 in protein homeostasis: increase in protein quantity at the expense of quality. Sci Signal 2013; 6(271):ra24. PMID: 23592839

Liu B, Han Y, and Qian SB. Co-translational response to proteotoxic stress by elongation pausing of ribosomes. **Mol Cell** 2013; 49(3):453-463. PMID: 23290916

Liu B, Conn CS, and Qian SB. Viewing folding of nascent polypeptide chains from ribosomes. Expert Rev Proteomics 2012; 9(6):579-81. PMID: 23256666

Stern-Ginossar N, Weisburd B, Michalski A, Le VT, Hein MY, Huang SX, Ma M, Shen B, Qian SB, Hengel H, Mann M, Ingolia NT, Weissman JS. Decoding human cytomegalovirus. **Science** 2012; 338(6110):1088-93. PMID: 23180859

Lee S, Liu B, Lee S, Huang SX, Shen B, and <u>Qian SB</u>. Global mapping of translation initiation sites in mammalian cells at single-nucleotide resolution. **Proc Natl Acad Sci USA**. 2012; 109(37):E2424-32. PMID: 22927429

Han Y, David A, Liu B, Magadán JG, Bennink JR, Yewdell JW, and <u>Qian SB</u>. Monitoring co-translational protein folding in mammalian cells at codon resolution. **Proc Natl Acad Sci USA**. 2012; 109(31):12467-72. PMID: 22802618

Park WJ, Kothapalli KS, Reardon HT, Lawrence P, <u>Qian SB</u>, Brenna JT. A novel FADS1 isoform potentiates FADS2-mediated production of eicosanoid precursor

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- Zhang X, and Qian SB. Chaperone-mediated hierarchical control in targeting misfolded proteins to aggresome. **Mol Biol Cell** 2011; 22(18):3277-88
- Conn CS and Qian SB. mTOR signaling in protein homeostasis: less is more? Cell Cycle 2011; 10(12):1940-7
- Sun J, Conn CS, Han Y, Yeung V, and Qian SB. PI3K-mTORC1 attenuates stress response by inhibiting cap-independent Hsp70 mRNA translation. **J Biol Chem** 2011; 286(8):6791-800
- Qian SB, Zhang X, Sun J, Bennink JR, Yewdell JW, Patterson C. mTORC1 links protein quality and quantity control by sensing chaperone availability. **J Biol Chem** 2010; 285(35):27385-95 (co-correspondence author)
- Qian SB, Waldren L, Choudhary N, Klevit RE, Chazin WJ, Patterson C. Engineering a ubiquitin ligase reveals conformational flexibility required for ubiquitin transfer. **J Biol Chem** 2009; 284(39):26797-802 (co-correspondence author)
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- Qian SB, and Chen SS. Transduction of human hepatocellular carcinoma cell lines transduced with human gamma-interferon gene via retroviral vector. **World J Gastroenter** 1998; 4(3): 210-213
- Qian SB, Zhang TF, and Chen SS. Enhanced expression of HLA class I molecules in human hepatocellular carcinoma cell lines transduced with human gamma-interferon gene. **Chin Med J (Eng)** 1998; 111(4): 319-322