



REVOLUTIONIZING SCIENCE. ENHANCING OUR LIVES.

INSTITUTE FOR SYSTEMS BIOLOGY 2006 ANNUAL REPORT

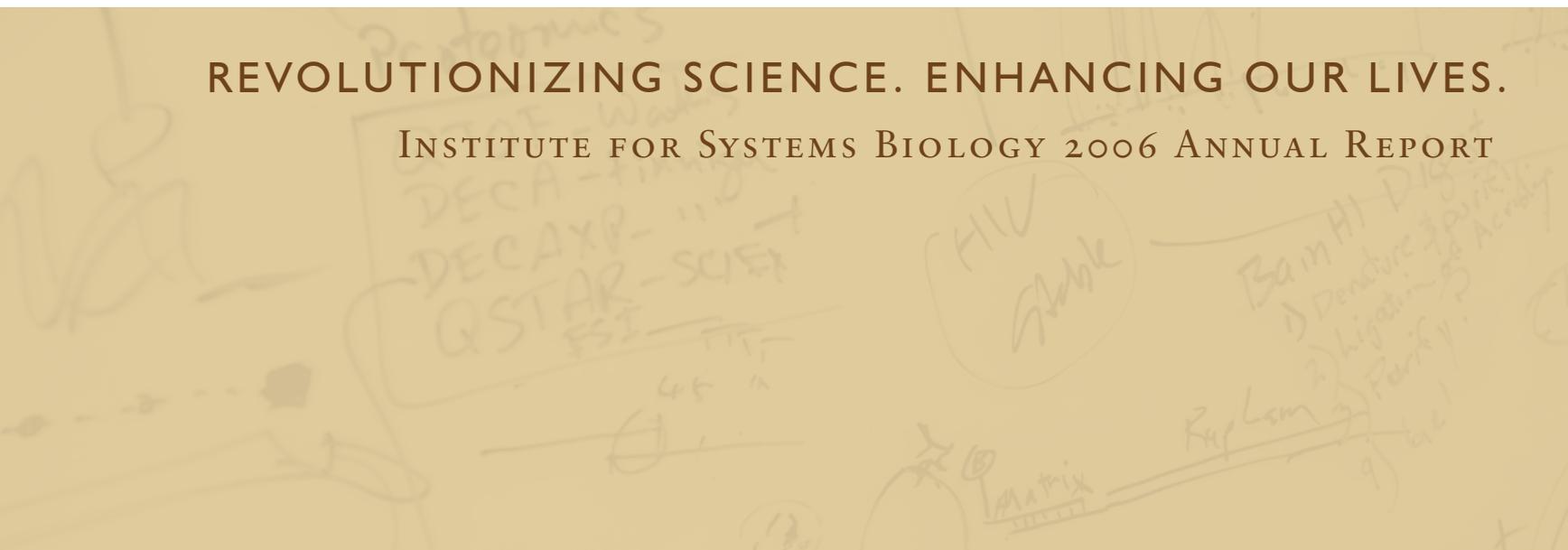


TABLE OF CONTENTS

Report from the Chairman	1
Report from the President	3
Scientific Achievements	5
Transference of Knowledge to the Community	9
Financial Tables	10
Financial Highlights	11
Development and Public Relations	13
Scientific Publications	14
Strategic Partnerships	16
Board of Directors, Executive Management, Faculty, Senior Scientists, Senior Engineers	17

REPORT FROM THE CHAIRMAN

The report that you have in your hands charts the remarkable progress that we have made at the Institute for Systems Biology (ISB). Indeed, in just a few short years, ISB has become a global leader in the advancement of “systems biology” research. In its simplest form, systems biology provides a framework for thinking about how the building blocks of biological machines interact to yield the properties that we associate with living organisms.

Having participated in the sequencing of human and other genomes in the 1990s, Lee Hood, together with immunologist Alan Aderem and proteomics specialist Ruedi Aebersold, recognized that a new sort of research entity would be required to fully realize the potential benefit to human health created by advances in our understanding of genetics. It would need to be a place where physicists, chemists, mathematicians, computer scientists and biologists could work together to build the tools necessary to understand the emergent properties of living systems.

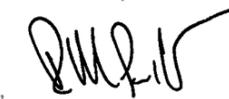
Lee, Alan and Ruedi created just such a place in ISB.

In six short years the Institute has tripled in size to more than 200 employees. It now occupies 65,000 square feet of state-of-the-art laboratories at the north end of Seattle’s Lake Union and in 2006 ISB maintained an operating budget of \$25 million. In addition, the faculty and senior scientists published 79 peer-reviewed articles and attracted \$25.4 million in new grants from government funding agencies. Private grants, gifts and commitments to the Institute yielded an additional \$6.2 million.

This annual report provides persuasive evidence of the importance of the vision originally articulated by Drs. Hood, Aderem and Aebersold. As Lee Hood notes in his letter, ISB is focusing increasing effort on the application of tools developed through systems biology research to the realization of personalized, predictive, preventive and participatory (P4) medicine.

Recognizing that research grants, especially given current federal budget constraints, cannot possibly support the full development of the Institute for Systems Biology, the ISB Board of Directors has, together with management, chartered a process aimed at developing new sources of long-range funding. I look forward to sharing the details of this plan with you next year. In the meantime, the attached summary of our activities in 2006 makes plain that the Institute for Systems Biology has adopted a truly pioneering approach, one that will dramatically advance biomedical research and, ultimately, the practice of medicine.

Sincerely,

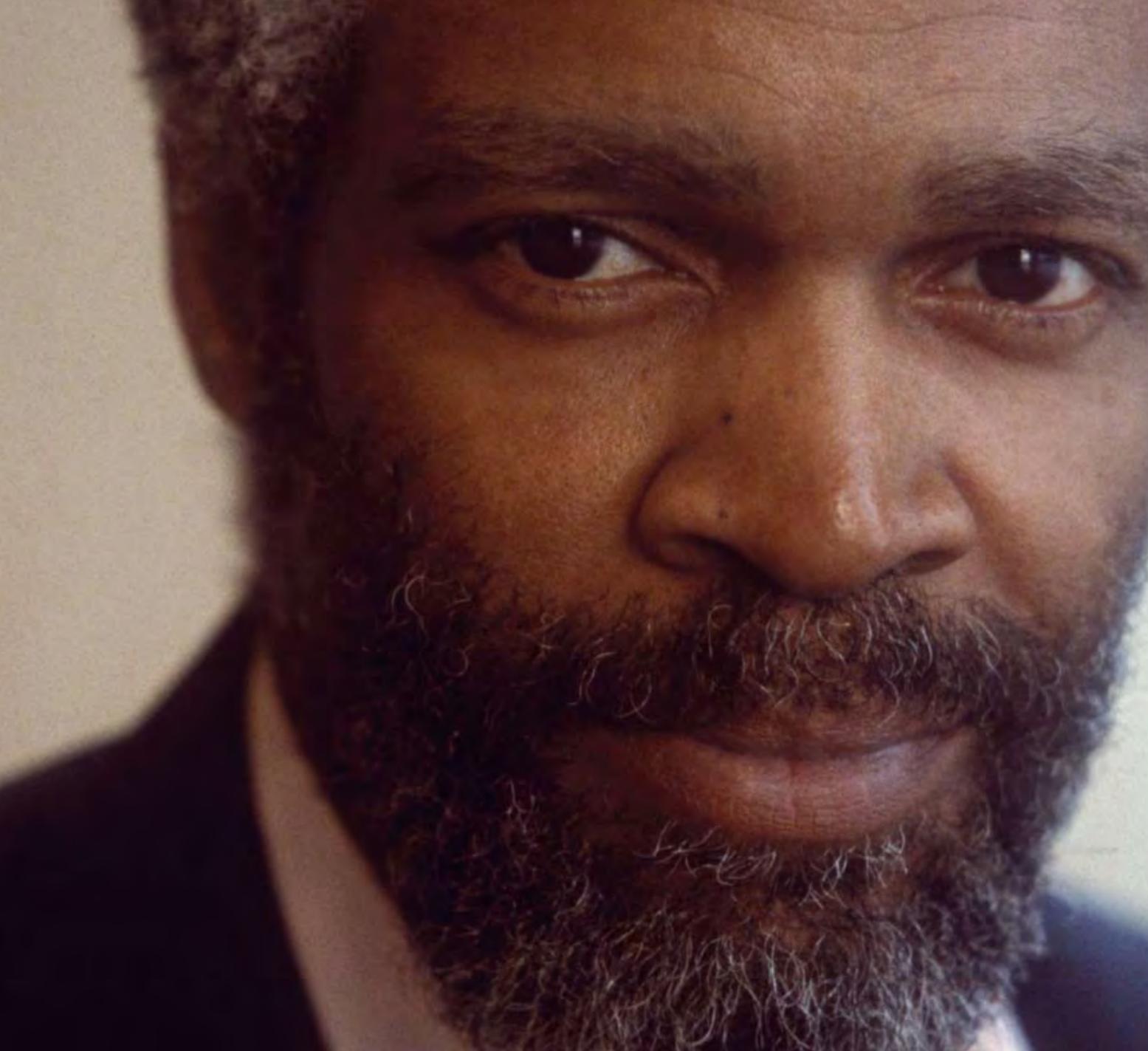


Roger M. Perlmutter, MD, PhD
Chairman of the Board



P4 MEDICINE

Medicine of the future will be personalized, predictive, preventive and participatory, examining the unique biology of an individual to assess their probability of developing various illnesses and then designing appropriate treatments, even before the onset of disease.



*“My DNA will be a crystal ball,
allowing me to see into my health future.”*

REPORT FROM THE PRESIDENT

One of the grand challenges of the 21st century is to decipher the complexity of the biological systems upon which our life depends. Systems biology, pioneered by ISB, is the most powerful approach to meeting that challenge, understanding the intricacy of biological function, and harnessing the body's biological networks to improve human health.

Until recently, biologists largely studied cellular and molecular phenomena without observing or measuring their extraordinary connectivity. The complexity of that connectivity within human biological networks requires approaching biology as an information science. Each component of the biological systems — proteins, DNA, RNA — can be represented as digital information. Using measurement technologies to generate data and computer analyses, and modeling to test hypotheses, scientists can develop strategies for predicting what happens to the functioning of a biological system when one or more elements is destroyed or mutates. That understanding will allow us to think about new strategies for early diagnosis (and hence the cure of most cancers) as well as new strategies for designing drugs for therapy, or even the prevention of cancers, neurological disorders, Type I Diabetes and many more diseases.

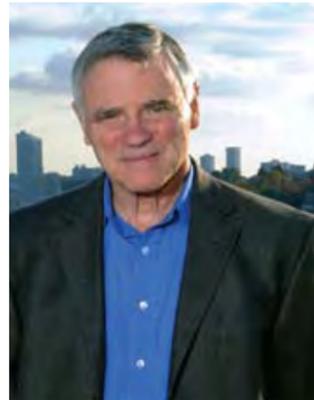
Disease arises from environmentally or genetically perturbed networks that do not return to their normal states. Thus, studying biological network malfunctions and determining how to re-engineer them to restore healthy function will provide new insights into diagnosis, therapy and prevention.

This is the basis of the new medicine based on systems biology: a medicine that will quickly evolve over the next generation from a reactive practice to a predictive, personalized, preventive and participatory (P4) process.

At ISB the systems approach is producing spectacular results. We have demonstrated how perturbed biological networks function abnormally, explaining many features of prostate cancer and mad cow disease (in mice). We are making significant progress on new technologies for vaccine development. We have demonstrated that our organs secrete protein biomarkers into the blood that allow one to distinguish health from disease, and in fact, which type of disease is present. Our scientists are developing nanotechnologies that will enable the measurement of thousands of these proteins from a droplet of blood. We are working to develop new DNA sequencing methods, which will enable us to sequence inexpensively entire individual genomes within 10 years. Finally, ISB is working on new computational and mathematical approaches to the analyses of genomes and blood protein content.

Together, these and many other advances represent the excitement and the immense promise of ISB's research community. They constitute a driving force for the emergence of P4 medicine.

Leroy Hood, MD, PhD
President



PREDICTIVE

One drop of blood is all it will take. One drop on a chip placed into a handheld device in your physician's office will tell you if you have an 85 percent chance of developing heart disease, a 12 percent chance of developing diabetes, a 70 percent chance of developing Parkinson's disease, or cancer, or virtually any other disease influenced by your genetic makeup. That knowledge will give you power.



SCIENTIFIC ACHIEVEMENTS

ISB researchers made significant progress in 2006, publishing nearly 80 peer-reviewed articles in scientific journals, receiving substantial government awards to advance systems biology approaches, broadening the scope of ISB's strategic partnerships and receiving worldwide recognition for the excellence of their work.

A new window on inflammatory disease

Using a systems approach, scientists at ISB discovered that the absence of the transcription factor ATF3 leads to inflammatory disease in mice. The inflammatory response is required for immune responses to infection, but uncontrolled, it leads to a variety of illnesses such as rheumatoid arthritis and lupus. ATF3 appears to control the balance between the beneficial and destructive arms of inflammation, and a complete understanding of this regulation could point to novel therapeutics for these diseases. These findings were published in *Nature*.

Mutations in immune system genes increase risk for infectious diseases and point the way to better drugs and vaccines

ISB researchers found significant links between mutations in innate immune genes and human susceptibility to a number of infectious diseases. The mutations are in a family of molecules called Toll-like receptors (TLRs), which alert the body to the presence of infectious agents. The results suggest new drug targets and vaccine strategies. The studies demonstrate, for example, that a mutation in TLR9 leads to rapid progression from HIV infection to the development of AIDS and a mutation in TLR2 predisposes patients to genital herpes, tuberculosis and leprosy. These observations were published in the *Proceedings of the National Academy of*

Sciences, The Journal of Experimental Medicine, The Journal of Infectious Diseases, The Journal of Immunology and AIDS.

Gene mutation a potential herald of Type 1 Diabetes

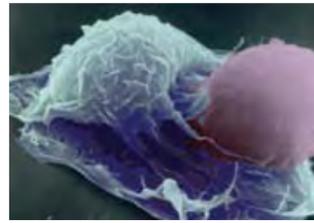
In the October 2006 issue of *The American Journal of Human Genetics* ISB researchers identified a gene, ITPR3, which is associated with Type 1 Diabetes. ITPR3 regulates calcium flow within cells and had not previously been associated with diabetes. This finding gives deeper insight into mechanisms underlying Type 1 Diabetes, a debilitating illness which generally strikes the young.

New means of defense against E. coli and Salmonella promise new therapies

ISB researchers identified two new mechanisms by which the immune system defends us from infection. In the first, reported in *Nature*, the scientists found that immune cells secrete a protein, lipocalin, which limits E. coli infections by competing with the bacterium for iron. In the second, reported in *Nature Immunology*, investigators identified a protein, Ipaf, which detects the flagella of Salmonella and triggers a robust immune response to the bacteria. Knowledge of these novel pathways could lead to the development of new antibiotics.

Computational methods facilitate development of new approaches to P4 medicine

The prediction of traits, including diseases and responses to therapies, is greatly complicated by the intricate ways in which combinations of gene variants interact. This complexity underlies the genetic challenges of predictive and personalized medicine. ISB researchers have developed computational methods that accurately predict genome-wide gene activity and precise trait outcomes



“My DNA may reveal that I have an 85 percent chance of developing breast cancer. Knowing that gives me options and could save my life.”

PREVENTIVE

If your genetic structure tells doctors you have a 95 percent chance of developing Parkinson's disease, and you know that by taking a pill and exercising starting at age 20 you could reduce that chance to less than three percent, what would you do? If protein biomarkers in your blood told physicians that your pancreas cells were beginning to mutate and become cancerous, would you take a drug that would tell your immune system to attack those cells and cure you before you experienced even a single symptom?



SCIENTIFIC ACHIEVEMENTS

(continued)

for combinations of mutations. This work, reported in *Molecular Systems Biology*, provides a core strategy for the systems biology of complex genetics.

Mathematical models lead to a better understanding of genetic diseases

ISB scientists developed a mathematical model for the genetic program leading to the formation of an essential organelle, the peroxisome. Peroxisomes have a variety of metabolic roles that implicate them in many human health concerns, including aging, neuropathology, cancer, heart disease, obesity and diabetes. These findings were reported in *Molecular Systems Biology*.

Metals research could lead to remedy for some genetic diseases

ISB researchers developed a model that explains how cells respond to metals in the environment. These results, and the systems biology approach used to map the process, may be useful in understanding the pathophysiology of a number of genetic maladies that are known to be linked to metals, such as Menkes and Wilson's diseases. This work was reported in *Genome Research*.

Quantitative methods developed to identify biomarkers

Blood biomarkers form one of the cornerstones of P4 medicine. In 2006 ISB scientists developed new methods to identify and quantify molecules in the blood that can serve as early diagnostic markers for cancer, heart disease, and the efficacy of vaccines. This work was reported in *Proteomics*, *Science*, *Genome Biology*, *Nature Biotechnology*, and *Nature Methods*.

Genetic diversity of TB bacteria influences disease transmission

ISB researchers demonstrated that mycobacterial lineages adapt to particular human populations, and that host-pathogen co-evolution impacts the transmission of tuberculosis. This led to the further observation that strain genetic diversity can influence the transmission of drug-resistant bacteria. These are important findings for the design of new tuberculosis diagnostics, drugs and vaccines. These observations were published in *Science*, the *Proceedings of the National Academies of Science* and *PLOS Pathogens*.

ISB selected by NIH to host national Center for Systems Biology

In spring 2006, ISB received a five-year, \$16.3 million grant from the National Institutes of Health (NIH) to fund a Center for Systems Biology. The Center provides professional development in systems biology approaches and fosters collaboration among the diverse researchers at the Institute: biologists, chemists, computer scientists, engineers, mathematicians and physicists. Core research funded by the grant includes genomics, proteomics, microfluidics and imaging, and informatics. The Center also provides support for K-12 education in inquiry-based science. The NIH is funding six additional centers to support systems biology work pioneered at institutes like ISB.



Alan Aderem, PhD
Co-Founder and Director

“In the future I will be cured by medications designed just for me.”

PERSONALIZED

In the future, having your genetic structure will allow physicians to provide you with medicine made for you and no one else. Not one of the hundreds of thousands of other people who share your disease but not your personal biology. These medicines will have no side-effects. They will eliminate the need for trial and error treatments that allow for disease progression, and they will cure you before you know you are sick. Having your genetic structure will give you knowledge, and that knowledge will give you the power to live intentionally rather than reactively, and to live longer and more fully.



TRANSFERENCE OF KNOWLEDGE TO THE COMMUNITY

ISB enhances people's lives and improves the human condition by transferring knowledge gained through its research to the community. It accomplishes this through support of science and math education in public schools and the transfer of new technologies to the commercial sector.

Growing new biotech companies: Accelerator Corporation

A key component of ISB's commitment to the community is Accelerator Corporation, a novel partnership among ISB and life science leaders Amgen Ventures, ARCH Venture Partners, MPM Capital, OVP Venture Partners, Versant Ventures and Alexandria Real Estate Equities, Inc. Accelerator creates early-stage companies arising from scientific discovery at the Institute and elsewhere.

Accelerator grew five emerging biotechnology companies in 2006: VieVax Corp., VLST Corp., Spaltudaq Corp., Homestead Clinical Corp. and Allozyne, Inc. These companies focus on:

- novel approaches to vaccine development;
- faster development of effective therapeutics for inflammatory and autoimmune diseases;
- developing tumor-specific therapeutic antibodies to treat cancer;
- commercializing ISB technologies with broad utility for developing diagnostic, prognostic and theranostic tools;
- commercializing technologies that improve the efficacy of protein-based therapeutics.

VLST Corp., focused on developing therapeutics for autoimmune and inflammatory disorders, raised \$55 million during 2006 as the first Accelerator company to receive series B financing.

Building scientific literacy: Center for Inquiry Science

ISB's Center for Inquiry Science (CIS) exemplifies the Institute's commitment to improving science education for every child in the state. Excellence in science education for K-12 students is critical for providing the skilled employees Washington's research and development sectors require in the next generation, as well as equipping an educated citizenry to make wise choices about the many scientifically related challenges facing our society.

Mission of CIS

To enable schools and districts to have the capacity to produce scientifically literate and capable students by creating and supporting a statewide infrastructure, comprised of collaborative and regional partnerships among schools, districts, and community partnerships that will train and support science educators.

CIS 2006 highlights

- Programming supported teachers and administrators in 21 districts representing 24 percent of Washington state's student population.
- CIS developed, coordinated and/or facilitated 76 professional development events serving 1,135 science educators.
- CIS generated more than \$800,000 in grants, gifts and fee-based revenue, approximately \$543,000 of which came from outside Washington state, to help schools build strong, inquiry-based scientific literacy among the state's student population. Grants were provided by such organizations as the National Institutes of Health, the Boeing Company, the National Science Foundation, the Washington State Office of the Superintendent of Public Instruction/Pacific Science Center and the Arthur Vining Davis Foundations.



“One day my DNA could reveal that I'm highly likely to die young of heart disease. If it does, I'll work with my doctors to create a different future.”

PARTICIPATORY

P4 Medicine will give you more information, more choices and more control than any other healthcare advance in history. You will control whether your genome is sequenced. You will control who gets to view your genetic makeup. You will sign up for biomarker tests every six months and you — no one else — will adhere to therapies (whether physical, nutritional or medicinal) to cheat your genetic destiny. Or you won't. That's the participatory part. You will have the power, and the responsibility, of using resources never before available to live your longest and best life.

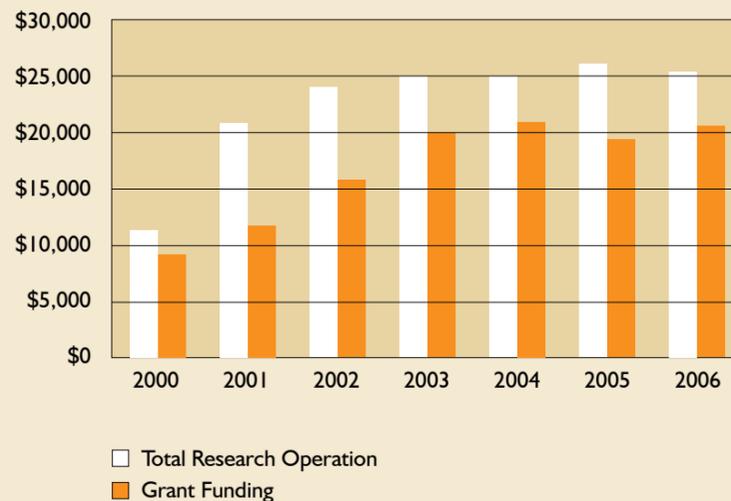
FINANCIAL TABLES

Fiscal Year Ended December 31, 2006

(Dollars in thousands)

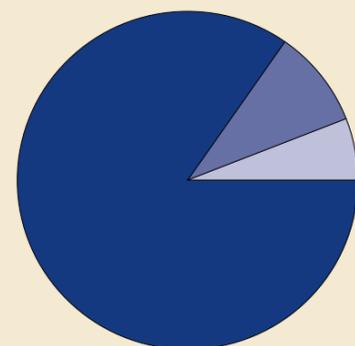
GROWTH IN OPERATIONS

Over A Seven Year Period



REVENUES

FY 2006



FINANCIAL HIGHLIGHTS

Revenue from grants and contracts, the Institute's major source of funding, increased \$1.4 million to \$20.4 million in 2006, a seven percent increase over 2005.

This positive outcome was very welcome, as 2006 was a fiscally challenging year for independent research institutions such as ISB, which heavily depend on government grants and contracts. Nationally, awards from the National Institutes of Health (NIH), the largest source of biomedical research funding, dropped slightly in inflation-adjusted terms for the second year in a row. Furthermore, it remained very difficult for younger investigators to receive grant funding, which made the climate particularly competitive for our relatively young cadre of investigators.

The excellence of the Institute's science countered these market forces, however.

ISB was awarded more than \$25 million in new government grants during 2006, including a prestigious center grant in systems biology from the NIH, which will bring ISB \$16.3 million over the next five years. In addition, ISB received new grants and contracts from private sources totaling more than \$6.2 million, including

seven-figure awards from the Bill and Melinda Gates Foundation and The Boeing Corporation. Unrestricted gifts totaled more than \$2.0 million, and investment and other income exceeded \$1.4 million, a 30 percent increase over 2005.

ISB invested just over \$800,000 in new equipment and equipment upgrades in 2006. Proteomics is a core competency of the Institute, and almost half of these funds were spent to acquire a triple quadrupole/linear ion trap mass spectrometer ideal for drug discovery and development, protein discovery, protein identification and biomarker validation. ISB also invested significantly in two state-of-the-art Leica microscopes to enhance image analysis.

Looking to grow and diversify its revenue sources to advance its mission, ISB began in 2006 to expand its fundraising and marketing efforts, hiring a Vice President for Development and other related staff. Building on the successes of 2006, ISB faculty and administrators began developing a new strategic plan for the next decade and beyond.



Gary Raisl, EdD
Vice President of Finance and Administration

FINANCIAL STATEMENTS

Balance Sheet

Cash and investments	11,662
Other assets	7,440
Property and equipment, net	8,094

Total Assets 27,195

Accounts payable and accrued expenses	1,964
Deferred revenues	913
Notes payable	11,337

Total Liabilities 14,213

Unrestricted net assets	891
Temporarily restricted net assets	3,419
Permanently restricted net assets	8,672

Total Net Assets 12,982

UNRESTRICTED REVENUES & EXPENDITURES

Revenues

Grant and contract revenue	20,364
Contributions	2,222
Investment and other income	1,414

Total Revenues 24,000

Expenditures

Research and direct costs	19,447
Management and general	5,305
Fundraising and other	433

Total Expenditures 25,185

Decrease in Net Assets (1,185)



AWARDS AND RECOGNITION

ISB co-founder and president Lee Hood received the 2006 Heinz Award in Technology, the Economy and Employment. Hood and ISB co-founder Reudi Aebersold were elected to the prestigious European Molecular Biology Organization. ISB co-founder Alan Aderem was recognized by the Society for Biomolecular Sciences for the most outstanding paper at the organization's 2006 conference in Seattle.

DEVELOPMENT AND PUBLIC RELATIONS

Since its founding in 2000, more than \$50 million in gifts and grants have been awarded to ISB by the private sector. This philanthropic investment has unleashed systems biology as the model for biological inquiry and discovery in the 21st century, leveraged millions of dollars in government grants and contracts for our scientists, and enabled ISB to build key strategic partnerships. ISB is truly a remarkable example of the power of philanthropy.

In 2006, the Institute recorded \$2.6 million in cash income from private sources. Nearly \$2.1 million was in unrestricted funds, which are invaluable in that they help launch the research and careers of our junior faculty and meet costs of other research and administrative opportunities. Program support totaled almost \$500,000, primarily advancing the work of ISB's Center for Inquiry Science, Washington's premier center of training and support for secondary school science teachers. In addition, we received a commitment of \$1 million from the Boeing Foundation, payable over four years, to support CIS. The McDonnell Foundation provided another major gift.

Sponsorship income of more than \$60,000 supported the 5th Annual ISB International Symposium, "Systems Biology and Medicine." This two-day event in April attracted more than 250 attendees, the Institute's most successful symposium ever.

Looking to ISB's future, the board of directors in 2006 decided to professionalize operations in development and public relations. The position of vice president of development was established, and I was delighted to join the Institute in September. ISB immediately began to increase substantially the number of individuals, corporations and foundations to whom the Institute can turn for gifts and grants, to boost unrestricted gift income, focus fund-raising on strategic objectives, and begin a concerted effort to raise endowment funds. All these endeavors are integrated with ISB's strategic planning, which began in late fall. The Institute is also launching a communications program to educate the public about ISB's achievements and the impact of our systems approaches on future medical care.

ISB intends to continue providing many opportunities for visionary and generous donors who want to enhance lives through our revolutionary science.

Laurence W. Herron
Vice President for Development



Major Donors

- | | |
|---------------------------------------|---------------------------------------|
| Anonymous | Invitrogen |
| Alexandria Real Estate Equities, Inc. | James S. McDonnell Foundation |
| Allen Institute for Brain Science | Lake Union Travel |
| Amgen Foundation | Valerie Logan and Leroy Hood |
| Bill and Melinda Gates Foundation | Merck & Co., Inc. |
| The Boeing Company | OVP Venture Partners |
| California Institute of Technology | Pacific Northwest National Laboratory |
| Cancer Research Institute | Roger M. Perlmutter |
| Fermentas, Inc. | Puget Sound Refrigeration |
| Frederick Frank | Henry Riggs |
| GVA Kidder Matthews | The Runstad Foundation |
| Harbor Properties, Inc. | University of Washington |
| Charles L. Hirsch | Vulcan, Inc. |
| IDT | |

2006 SCIENTIFIC PUBLICATIONS

A

Abnizova I, Rust AG, Robinson M, te Boekhorst R, Gilks WR: Transcription Binding Site Prediction Using Markov Models. *Journal of Bioinformatics and Computational Biology* 4: 425-441, 2006.

Aga RS, Fair E, Abernethy N, DeRiemer K, Paz A, Kawamura LM, Small PM, Kato-Maeda M: Microevolution of the direct repeat locus of Mycobacterium tuberculosis in a strain prevalent in San Francisco. *Journal of Clinical Microbiology* 44: 2006.

B

Bonneau R*, Reiss D, Shannon P, Facciotti MT, Hood L, Baliga NS, Thorsson V: The Inferelator: an algorithm for learning parsimonious regulatory networks from systems-biology data-sets. *Genome Biology* 7: R36, May 2006.

Bradshaw RA, Burlingame AL, Carr S, Aebersold R: Reporting protein identification data: the next generation of guidelines. *Molecular & Cellular Proteomics* 5: 787-8, 2006.

Brown V, Brown RA, Ozinsky A, Hesselberth JR, Fields S: Binding specificity of Toll-like receptor cytoplasmic domains. *European Journal of Immunology* 36: 742-753, 2006.

C

Cameron RA, Rowen L, Nesbitt R, Bloom S, Rast JP, Berney K, Arenas-Mena C, Martinez P, Lucas S, Richardson PM, Davidson EH, Peterson KJ, Hood L: Unusual gene order and organization of the sea urchin hox cluster. *Journal of Experimental Zoology, Part B, Molecular and Developmental Evolution* 15: 45-58, January 2006.

Carter GW, Rupp S, Fink GR, Galitski T: Disentangling information flow in the Ras-cAMP signaling network. *Genome Research* 16: 520-526, 2006.

Chen R, Pan S, Cooke K, White KN, Bronner MP, Goodlett DR, Aebersold R, Brentnall TA: Comparison of pancreas juice proteins from cancer versus pancreatitis using quantitative proteomic analysis. *Pancreas* (in press): 2006.

Chen R, Pan S, Crispin DA, Brentnall TA: Gene expression and proteomic analysis of pancreatic cancer: a recent update. *Cancer Genomics and Proteomics* 3: 1-10, 2006.

Chen R, Pan S, Yi EC, Donohoe S, Bronner MP, Potter JD, Goodlett DR, Aebersold R, Brentnall TA: Quantitative proteomic profiling of pancreatic cancer juice. *Proteomics* 6: 3871-9, 2006.

Chung Y, Yang X, Chang SH, Ma L, Tian Q, Dong C: Expression and regulation of IL-22 in CD4+ T lymphocyte. *Cell Research* 16: 902-7, November 2006.

D

de Jong BC, Hill PH, Brookes RH, Gagneux S, Burl S, Jeffries D, Out JK, Donkor S, Fox A, McAdam K, Small PM, Adegbola RA: Mycobacterium africanum elicits an attenuated T-cell response to Early Secreted Antigenic Target, 6kDa, in patients with tuberculosis and their household contacts. *Journal of Infectious Diseases* 193: 1279-86, 2006.

Deutsch EW, Ball CA, Bova GS, Brazma A, Bumgarner RE, Campbell DA, Causton HC, Christiansen J, Davidson D, Eichner L J, et al.: Development of the Minimum Information Specification For In Situ Hybridization and Immunohistochemistry Experiments (MISFISHIE). *OMICS* 10: 205, 2006.

Domon B, Aebersold R: Challenges and opportunities in proteomics data analysis. *Molecular & Cellular Proteomics* 5: 1921-6, 2006.

Domon B, Aebersold R: Mass spectrometry and protein analysis. *Science* 312: 212-7, 2006.

Dudley AM, Cardozo DL: Introduction to biological research: a laboratory course. *American Biology Teacher* 68: 14-24, 2006.

E

Ehlers M, Fukuyama H, McGaha TL, Aderem A, Ravetch JV: TLR9/MyD88 signaling is required for class switching to pathogenic IgG2a and 2b autoantibodies in SLE. *Journal of Experimental Medicine* 203: 553-61, 2006.

F

Fagarasanu A, Fagarasanu M, Eitzen GA, Aitchison JD, Rachubinski RA: The peroxisomal membrane protein Inp2p is the peroxisome-specific receptor for the myosin V motor Myo2p of Saccharomyces cervisiae. *Developmental Cell* 10: 587-600, 2006.

Fischback MA, Lin H, Zhou L, Yu Y, Abergel RJ, Liu DR, Raymond KN, Wanner BL, Strong RK, Walsh CT, Aderem A, Smith KD: The pathogen-associated iroA gene cluster mediates bacterial evasion of lipocalin 2. *Proceedings of the National Academies of Science USA* 103: 16502-7, 2006.

Flory MR, Lee H, Bonneau R, Mallick P, Serikawa K, Morris DR, Aebersold R: Quantitative proteomic analysis of the budding yeast cell cycle using acid-cleavable isotope-coded affinity tag reagents. *Proteomics* 6: 6146-6157, 2006.

Foltz G, Ryu GY, Yoon JG, Nelson T, Fahey J, Frakes A, Lee H, Field L, Zander K, Sibenaller Z, Ryken TC, Vibhakar R, Hood L, Madan A: Genome-Wide Analysis of Epigenetic Silencing Identifies BEX1 and BEX2 as Candidate Tumor Suppressor Genes in Malignant Glioma. *Cancer Research* 66: 1-10, 2006.

Fu E, Ramsey SA, Thariani R, Yager P: One-dimensional surface plasmon resonance imaging system using wavelength interrogation. *Review of Scientific Instruments* 77: 076106, 2006.

Fu E, Ramsey SA, Chen J, Chinowsky TM, Wiley B, Xia Y, Yager P: Resonance wavelength-dependent signal of absorptive particles in surface plasmon resonance-based detection. *Sensors & Actuators B: Chemical*: 2006.

G

Gagneux S, Davis Long C, Small PM, Van T, Schoolnik GK, Bohannan B: The Competitive Cost of Antibiotic Resistance in Mycobacterium Tuberculosis Resistance. *Science* 312: 1944-6, June 2006.

Gagneux S, Burgos M, DeReimer K, Enciso A, Munoz S, Hopewell PC, Small PM, Pym A: Impact of bacterial genetics on the transmission of isoniazid-resistant Mycobacterium tuberculosis. *PLoS Pathogens* 2: e61, 2006.

Gagneux S, DeReimer K, Van T, Kato-Maeda M, de Jong B, Narayanan S, Nicol M, Niemann S, Gutierrez C, Kremer K, Hilty M, Hopewell PC, Small PM: Variable host-pathogen compatibility in Mycobacterium tuberculosis. *Proceedings of the National Academies of Science USA* 103: 2869-73, 2006.

Gan RR, Yi C, Chiu Y, Lee H, Kao YP, Wu TH, Aebersold R, Goodlett DR, Ng WP: Proteome analysis of Halobacterium sp. NRC-1 facilitated by the biomodules analysis tool BM Sorter. *Molecular & Cellular Proteomics* 5.6: 987-997, 2006.

Gessler D, Dye C, Farmer P, Murray M, Navin T, Reves R, Shinnick T, Small P, Yates T, Simpson G: A national tuberculosis archive. *Science* 311: 1245-6, 2006.

Gilchrist M, Thorsson V, Li B, Rust AG, Korb M, Kennedy K, Hai T, Bolouri H, Aderem A: Systems Biology Approaches Identify ATF3 as a Negative Regulator of Innate Immunity. *Nature* 441: 173-8, 2006.

Glusman G, Qin S, El-Gewely R, Siegel A, Roach J, Hood L, Smit AF: A third approach to gene prediction suggests thousands of additional human transcribed regions. *PLoS Computational Biology* 2: e18, 2006.

Guo Z, Hood L, Malkki M, Petersdorf EW: Long-Range Multi-Locus Haplotype Phasing of the MHC. *Proceedings of the National Academies of Science USA* 103: 6964-6969, 2006.

Gupta N, Wollscheid B, Watts JD, Scheer B, Aebersold R, DeFranco AL: Quantitative proteomic analysis of B cell lipid rafts reveals that ezrin regulates antigen receptor-mediated lipid raft dynamics. *Nature Immunology* 7: 625-33, 2006.

H

Hardwidge PR, Donohoe S, Aebersold R, Finlay BB. Proteomic analysis of the binding partners to enteropathogenic Escherichia coli virulence proteins expressed in Saccharomyces cerevisiae. *Proteomics*: 6: 2174-9, 2006.

Hawn TR, Smith KD, Aderem A, Skerrett SJ: Myeloid Differentiation Primary Response Gene (88) - and Toll-Like Receptor 2-Deficient Mice Are Susceptible to Infection with Aerosolized Legionella pneumophila. *Journal of Infectious Diseases* 193: 1693-702, 2006.

Hawn TR, Dunstan SJ, Thwaites GE, Simmons CP, Thuong NT, Lan NT, Quy HT, Chau TT, Hieu NT, Rodrigues S, Janer M, Zhao LP, Hien TT, Farrar JJ, Aderem A: A polymorphism in toll-interleukin 1 receptor domain containing adaptor protein is associated with susceptibility to meningeal tuberculosis. *Journal of Infectious Diseases* 194: 1127-34, 2006.

Hongay CF, Grisafi PL, Galitski T, Fink GR: Antisense transcription controls cell fate in Saccharomyces cerevisiae. *Cell* 127: 735-45, 2006.

I

Iyer S, Deusch K, Lin B: Batch RNA Selector: a standalone siRNA oligo prediction program with improved sensitivity. *Computer Methods and Programs in Biomedicine*: 17197051, December 2006.

J

Jasavala R, Martinez H, Thumar J, Andaya A, Gingras AC, Eng JK, Aebersold R, Han DK, Wright ME: Identification of putative androgen receptor interaction protein modules: cytoskeleton and endosomes modulate AR signaling in prostate cancer cells. *Molecular & Cellular Proteomics* (accepted), 2006.

Jimenez-Corona ME, Garcia-Garcia L, DeRiemer K, Ferreyra-Reyes L, Bobadilla-Del-Valle M, Cano-Arellano B, Canizales-Quintero S, Martinez-Gamboa A, Small PM, Sifuentes-Osornio J, Ponce-de-Leon A: Gender differentials of pulmonary tuberculosis transmission and reactivation in an endemic area. *Thorax* 61: 384-53, 2006.

K

Kaur A, Pan M, Meislin M, El-Gewely R, Baliga NS: A systems view of haloarchaeal strategies to withstand stress from transition metals. *Genome Research* 16: 841-854, July 2006.

King N, Deusch EW, Ranish J, Nesvizhskii AI, Eddes J, Mallick P, Eng J, Desiere F, Flory M, Martin D, Kim B, Lee H, Raught B, Aebersold R: Analysis of the S. cerevisiae proteome with PeptideAtlas. *Genome Biology* 7: R106, 2006.

Kowalewska J, Smith KD, Hudkins KL, Chang A, Fogo AB, Houghton D, Leslie D, Aitchison JD, Nicosia RF, Alpers CE: Membranous glomerulopathy with spherules: an uncommon variant with obscure pathogenesis. *American Journal of Kidney Diseases* 47: 983-992, 2006.

L

Lähdesmäki H, Hautaniemi S, Shmulevich I, Yli-Harja O: Relationships Between Probabilistic Boolean Networks and Dynamic Bayesian Networks as Models of Gene Regulatory Networks. *Signal Processing* 86: 814-834, 2006.

Lee S, Pe'er D, Dudley AM, Church GM, Koller D: Identifying regulatory mechanisms using individual variation reveals key role for chromatin modification. *Proceedings of the National Academy of Sciences* 103: 14062-7, 2006.

Leslie D, Timney B, Rout MP, Aitchison JD: Studying nuclear protein import in yeast. *Methods* 39: 291-308, August 2006.

Lu W, Zhou D, Glusman G, Utleg AG, White JT, Nelson PS, Vasicek TJ, Hood L, Lin B: KLK3IP is a novel androgen regulated and transcribed pseudogene of kallikreins that is expressed at lower levels in prostate cancer cells than in normal prostate cells. *Prostate* 66: 936-44, 2006.

M

Maccarana M, Olander B, Malmstrom J, Tiedemann K, Aebersold R, Lindahl U, Li JP, Malmstrom A: Biosynthesis of dermatan sulfate: Chondroitin glucuronate C5-spimerase is identical to SART2. *Journal of Biological Chemistry* 17: 11560-11568, 2006.

Mallick P, Schirle M, Chen SS, Flory M, Lee H, Martin D, Ranish J, Raught B, Schmitt R, Werner T, Kuster B, Aebersold R: A computational approach for identifying and predicting proteotypic peptides for quantitative proteomic experiments. *Nature Biotechnology* (accepted): 2006.

Malmström J, Lee H, Nesvizhskii AI, Shteynberg D, Mohanty S, Brunner E, Ye M, Weber G, Eckerskorn C, Aebersold R: Optimized peptide separation and identification for mass spectrometry based proteomics via free-flow electrophoresis. *Journal of Proteome Research* 5: 2241-2249, 2006.

Marzolf B, Deutsch EW, Moss P, Campbell D, Johnson MH, Galitski T: SBEAMS Microarray: Database software supporting genomic expression analyses for systems biology. *BMC Bioinformatics*: 7: 286, 2006.

Miao EA, Alpuche-Aranda CM, Dors M, Clark AE, Bader MW, Miller SI, Aderem A: Cytoplasmic flagellin activates Caspase 1 and IL-1 secretion through Ipaf. *Nature Immunology* 7: 569-75, 2006.

N

Niemistro A, Selinmumi J, Saleem R, Shmulevich I, Aitchison JD, Yli-Harja O: Extraction of the Number of Peroxisomes in Yeast Cells by Automated Image Analysis. *28th IEEE EMBS Annual International Conference, Fra10.6*, September 2006.

Nykter M, Hunt KK, Pollock RE, El-Naggar AK, Taylor E, Shmulevich I, Yli-Harja O, Zhang W: Unsupervised analysis uncovers changes in histopathologic diagnosis in supervised genomic studies. *Technology in Cancer Research and Treatment* 5: 2006.

O

Orrell D, Ramsey S, Marelli M, Smith JJ, Petersen TW, de Atauri P, Aitchison JD, Bolouri H: Feedback control of stochastic noise in the yeast galactose utilization pathway. *Physica D* 217: 64-76, 2006.

P

Pan S, Wang Y, Quin JF, Pesking ER, Waichunas D, Wimberger JT, Jin J, Li J, Zhu D, Pan C, Zhang J: Identification of glycoproteins in human cerebrospinal fluid with a complementary proteomic approach. *Journal of Proteome Research* 5: 2769-2779, 2006.

Pedrioli PG, Raught B, Zhang XD, Rogers R, Aitchison J, Matunis M, Aebersold R: Automated identification of SUMOylation sites using mass spectrometry and SUMMoN pattern recognition software. *Nature Methods* 3: 533-9, 2006.

Purcell MK, Smith KD, Aderem A, Hood L, Winton JR, Roach JC: Conservation of Toll-Like Receptor Signaling Pathways in Teleost Fish. *Comparative Biochemistry and Physiology Part D: Genomics and Proteomics* 1: 77-88, 2006.

R

Ramsey S, Smith JJ, Orrell D, Marelli M, Petersen TW, de Atauri P, Bolouri H, Aitchison JD: Dual feedback loops in GAL regulon suppress cellular heterogeneity in yeast. *Nature Genetics* 38: 1082-1087, 2006.

Ramsey S, Ozinsky A, Clark A, Smith KD, de Atauri P, Thorsson V, Orrell D, Bolouri H: Transcriptional noise and cellular heterogeneity in mammalian macrophages. *Philosophical Transactions of the Royal Society London, Biological Sciences* 361: 495-506, 2006.

Reiss DJ, Baliga NS, Bonneau R: Integrated biclustering of heterogeneous genome-wide datasets for the inference of global regulatory networks. *BMC Bioinformatics*: 7: 280, June 2006.

Roach JC, Deusch K, Li S, Siegel AF, Bekris LM, Einhaus DC, Sheridan CM, Glusman G, Hood L, Lernmark A, Janer M, on behalf of the Swedish Childhood Diabetes Study Group and the Diabetes Incidence in Sweden Study Group:

Genetic mapping at 3-kilobase resolution reveals inositol 1,4,5-triphosphate receptor 3 as a risk factor for type 1 diabetes in Sweden. *American Journal of Human Genetics*: 79: 614-627, 2006.

Rowen L: Sequencing the human genome: A historical perspective on challenges for systems integration. *Micro/Nano Technology for Genomics and Proteomics*: Ed. M Ozkan and MJ Heller, Spinger: 365-398, 2006.

Rundle NT, Nelson J, Flory MR, Joseph J, Th'ng J, Aebersold R, Dasso M, Andersen RJ, Roberge M: An ent-Kaurene That Inhibits Mitotic Chromosome Movement and Binds the Kinetochores Protein Ran-Binding Protein 2. *ACCS Chemical Biology* 1: 443-450, 2006.

S

Saleem RA, Smith JJ, Aitchison JD: Proteomics of the peroxisome. *Biochimica et Biophysica Acta*: 2006.

Seebacher J, Mallick P, Zhang N, Eddes J, Aebersold R, Gelb M: Protein crosslinking analysis using mass spectrometry, isotope-coded crosslinkers and computational data processing. *Journal of Proteome Research* 5: 2270-2282, 2006.

Shannon P, Reiss D, Bonneau R, Baliga NS: The Gaggles: A system for the integrating bioinformatics and computational biology software and data sources. *BMC Bioinformatics* 7: 176, March 2006.

Shio Y, Rose DW, Donohoe S, Aebersold R, Eisenman RN: Identification and characterization of SAP25, a novel component of the mSin3 corepressor complex. *Molecular Cell Biology* 26: 1386-1397, 2006.

Shio Y, Aebersold R: Quantitative proteome analysis using isotope-coded affinity tags and mass spectrometry. *Nature Protocols* 1: 139-45, 2006.

Smith JJ, Sydorskyy Y, Marelli M, Hwang D, Bolouri H, Rachubinski RA, Aitchison JD: Expression and functional profiling reveal distinct gene classes involved in fatty acid metabolism. *Molecular Systems Biology* 2: 0009, 2006.

Stewart JJ, White JT, Yan X, Collins S, Drescher CW, Urban ND, Hood L, Lin B: Proteins Associated with Cisplatin Resistance in Ovarian Cancer Cells Identified by Quantitative Proteomic Technology and Integrated with mRNA Expression Levels. *Molecular & Cellular Proteomics* 5: 433-443, 2006

T

Tian Q: Proteomic exploration of the Wnt/ beta-catenin pathway. *Current Opinion in Molecular Therapeutics* 8: 191-7, June 2006.

True L, Coleman I, Hawley S, Huang A, Gifford D, Coleman R, Beer TM, Gelmann E, Datta M, Mostaghel E, Knudsen B, Lange P, Vessella R, Lin D, Hood L, Nelson PS: A Molecular Correlate to the Gleason Grading System for Prostate Adenocarcinoma. *Proceedings of the National Academies of Sciences USA* 103: 10991-10996, 2006.

V

Vizeacoumar FJ, Vreden WN, Fagarasanu M, Eitzen GA, Aitchison JD, Rachubinski RA: The dynamin-like protein Vps1p of the yeast Saccharomyces cerevisiae associates with peroxisomes in a Pex19p-dependent manner. *Journal of Biological Chemistry* 281: 12817-12823, 2006.

Vizeacoumar FJ, Vreden WN, Aitchison JD, Rachubinski RA: Pex19p binds Pex30p and Pex32p at regions required for their peroxisomal localization but separate from their peroxisomal targeting signals. *Journal of Biological Chemistry* 281: 14805-14812, 2006.

W

Whitehead K, Kish A, Pan M, Kaur A, Reiss DJ, King N, Hohmann L, DiRuggiero J, Baliga NS: An integrated systems approach for understanding cellular responses to gamma radiation. *Molecular Systems Biology* 2: 47, 2006.

Z

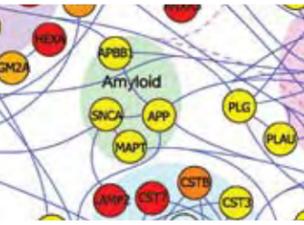
Zhang H, Liu AY, Loriaux P, Wollscheid B, Zhou Y, Watts JD, Aebersold R: Mass spectrometric detection of tissue proteins in plasma. *Molecular & Cellular Proteomics* 1: 64-71, 2006.

Zhang H, Loriaux P, Eng J, Campbell D, Keller A, Moss P, Bonneau R, Zhang N, Zhou Y, Wollscheid B, Cooke K, Yi EC, Lee H, Peskind ER, Zhang J, Smith RD, Aebersold R: UniPep, a database for human N-linked glycosites: a resource for biomarker discovery. *Genome Biology* 7: R73, 2006.

Zhang W, Shmulevich I: [Computational and Statistical Approaches To Genomics](#), 2nd Edition, Springer, 2006.

Zody MC, Garber M, Sharpe T, Young SK, Rowen L, O'Neill K, Whitaker CA, Kamal M, Chang JL, Cuomo CA, Dewar K, FitzGerald MG, Kodira CD, Madan A, Qin S, Yang X, Abbasi N, Abouelleil A, Arachchi HM, Baradarani L, Birditt B, Bloom S, Bloom T, Borowsky ML, Burke J, Butler J, Cook A, DeArellano K, DeCaprio D, Dorris L 3rd, Dors M, Eichler EE, Engels R, Fahey J, Fleetwood P, Friedman C, Gearin G, Hall JL, Hensley G, Johnson E, Jones C, Kamat A, Kaur A, Locke DP, Madan A, Munson G, Jaffe DB, Lui A, Macdonald P, Mauceli E, Naylor JW, Nesbitt R, Nicol R, O'Leary SB, Ratcliffe A, Rounsley S, She X, Sneddon KM, Stewart S, Sougnez C, Stone SM, Topham K, Vincent D, Wang S, Zimmer AR, Birren BW, Hood L, Lander ES, Nusbaum C: Analysis of the DNA sequence and duplication history of human chromosome 15. *Nature* 440: 671-5, March 2006.

STRATEGIC PARTNERSHIPS: BRIDGING ACADEMIA AND INDUSTRY



As a private, non-profit institute, ISB is positioned between academia and industry, leveraging the strengths and advantages of both. It is an organization without departments, which enables the cross-disciplinary science, training and teamwork required today to pioneer technologies and address fundamental problems in biology.

ISB's relatively small size and open organization facilitates internal collaboration and the agility to tackle complexity in those areas of technology development and computation where we enjoy unparalleled expertise. ISB cannot by itself, however, pursue all the systems biology approaches necessary for P4 medicine. As a visionary and integrating force in biology, ISB has established strategic partnerships with private companies, universities and other institutions.

ISB's academic collaborations leverage an array of expertise to undertake comprehensive analyses of scientific problems. ISB's partnerships with industry demonstrate how non-profit and for-profit alliances can rapidly develop and disseminate new methods and technologies with global impact in predicting and preventing diseases.

In 2006, ISB maintained partnerships with the following organizations:

- Applied Biosystems*
- Aviva Systems Biology*
- Battelle Memorial Institute*
- California Institute of Technology*
- Cytellect, Inc.*
- The Fred Hutchison Cancer Research Institute*
- Helicos*
- IBM*
- Lumera Corporation*
- NimbleGen Systems, Inc.*
- The Nanosystems Biology Alliance*
- The Rockefeller University*
- The Scripps Research Institute*
- University of California at Los Angeles*

Examples of 2006 strategic partner initiatives

- GenoLogics Life Sciences Software Inc., in partnership with ISB, announced integrated open source software that broadened the dissemination of new technologies and computational tools for proteomics research to scientists and labs worldwide.
- In October, Lumera Corporation and ISB extended their productive relationship to develop detection methods for an array of diagnostic biomarkers aimed at various types of cancer.
- Helicos Biosciences Corporation and ISB established a partnership that employs the company's new high throughput DNA sequencing platform to study the transcriptomes (populations of mRNAs in cells or tissues) of normal and cancer cells.

Institute for Systems Biology 2006 Board of Directors

Roger Perlmutter, MD, PhD
Chairman of the Board
Executive Vice President,
Research and Development
Amgen, Inc.

Fred Frank
Vice Chairman
Lehman Brothers, Inc.

George Rathmann, PhD
Director Emeritus
Chairman
Nuvelo, Inc.

Alan Aderem, PhD
Director
Institute for Systems Biology

Chuck Hirsch
Managing Director
Madrona Venture Group

Henry E. Riggs
President Emeritus
Keck Graduate Institute of
Applied Life Sciences

Steve Clifford
Secretary/Treasurer
Chairman
National Mobile Television

Leroy Hood, MD, PhD
President
Institute for Systems Biology

H. Jon Runstad
Co-founder, Chairman and Chief
Executive Officer
Wright Runstad & Company

Garry Menzel, PhD
Managing Director
Global Head of Life Sciences
Credit Suisse

Institute for Systems Biology 2006 Executive Management

Leroy Hood, MD, PhD
President

John Aitchison, PhD
Associate Director

Gary Raisl, EdD
Vice President of Finance

Alan Aderem, PhD
Director

Laurence W. Herron
Vice President for Development

Faculty

Alan Aderem, PhD
Ruedi Aebersold, PhD
John Aitchison, PhD
Nitin Baliga, PhD
Aimée Dudley, PhD

David Galas, PhD
Tim Galitski, PhD
Leroy Hood, MD, PhD
Dan Martin, MD
Adrian Ozinsky, MD, PhD

Jeff Ranish, PhD
Ilya Shmulevich, PhD
Peter Small, PhD

Senior Research Scientists

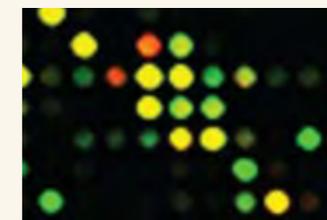
Pierre-Yves Bochud, MD
John Boyle, PhD
Greg Carter, PhD
Eric Deutsch, PhD
Alan Diercks, PhD
Sebastien Gagneux, PhD
Mark Gilchrist, PhD
Gustavo Glusman, PhD
Liz Gold, MD
Nat Goodman, PhD

Marta Janer, PhD
Christopher Lausted, MS
Simon Letarte, PhD
Bin Li, PhD
Biaoyang Lin, PhD
Monica Orellana, PhD
Shizhen Qin, PhD
Stephen Ramsey, PhD
David Reiss, PhD
Jared Roach, MD, PhD

Lee Rowen, PhD
Alistair Rust, PhD
Arian Smit, PhD
Jennifer Smith, PhD
James Spotts, PhD
Vesteinn Thorsson, PhD
Qiang Tian, MD, PhD
Julian Watts, PhD

Senior Software Engineers

Bill Longabaugh
Paul Shannon



Proteomics

QTOF - Waters
DECA - Finnigan
DECA XP - " "
QSTAR - SCIEX
ESI

HIV
Stable

Bain HI
1) Denat
2) High



Matrix

7000

3000

300



RNA ex



Institute for Systems Biology
1441 North 34th Street
Seattle, WA 98103-8904

www.systemsbiology.org

INSTITUTE FOR
**Systems
Biology**

Revolutionizing science. Enhancing life.