The Stanford Bio-X Undergraduate Summer Research Program (Stanford Bio-X USRP) is now 12 years old and has partnered with 222 Stanford faculty mentors in order to provide a ten-week summer research opportunity to 436 students.

The program aims to foster the interdisciplinary spirit of Stanford Bio-X in a new generation of up-and-coming scientists by exposing Stanford undergraduates to ten weeks of hands-on laboratory research experience. In addition to the ten weeks of laboratory research, students attend weekly faculty talks by thirty Bio-X faculty affiliates to introduce them to the cutting-edge research taking place in laboratories across campus. The program concludes with a scientific poster session alongside graduate students, faculty, and Stanford Bio-X community members from across campus and beyond.

Funding for the support of our program was provided by generous contributions from The Rose Hills Foundation, Vice Provost for Undergraduate Education, Burroughs Wellcome Fund, Tom & Dana Hayse, Ted & Colleen Friedel, Victoria Reed, William J. and Jill H. Shepherd, Vicky Rogers, Brian & Karen Mariscal, Stanford University Dean of Research Office, Mr. Vipool M. Patel and Mrs. Sharon L. Patel, Bio-X, and an Anonymous Donor.

This year, 65 students participated in the program.
2016 Stanford Bio-X Undergraduate Research Talks by Stanford Faculty:

June 22
Wolf Frommer “Interfering with a major threat: Plant pathogens”
Paul Wang “How to form your career around medical device innovation”
Melanie Hayden-Gephart “Malignant brain tumor surgery and science”

June 29
Liqun Luo “How do neurons connect with each other?”
Sean Mackey “The strain in pain lies mainly in the brain: Lessons learned from neuroimaging of pain”
Zhen Cheng “Lighting up diseases with dark materials”

July 6
Michael Snyder “Precision medicine: Managing health and disease using genomics and big data”
E.J. Chichilnisky “Electrical stimulation of the retina for design of retinal prosthesis”
Tom Clandinin “Dissecting motion processing pathways in fruit flies”

July 13
Thomas Südhof “How to construct a synapse”
Joseph Lipsick “Using Drosophila to understand cancer: The DREAM complex”

July 20
Paul Khavari “Stem cell differentiation and cancer”
Jonathan Pollack “Ameloblastoma: Oncogenesis recapitulates odontogenesis”
Lei Stanley Qi “CRISPR genome engineering for therapeutics”

July 27
Tony Wyss-Coray “Young blood for old brains”
John Oghalai “How hearing works”
Justin Gardner “Human systems neuroscience”

August 3
Noah Rosenberg “Theory of gene trees and species trees”
H. Craig Heller “Roles of sleep and circadian rhythms in learning and memory”
Erin Mordecai “Effect of temperature on Dengue, Chikungunya, and Zika transmission”

August 10
William Talbot “Using the zebrafish to investigate the vertebrate nervous system”
Manu Prakash “Simple science: Toys and Toy systems in biology, global health and science education”
Eric Gross “Peptide modulators to break specific calcineurin protein-protein interactions”

August 17
John Boothroyd “What makes Toxoplasma such a successful parasite/commensal?”
Kathleen Sakamoto “Targeting CREB for leukemia therapy”
David Relman “Better living through microbes”

August 24
Michelle Monje-Deisseroth “Myelin plasticity in health and disease”
Ada Poon “Bioelectronics”
Anthony Ricci “How hearing happens?”
2016 Program Participants:

Gunes Ates Akgun, Biology
Supported by: Stanford University Dean of Research Office
Mentor: Zhen Cheng, Radiology
Super enhancers are regions on the mammalian genome which affect the expression of more than 20 genes. Ates hypothesizes that a region of the human genome is a candidate super enhancer sequence for cardiac development that turns on multiple genes to differentiate heart progenitor cells into cardiomyocytes. Ates will implement the CRISPR-Cas 9 system to create mutant embryonic stem (ES) cell lines by knocking out this target super enhancer. By comparing the mutant cell lines to wild type, or non-mutant, ES cell lines, Ates aims to shed light on the role of this target genetic region on cardiomyocyte differentiation efficiency.

Daniel Alber, undeclared
Supported by: Stanford University Dean of Research Office
Mentor: Anthony Oro, Dermatology
Basal cell carcinoma (BCC) is the most common cancer and is derived from inappropriate signaling of the hedgehog pathway. The exceptionally high rate of mutation of advanced BCCs anoints them as an excellent model system to study mechanisms of drug resistance. Daniel is studying the role of atypical protein kinase C (aPKC) in altering the genetic sequence selectivity of Gli1, which mediates hedgehog target gene transcription.

Alan Aw, Mathematical & Computational Science
Supported by: Stanford University Dean of Research Office
Mentor: Noah Rosenberg, Biology
Alan’s research focuses on improving methods for making inferences about evolutionary trees—which appear in many areas of biology, from tracing epidemics to understanding cell lineages in tumors. Specifically, Alan is utilizing probabilistic models and mathematical structures and tools to provide novel insights into the genealogical histories of lineages sampled across closely related species.

Dylan Cable, Mathematics
Supported by: Vicky Rogers
Mentor: Justin Gardner, Psychology
Imagine you are at a baseball game: you hear the crack of the bat and the crowd cheer, and you catch a glimpse of the ball moving out of the park. You would be able to report high confidence of the ball moving out of the park, and this confidence would be well rooted not in your visual evidence of the ball moving, but rather in a strong prior expectation that when balls are hit, they move out of the park. Dylan’s project aims to dissect these two sources of confidence (prior expectations and sensory evidence) using a behavioral experiment in which a subject’s perception is biased by their priors.

2016 USRP participant Aris Kare completed his summer research training in Dr. Stanley Qi’s lab
Taylor Chavez, Bioengineering  
**Supported by: Anonymous Donor**  
**Mentor: Wolf Frommer, Biology**  
Taylor is developing a sugar sensor to measure sugar transporter activity in plant cells. The sensor designed will consist of a fusion between fluorescent proteins and a sugar transporter. The application of the sensor in glucose metabolism can advance the understanding of how sugar is used by different tissues during plant development, an essential knowledge required for the improvement of crops yields.

Annabel Chen, Biology  
**Supported by: Anonymous Donor**  
**Mentor: Eric Gross, Anesthesia**  
Annabel is studying a receptor called the “chili pepper receptor”, or transient receptor potential vanilloid 1, that is responsible for pain sensation. She is researching the effects of modifying this receptor in order to develop pain therapeutics that are non-narcotic and cardiac-safe.

Kelly Chen, Biology  
**Supported by: Vicky Rogers**  
**Mentor: Tony Wyss-Coray, Neurology**  
Kelly is investigating the mechanism by which seizures impair hippocampal function. Seizures increase secretion of vascular endothelial growth factor (VEGF) in the adult hippocampus, and a significant amount of adult hippocampal VEGF comes from neural stem and progenitor cells (NSPCs). Kelly uses the Novel Object Location task in mice to test if NSPC-VEGF is protective against hippocampal-dependent spatial-memory loss after seizure.

Michael Chen, Chemistry  
**Supported by: Anonymous Donor**  
**Mentor: Karl Deisseroth, Bioengineering and Psychiatry & Behavioral Science**  
Michael is investigating the neural basis of stress to gain a better understanding of the neural mechanisms controlling the stress response, which is critical to developing effective treatments for diseases caused by stress such as depression and PTSD. To begin answering this question, he is identifying patterns of neural activity that are linked to stressful behaviors using fiber photometry, a technique that enables optical recording of the activity of genetically targeted populations of neurons.

Dominique Cooper, Biology  
**Supported by: Anonymous Donor**  
**Mentor: Michelle Monje-Deisseroth, Neurology**  
DIPG is a deadly brain tumor in children that currently has a 0% survival rate. Dominique will be investigating the mechanism through which DIPG spreads in the brain to hopefully aid in finding a way to prevent the tumor metastasis of DIPG. To do this, she will be pulling from various disciplines, namely neuroscience, stem cell research, and cancer research.

Kristina Correa, undeclared  
**Supported by: Anonymous Donor**  
**Mentor: Theo Palmer, Neurosurgery**  
Genetic research has shown that Autism Spectrum Disorder (ASD) is associated with both environmental effects, such as an illness in the mother, and/or specific genetic conditions, such as the loss of one cMet allele. Using mice as a model organism, Kristina will use fluorescent tagging to investigate the concentration of specific neurons that have been associated with ASD in subjects who have only one cMet allele and/or whose mothers have been exposed to an infection. By studying the connection of genetics and infection of the mother to ASD, we can better understand the risk factors for, and potential causes of, ASD.
Julia Eberhard, undeclared  
Supported by: Anonymous Donor  
Mentor: Steven Artandi, Medicine (Hematology) and Biochemistry  
Julia is using a biochemical strategy to discover new proteins that are associated with a mutated protein, NPM1, specific to human leukemias. Identification of a novel pathway could lead to improvements in treatment for patients with NPM1 mutated leukemia.

Juleh Eide, Biology  
Supported by: Anonymous Donor  
Mentor: Alan Cheng, Medicine (Otolaryngology)  
Juleh is studying inner ear progenitor cells that are capable of regenerating lost sensory cells, which are required for hearing and balance functions. By damaging sensory cells in transgenic mice and staining for various markers in these progenitors, Juleh plans on mapping the steps of sensory cell regeneration in mammals. Through this work, she hopes to build the foundation of finding a cure for hearing loss and balance dysfunction in humans.

Eric Cramer, Biomedical Computation  
Supported by: Anonymous Donor  
Mentor: Sean Mackey, Anesthesia  
Prescription opioid abuse has reached epidemic proportions in our country. Many problems with abuse start with exposure to opioids during the surgical period. Eric will be creating better models to predict which people will be more vulnerable to the addictive nature of opioids. This will be followed by helping these patients with safer approaches to pain management during their surgeries.

Tai Dinger, Biology  
Supported by: Anonymous Donor  
Mentor: Aaron Gitler, Genetics  
The neurodegenerative disease Amyotrophic Lateral Sclerosis (ALS) has recently been associated with a genetic mutation that causes the expression of very long nucleotide repeats that are then translated to aberrant proteins. Very little is known about how these aberrant proteins are produced, so Tai will be performing genetic screens on a yeast model to determine the proteins that enhance translation of the aberrant protein products. This knowledge will be important in devising strategies to combat not only ALS, but other diseases associated with long nucleotide repeats.

Julia Eberhard, undeclared  
Supported by: Anonymous Donor  
Mentor: Steven Artandi, Medicine (Hematology) and Biochemistry  
Julia is using a biochemical strategy to discover new proteins that are associated with a mutated protein, NPM1, specific to human leukemias. Identification of a novel pathway could lead to improvements in treatment for patients with NPM1 mutated leukemia.
**Microglia** are specialized immune cells that regulate inflammation in the central nervous system (CNS) of vertebrates and defend the brain against foreign pathogens. Using targeted gene-editing techniques, Scott will investigate the mechanism by which a recently discovered protein in microglia, NLRC3-like, protects the CNS from inappropriate hyperinflammation. A better understanding of NLRC3-like and the inflammation pathway has important implications for diseases associated with neuroinflammation, such as Parkinson’s disease and Alzheimer’s disease.

**Victoria Fan**, Biology  
**Supported by:** Vicky Rogers  
**Mentor:** E.J. Chichilnisky, Ophthalmology and Neurosurgery  
The goal of Victoria’s research project is to contribute to the development of advanced artificial retinas for treating blindness. Current devices stimulate retinal neurons to produce crude artificial vision; however, the unintended activation of many neurons of diverse types limits their usefulness for patients. Victoria’s work is aimed at developing specific electrical stimulation patterns in the laboratory that can reduce this problem and eventually improve clinical outcomes.

**Johannah Farner**, Biology  
**Supported by:** The Rose Hills Foundation  
**Mentor:** Erin Mordecai, Biology  
Serpentine soil is toxic to most plant species, but a select few native plant species have evolved to tolerate these generally poisonous growing conditions. Has this unique chemical environment led to the evolution of a similarly specific fungal species community in serpentine grasslands? Johannah hopes to answer this question, as it will provide insight into the role that fungal pathogens play in the ecosystem dynamics of California grasslands.

**Alex Feldman**, Biology  
**Supported by:** Burroughs Wellcome Fund  
**Mentor:** Judith Frydman, Biology and Genetics  
Huntington’s Disease (HD) is one of many neurological disorders caused by the aggregation of misfolded proteins. Specifically, HD results from the aggregation of the pathogenic protein Huntingtin (Htt). Alex will investigate the mechanisms by which a specific amino acid sequence (N17) on pathogenic Htt contributes to the formation of early-stage Huntingtin oligomers and how these oligomers affect HD’s pathogenesis.

**Scott Fleming**, Management Science & Engineering  
**Supported by:** Vice Provost for Undergraduate Education  
**Mentor:** William Talbot, Developmental Biology  
Microglia are specialized immune cells that regulate inflammation in the central nervous system (CNS) of vertebrates and defend the brain against foreign pathogens. Using targeted gene-editing techniques, Scott will investigate the mechanism by which a recently discovered protein in microglia, NLRC3-like, protects the CNS from inappropriate hyperinflammation. A better understanding of NLRC3-like and the inflammation pathway has important implications for diseases associated with neuroinflammation, such as Parkinson’s disease and Alzheimer’s disease.

**Isabel Goronzy**, Chemistry  
**Supported by:** Vice Provost for Undergraduate Education  
**Mentor:** Steven Boxer, Chemistry  
To enter and infect human cells, the influenza virus must bind to molecules on host cell membranes. To identify factors that govern viral binding and determine means to interfere with viral infectivity, Isabel will examine the sterol composition and physical properties of host cell membranes.
Katie Gu, Biology
Supported by: Bio-X  
Mentor: Paul Khavari, Dermatology  
Skin diseases ranging from eczema to skin cancer affect millions of Americans each year. Using a variety of groundbreaking techniques in the Khavari lab, Katie will focus on identifying new biomolecules, termed noncoding RNAs, that control many aspects of skin development and differentiation.

Ricardo Guajardo, undeclared  
Supported by: Vice Provost for Undergraduate Education  
Mentor: Liqun Luo, Biology  
Ricardo is studying methods by which neurons are wired during development. Using fruit flies as a model organism, he aims to identify the molecular partners of Toll-like receptors (a class of proteins), which mediate a specific pathway of neural targeting. Identifying these molecules will allow for a better understanding of the molecular mechanisms behind this particular pathway and will shed light on how a structure as intricate as the brain is wired to such a degree of exactitude.

Daniel Hart, Bioengineering  
Supported by: Bio-X  
Mentors: Calvin Kuo, Medicine (Hematology); and Stanley Qi, Bioengineering and Chemical & Systems Biology  
To advance the development of cancer treatments, The Cancer Genome Atlas (TCGA) has carried out genome scale DNA sequencing efforts of a variety of cancers to identify novel druggable targets; however, these data require functional validation in a biological system. Daniel Hart will use the CRISPR/Cas9 system for genetic screens in primary organoid culture to identify key drug targets in esophageal cancer by conducting loss of function and gain of function screens.

Valerie Hau, Computer Science  
Supported by: Vice Provost for Undergraduate Education  
Mentor: Oussama Khatib, Computer Science  
The primary objective of Valerie’s research is to pinpoint and map the areas of the brain responsible for coordinated, bimanual motor movement. By using the Haptic fMRI Interface (HFI-6) in conjunction with functional magnetic resonance imaging, she hopes to determine which areas control independent arm movement and whether these align with those that control coordinated arm movement.
Karen Huynh, Electrical Engineering
Supported by: Bio-X
Mentor: Ada Poon, Electrical Engineering
Karen’s research project works to extend the use of optogenetic tools to cardiac applications. This includes designing and building soft, stretchable, implantable wireless devices to deliver light to cells in the heart. The development of these new tools will contribute greatly to the use of cardiac optogenetics to understand health functions in health and disease.

“I felt like I gained some valuable lab experience that isn’t necessarily exposed in a class-lab setting or in my previous research through the Bio department. Working on a distinct project that was my own (and not just working on running experiments for a mentor’s project) relies on a lot of skills beside experimentation and technique.”
—USRP Participant Lana Ho

Nicolas Herrera, Biology
Supported by: Bio-X
Mentor: Yanmin Yang, Neurology
Using the hypothesis that neurodegeneration in Huntington’s disease (HD) is exacerbated by a transport failure in neurons, Nicolas’s project involves designing and implementing microfluidic chambers to organize cell cultures and study intracellular trafficking. Through these chambers, drugs and viruses that support neurons can be tested for future HD therapies.

Kathleen Howell, Biology
Supported by: Bio-X
Mentor: Joe Lipsick, Pathology and Genetics
Kathleen’s research uses fruit flies as a genetic model to help understand the roles specific genes play in the cell cycle. By turning on and off these genes at various points in development, she is working to determine the relationships between these genes, how these genes are critical to normal cell cycle proceedings, and how problems with these genes can ultimately lead to cancer.

Annie Hu, Biology
Supported by: The Rose Hills Foundation
Mentor: Samuel Yang, Emergency Medicine
Sepsis is the primary cause of infection-related death in the world and accounts for 60-80% of all deaths in developing countries; however, current diagnostics for the condition are time-consuming and often lead to false-negative results, making them insufficient for proper patient treatment. Annie is developing a rapid test for sepsis using high-resolution melt and machine learning to identify pathogens directly from clinical samples while simultaneously profiling their antibiotic susceptibility. This project will improve not only patient outcomes in a clinical setting, but also the state of pathogen profiling in the scientific community.

2016 USRP participant Luladay Price completed her summer research training in Dr. Russell Fernald’s lab
Anna Jaffe, Bioengineering
Supported by: Ted & Colleen Friedel
Mentor: Karl Deisseroth, Bioengineering and Psychiatry & Behavioral Sciences
Anna’s project focuses on screening a maximally diverse population of mice during behavioral tests to discover genetic determinants of specific psychiatric disorders. She will also be mapping the corresponding structural deficits by visualizing morphological changes in the brain.

Zina Jawadi, undeclared
Supported by: Ted & Colleen Friedel
Mentor: John Oghalai, Otolaryngology
Zina is examining the role of efferent activity, which sends signals from the brain to the cochlea (the auditory portion of the inner ear), on frequency discrimination. Behavioral training will be performed to help measure frequency discrimination, which is the minimum frequency difference between two tones a mouse can differentiate. Previous experiments on anesthetized mice have found that background noise modulates efferent activity, altering how the cochlea detects sound and impacting speech perception with background noise. This experiment will lead to a better understanding of auditory efferent function, which is impaired in mice with hearing loss.

Ariana Johnson, Symbolic Systems
Supported by: Ted & Colleen Friedel
Mentor: Vinod Menon, Psychiatry & Behavioral Sciences
Ariana is investigating how children with autism process different types of memories by examining brain region activity and connectivity during the completion of memory tasks. Her project will be focused on compiling and analyzing neuropsychological, cognitive, affective, structural, and functional brain imaging data collected from children with autism.

Sharon Kam, Biology
Supported by: Burroughs Wellcome Fund
Mentor: Kathleen Sakamoto, Pediatrics
Hematopoietic stem cells (HSCs) are used to treat hematologic diseases and cancer in patients who receive stem cell transplantation. HSCs are difficult to sustain and cultivate in vitro and can be challenging to obtain from patients receiving their own stem cells due to damage from chemotherapy or radiation therapy. Sharon will optimize conditions that will allow HSCs to proliferate in culture and create various types of cocultures that mimic the bone marrow niche using mesenchymal stem cells, which reside in the bone marrow, and growth factors in both human and murine models.
Joyce Kang, Computer Science
Supported by: Burroughs Wellcome Fund
*Mentor: Ami Bhatt, Medicine (Hematology) and Genetics*
Joyce is investigating the role of the human gut microbiome, the collective genomes of all the microorganisms in the intestine, in immunocompromised patients undergoing bone marrow transplantation (BMT), a treatment that aims to cure certain blood-related cancers. Specifically, Joyce plans to use a combination of wet lab techniques, next-generation DNA sequencing, and genetic analyses of genomic data to examine the link between changes in the microbiome and graft-versus-host disease, a common and debilitating complication of BMT in which the donor stem cells attack the recipient.

Aris Kare, Bioengineering
Supported by: The Rose Hills Foundation
*Mentor: Stanley Qi, Bioengineering and Chemical & Systems Biology*
Aris is using a novel CRISPRi system to study pathological neuron cells afflicted by Huntington’s disease. He is particularly interested in down-regulating the “CAG” repeat sequence of the HTT gene that leads to the development of mutant Huntingtin protein. Ultimately, he aims to suppress the defective protein enough to halt the onset of Huntington’s disease, allowing for insight as to how mutant Huntingtin affects neuron cell physiology.

Sawa Keymeulen, Biology
Supported by: The Rose Hills Foundation
*Mentor: Stefan Heller, Otolaryngology*
Sawa is working to address the current limitations of inner-ear stem cell research by using a specific genetic marker to sort and select certain otic (relating to the ear) lineage cells. With these cells, she hopes to be able to understand their optimal conditions and bioengineer sensory epithelia for the inner ear. This research will help to better understand inner ear development and regeneration to help affected patients.

Lina Khoeur, Human Biology
Supported by: Bio-X
*Mentor: Melanie Hayden-Gephart, Neurosurgery*
Lina is testing the use of cerebro-spinal fluid (CSF) to detect mutations in the BCR-ABL1 gene, found in about 20-30% of Acute Lymphoblastic Leukemia (ALL) patients. In cases of brain metastases, the CSF is a better method of tracking mutation than the blood, since it is difficult to bypass the blood-brain barrier. If mutations can be tracked in the CSF, it could provide a much less painful and invasive way to monitor mutation and determine the most effective treatments.

Joseph Kirollos, Biology
Supported by: Bio-X
*Mentor: Giles Plant, Neurosurgery*
Joseph’s project will be exploring the changes that take place in a model of cervical spinal cord injury after transplanting human stem cells. These specialized cells have been changed to respond to light stimulation; it is hoped that once transplanted, they will be able to cause changes or even repair the circuits of the brain and spinal cord.

Nira Krasnow, Human Biology
Supported by: Bio-X
*Mentor: Ron Kopito, Biology*
Nira is studying the molecular basis of Huntington’s Disease, a neurodegenerative disease strongly associated with the misfolding and subsequent aggregation of mutant huntingtin protein. Specifically, she is testing the hypothesis that mutant huntingtin aggregation causes the destabilization of diverse, unrelated proteins within the proteome which may have deleterious cellular effects associated with Huntington’s Disease pathogenesis. This work will enhance our understanding of the functional relationship between protein aggregation and neurodegenerative disease pathogenesis.
Anna Lai, Mechanical Engineering  
Supported by: Bio-X  
**Mentor:** Manu Prakash, Bioengineering  
Chemotaxis is the movement of an organism in response to a chemical stimulus. Anna is studying a non-biological analog of chemotaxis systems—implemented in a recently discovered dynamics of two component droplets, pioneered by the Prakash Lab. These simple systems demonstrate autonomous behavior of chemotaxis (sensing and motility) and provide the first example of complex behavior of chemotaxis implemented in non-living materials. Anna will be further exploring the physical dynamics of this system, specially looking at multi-droplet systems and exploring relationships of these ideas to condensed matter physics (spin glass systems).

Andrew Lee, Bioengineering  
Supported by: Bio-X  
**Mentor:** Paul Wang, Medicine (Cardiovascular)  
Catheter ablation is an invasive procedure that has been the gold standard for treating atrial fibrillation (AF), a heart disease that affects more than two million patients in the U.S.; however, the success rate among patients with enlarged atria is less than 30% due to incomplete lesions that do not span the entire thickness of the heart tissue. To tackle this, Andrew is designing and testing a magnetic monorail ablation device that will benefit many patients who suffer from persistent AF and cannot undergo traditional catheter ablation therapy.

Angela Li, Biomedical Computation  
Supported by: Bio-X  
**Mentor:** Michael Snyder, Genetics  
Mutations that lead to misregulated translation have significant consequences for human disease. In this project, Angela will develop a mathematical model that relates mutations to regulation of translation. Eventually, she hopes to validate this mathematical model using experimental methods. This project will have a major positive impact by enhancing our understanding of the genetic basis of translation regulation in disease.

Lillian Liao, Biology  
Supported by: The Rose Hills Foundation  
**Mentor:** Calvin Kuo, Medicine (Hematology)  
*In vitro* cancer modeling is challenging due to the complex architecture of tumors and various tumor interactions with parenchymal and stromal compartments. Using tissue-culture methods developed by the Kuo lab, Lillian will be characterizing immune cell populations and optimizing their preservation. The development and characterization of this system holds promise for studying tumor immunity and developing diagnostic assays for personalized cancer therapies.
Aidan McCarty, undeclared  
**Supported by: Bio-X**  
**Mentor: Richard Zare, Chemistry**  
Aidan is investigating the use of a polymer called polypyrrole to facilitate insulin administration for diabetes. Polypyrrole is a conductive polymer that can sustainably hold the drug and release it only in response to an electric stimulus. Understanding and optimizing this process could vastly improve drug administration in diabetic patients, and the method could be applied more broadly to ailments as disparate as the common migraine and malignant cancers.

Daniel Lowet, Human Biology  
**Supported by: Tom & Dana Hayse**  
**Mentor: Ian Gotlib, Psychology**  
Daniel proposes to analyze the association between suicidal behaviors, including suicidal ideation and attempts, and white matter integrity in the brain in 11- to 14-year-old adolescents. To investigate this relation, he will analyze brain scans, model the brain’s white matter tracts with a method called tractography, and examine data from self-report measures. These analyses will increase our understanding of the brain basis of suicidality in youth.

Eric Marceau, Biology  
**Supported by: Tom & Dana Hayse**  
**Mentor: Joseph Wu, Medicine (Cardiovascular) and Radiology**  
Eric is interested in studying autophagy, a biological process by which cells in the face of stress, such as during starvation, degrade their internal parts to recycle remains as nutrients needed for survival. This process is implicated in many important diseases such as heart attack, heart failure, stroke, cancer, and diabetes. Because currently there is not a way to investigate autophagy in mice without sacrificing them, Eric is developing a novel imaging tool that would enable the long-term monitoring of this process in living mice.

Alexander Lopez, Human Biology  
**Supported by: Tom & Dana Hayse**  
**Mentor: Theo Palmer, Neurosurgery**  
Mutations in genes implicated in fetal development in combination with maternal immune challenges have the potential for increased fetal vulnerability and worsened neurological outcomes in mice. Alexander’s project investigates cell proliferation in the brains of fetuses from immune challenged mothers, with or without a genetic alteration. Investigating problems in the neurological development of the fetal brain is critical to be able to address rising diagnoses of autism spectrum disorders.

Thi Nguyen, Biology  
**Supported by: Bio-X**  
**Mentor: Calvin Kuo, Medicine (Hematology)**  
The brain possesses a highly specialized vasculature to meet its extremely active metabolic demand. Thi will be working to elucidate one of the various pathways that brain angiogenesis regulates during embryonic development. The ability to manipulate these pathways will become critical to mitigating and curing diseases like stroke and cancer.

“I realized that to a very large degree, science is a collaborative effort. My interactions with my lab members helped me understand how strongly we rely on one another for technical advice and constructive criticism. As a microcosm of the scientific community, I felt that my lab experience helped me understand that lab work is as much about the science as it is about the interpersonal relationship that one builds in the process of collaboration and learning. Furthermore, I appreciated that the Bio-X talks gave me a survey of the fruits of scientific collaboration.”

—USRP Participant Jeffrey Kwong
Persiana Saffari, Electrical Engineering  
**Supported by:** Victoria Reed  
**Mentor:** Jonathan Pollack, Pathology  
Ameloblastomas are tooth bud tumors rare in humans but remarkably common in dogs. Persiana proposes to evaluate the similarities of human and canine ameloblastomas at the genetic level. If similarities can be demonstrated, then both humans and our canine companions will benefit from one another, through the collective knowledge and therapy studies done.

Zachary Rosenthal, Chemistry  
**Supported by:** Victoria Reed  
**Mentor:** James Chen, Chemical & Systems Biology and Developmental Biology  
Zachary will be investigating the Hedgehog signaling pathway, a pathway which, when defective, can be responsible for extreme cell proliferation. Abberant Hedgehog-activated cellular growth can ultimately result in a number of cancers, including basal cell carcinoma, medulloblastoma, and meningioma. Zachary will be responsible for the development of potent inhibitors of the pathway and the characterization of their mechanisms of action.

Luladay Price, Symbolic Systems  
**Supported by:** Vice Provost for Undergraduate Education  
**Mentor:** Russell Fernald, Biology  
Using gene knockout technology, Luladay will be studying the effect of the hormone arginine vasopressin in regulating social behavior in a fish model system (*Astatotilapia burtoni*). As dominance hierarchies are conserved between humans and the model system, investigating the effects of this hormone will shed light on how aggression is regulated and how social status is maintained.

Preethi Raghavan, Bioengineering  
**Supported by:** Anonymous Donor  
**Mentor:** Michelle Monje-Deisseroth, Neurology  
Preethi is researching the mechanisms underlying cognitive impairments in patients with Neurofibromatosis type I. She is analyzing structural changes of white matter in Nf1 mice and tracking the cell lineage involved in myelination, which is essential for the functioning of the nervous system. Through this, she hopes to obtain an understanding at the cellular level of how white matter is altered in these patients and how this could cause cognitive defects.

Christina Ren, Biology  
**Supported by:** Anonymous Donor  
**Mentor:** David Relman, Medicine (Infectious Diseases)  
It has been shown that the gut microbiota has significant consequences on human health; however, much of the research to date has focused on bacteria. For this project, Christina plans to critically examine the role of fungi, especially the interactions between fungi, bacteria, and the host that could contribute to predicting gut microbiota stability and resistance to disturbances. Towards this end, she will collect and analyze samples from human participants with and without introduction of certain perturbations such as antibiotics, diet modification, and colonic cleansing.

Walter Roper, Bioengineering  
**Supported by:** Victoria Reed  
**Mentor:** Eugene Butcher, Pathology  
Walter is studying the gut region and the leukocyte interactions within mice. Through this model, he hopes to identify the factors involved in the specification of peyer patches (small masses of lymphatic tissue found within the small intestine) and the spleen. This research could impact medical procedures and testing for diseases that target the stomach, including some types of cancer.

Zachary Rosenthalal, Chemistry  
**Supported by:** Victoria Reed  
**Mentor:** James Chen, Chemical & Systems Biology and Developmental Biology  
Zachary will be investigating the Hedgehog signaling pathway, a pathway which, when defective, can be responsible for extreme cell proliferation. Abberant Hedgehog-activated cellular growth can ultimately result in a number of cancers, including basal cell carcinoma, medulloblastoma, and meningioma. Zachary will be responsible for the development of potent inhibitors of the pathway and the characterization of their mechanisms of action.
Megha Srivastava, Computer Science  
**Supported by:** Mr. Vipool M. Patel & Mrs. Sharon L. Patel  
**Mentor:** Kalanit Grill-Spector, Psychology  
Megha will seek to use computational models to explain the biological foundation of facial perception. Based on fMRI data, she hypothesizes that the size of receptive fields corresponds to our ability to distinguish human faces. By training convolutional neural networks on images of human faces, she plans to study how reducing the receptive fields in the networks affects the ability to successfully distinguish faces.

Gabriela Steiner, Human Biology  
**Supported by:** William J. & Jill H. Shepherd  
**Mentor:** Anthony Ricci, Medicine (Otolaryngology)  
Gaby is studying the cytoarchitecture of the Organ of Corti (a structure in the cochlea of the inner ear) in gerbils to better understand how mechanical interactions between cells within the Organ of Corti amplify sound waves and give the mammalian cochlea tremendous sensitivity and frequency selectivity. She aims to obtain 3D quantitative data to generate a mechanical model of the gerbil cochlea with details about the arrangement and behavior of different Organ of Corti cells. Mechanical models of the cochlea allow for functional characterization of changes arising from pathologies, genetic alterations, and future interventions, such as implantations and the regeneration of cochlear substructures using stem cells.

Grace Tam, Biology  
**Supported by:** William J. & Jill H. Shepherd  
**Mentor:** Allan Reiss, Psychiatry & Behavioral Sciences and Radiology  
Grace will be analyzing how posture plays a role in the changes of oxygenated and deoxygenated hemoglobin levels in the brain as measured with fMRI. It has been proposed that posture within the magnet can vary cognitive functioning. In order to test the hypothesis that different lying positions affect neural activity, she will be analyzing the brain scans of healthy adults. Ultimately, this research aims to provide new understandings of neuroimaging to the field of medicine.

Sonia Targ, Mathematical & Computational Science  
**Supported by:** William J. & Jill H. Shepherd  
**Mentor:** Josef Parvizi, Neurology  
Sonia’s research explores the dynamics of information exchange between brain regions when we complete computations such as $2 + 2 = 4$. Specifically, she is analyzing electrocorticographical (ECoG) data to investigate interconnectivity between the visual numeral area and the intraparietal sulcus of the human brain, elucidating the process of numerical abstraction.
Aileen Wang, undeclared  
**Supported by:** The Rose Hills Foundation  
**Mentor:** Edward Graves, Radiation Oncology  
Aileen is developing methods to treat small animal models of disease with clinically relevant radiotherapy. Her project will engineer treatment planning approaches for the X-RAD SmART system to determine the optimal set of radiation beams to produce a radiation dose to treat a specific target while sparing other tissues. The software she will develop will allow researchers to study radiotherapies matching those given to human patients.

Benjamin Thomson, Chemistry  
**Supported by:** Brian & Karen Mariscal  
**Mentor:** Steven Boxer, Chemistry  
Benjamin is using a model protein to test hypotheses about the nature of hydrogen bonds in protein structure and function. Hydrogen bonds are responsible for protein folding, assembly, and drug binding, and they play key roles in enzyme catalysis. This work is directed towards a more fundamental understanding of these non-covalent interactions which could have broad implications in protein and drug design.

Paul Tran, Biology  
**Supported by:** Brian & Karen Mariscal  
**Mentor:** Russell Fernald, Biology  
Paul is studying how animals make the most important decision of their lives: who to mate with. Studying the mate preferences of hybrid cichlid fish will provide insights into how species differ in their preferences, the mechanisms they use for deciding, and ultimately the genetic and neural processes governing mate choice.

Rebecca Triplett, Human Biology  
**Supported by:** Bio-X  
**Mentor:** Thomas Südhof, Molecular & Cellular Physiology  
The mechanisms responsible for dynamic structural and functional changes in the synapses of neurons, which are necessary for the transmission, storage, and retrieval of information in the brain, are still poorly understood. Rebecca’s proposed work addresses this important knowledge gap through employing new tools to track synaptic molecules through time and space.

Mashbayar Tugsbayar, Biology  
**Supported by:** Bio-X  
**Mentor:** H. Craig Heller, Biology  
Smith Magenis syndrome is a developmental disorder whose most prominent features include an inverted circadian rhythm and moderate learning disability. Mashbayar will be analyzing the mouse model of Smith Magenis syndrome not only to gain further understanding of this difficult yet under-researched disease, but also to explore the link between circadian rhythm disturbances and learning deficits. Learning deficits present in many other diseases such as Down syndrome and Alzheimer’s disease are hypothesized to be caused by misfiring circadian neurons; comprehensive understanding of Smith Magenis syndrome and its mechanism may help us learn how to restore not only a regular circadian rhythm but also learning and memory in patients with such diseases.

Paul Tran, Biology  
**Supported by:** Brian & Karen Mariscal  
**Mentor:** Russell Fernald, Biology  
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Rebecca Triplett, Human Biology  
**Supported by:** Bio-X  
**Mentor:** Thomas Südhof, Molecular & Cellular Physiology  
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Mashbayar Tugsbayar, Biology  
**Supported by:** Bio-X  
**Mentor:** H. Craig Heller, Biology  
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Aileen Wang, undeclared  
**Supported by:** The Rose Hills Foundation  
**Mentor:** Edward Graves, Radiation Oncology  
Aileen is developing methods to treat small animal models of disease with clinically relevant radiotherapy. Her project will engineer treatment planning approaches for the X-RAD SmART system to determine the optimal set of radiation beams to produce a radiation dose to treat a specific target while sparing other tissues. The software she will develop will allow researchers to study radiotherapies matching those given to human patients.

“The most important lesson that I learned was how to critically think about research to develop appropriate questions. Then from the questions I learned how to design experiments that would hopefully address the question… Finally I learned how to implement the experiments I have designed and interpret the results.”  
—USRP Participant Tally Buckstaff
David Zimmerman, Physics
Supported by: The Rose Hills Foundation
Mentor: Tom Clandinin, Neurobiology
Every cell packs a meter-long molecule of DNA into a micron scale nucleus by carefully folding it up, while dynamically accessing the critical genes it needs to express. However, at present, no existing technology can describe this complicated folding pattern in individual cells. David is developing a new technology that will make it possible to probe this architecture with unprecedented spatial and temporal resolution.

Eileen Williams, Psychology
Supported by: Bio-X
Mentor: Ian Gotlib, Psychology
Eileen’s research project is designed to investigate neurological correlates and potential predictors of suicidal ideation and attempts in depressed youth. Using FSL, a comprehensive library of analysis tools for brain imaging data, Eileen will contrast differences in structural and functional connectivity in depressed adolescents with a history of suicidal ideation compared to those with no such history. Ultimately, such knowledge could contribute to early identification of at-risk individuals and possible avenues of treatment.

Timothy Wu, Biology and Symbolic Systems
Supported by: Anonymous Donor
Mentor: Peter Kao, Medicine (Pulmonary & Critical Care Medicine)
Timothy is studying mechanisms by which cancers inactivate normal immune surveillance including activating the programmed cell death receptor pathway (PD-1). Timothy will use CRISPR/Cas9 technology to disrupt specific nuclear proteins, NF45 and NF90, that control PD-1 expression. The strategic goal is to restore immune responses against endogenous cancers.

Brian Wei, Science, Technology & Society
Supported by: Bio-X
Mentor: John Boothroyd, Microbiology & Immunology
Brian’s project aims to understand how the important human parasite, Toxoplasma gondii, can invade almost any cell it encounters. He will do this using a modern version of a classical genetic approach, isolating mutants that are defective in this ability and then sequencing the DNA of those mutants to identify the genes involved. This work will impact our understanding of not only Toxoplasma but also its cousin parasite, Plasmodium falciparum, that invades similarly and is the pathogen responsible for human malaria.
"Investigating Possible Super Enhancer Sequences for Cardiac Development"
Gunes Ates Akgun, Jaecheol Lee, Ningyi Shao, Joseph Wu, Zhen Cheng
Departments of Biology, Cardiology, and Radiology, Stanford University

"Visualizing the Role of Cell Cycle in Epithelial Differentiation"
Daniel Alber, Samantha Piekos, Sandra Melo, Jessica Torkelson, Lingjie Li, Gautam Shankar, Anthony Oro
Department of Dermatology, Stanford University

"Bounds on the Number of Loci Required for All Splits of a Species Tree to Appear in a Set of Gene Trees"
Alan J. Aw, Rohan S. Mehta, Lawrence H. Uricchio, David Bryant, Noah A. Rosenberg
Departments of Mathematical & Computational Science and Biology, Stanford University; Department of Mathematics & Statistics, University of Otago, New Zealand

"Inferring Orientation Tuning from fMRI Data with the Forward Encoding Model Suffers from Biased Estimation"
Dylan Cable, Taosheng Liu, Justin L. Gardner
Department of Psychology, Stanford University; Department of Psychology, Michigan State University

"Fluorescent Biosensors for Sugar Transport"
Taylor M. Chavez, Lily S. Cheung, Wolf B. Frommer
Department of Biology, Stanford University; Department of Plant Biology, Carnegie Institution for Science

"Regulation of TRPV1 and TRPV4 Membrane Trafficking"
Annabel Chen, Carl Hurt, Eric Gross
Department of Anesthesia, Stanford University

"The Role of SPARC and Its Binding Partners in Diffuse Intrinsic Pontine Glioma Invasion to the SVZ"
Dominique Cooper, Elizabeth Qin, Michelle Monje
Departments of Neuroscience, Neurology, and Biology, Stanford University

"Synergistic Effects of Maternal Immune Activation and cMet Deletion on Embryonic Neuronal Subtype Distribution"
Kristina Correa, Alex Lopez, Aditi Narayan, Brooke Babineau, Theo Palmer
Department of Neurosurgery and Institute for Stem Cell Biology & Regenerative Medicine, Stanford University

"Predicting Remote Pain and Opioid Use Cessation Using Early Trajectory Clustering"
Eric Cramer, Sean Mackey, Ian Carroll, Jennifer Hah
Department of Anesthesiology (Division of Pain Management), Stanford University

"Designing Reporters for Repeat Associated Non-ATG (RAN) Translation in HEK 293T"
Tai Dinger, Shizuka Yamada, Aaron Gitler
Department of Genetics, Stanford University
“Biochemical Purification of Protein Complexes Associated with Mutant Nucleophosmin (NPM1)”
Julia Eberhard2, Marisa Juntilla1,2, Caitlin Roake2, Natalie Ortiz2, Steven Artandi2
Departments of Pathology1 and Hematology2, Stanford University

“In Vivo Characterization of Murine Inner Ear Hair Cell Progenitors”
Juleh Eide1, Patrick J. Atkinson1, Alan G. Cheng1
Department of Otolaryngology – Head & Neck Surgery1, Stanford University

“Optimizing Single-Cell Activation in Epiretinal Prostheses to Restore Vision in People Blinded by Photoreceptor Diseases”
Victoria H. Fan1,2, Lauren E. Grosberg1,2, E.J. Chichilnisky1,2
Department of Neurosurgery1 and Hansen Experimental Physics Laboratory2, Stanford University

“Effects of Soil Chemistry on Plant Pathogen Communities”
Johannah Farner1, Erin R. Spear1, Caroline Daws1, Erin A. Mordecai1
Department of Biology1, Stanford University

“The Role of N17 in Huntingtin Pathogenesis”
Alex Feldman1, Koning Shen1, Judith Frydman1
Department of Biology1, Stanford University

“Mechanistic Investigation of Inflammation Regulation by Novel Protein NLRC3-like”
Scott L. Fleming1, Ana Meireles Sousa1, William S. Talbot1
Department of Developmental Biology1, Stanford University

“Sterol-Dependent Membrane Dynamics Regulate Influenza Virus Binding”
Isabel Goronzy1, Robert Rawle2, Peter Kasson2, Steven Boxer1
Department of Chemistry1, Stanford University; Department of Molecular Physiology & Biological Physics2, University of Virginia

“Characterizing C/D Box Small Nucleolar RNA Interactions with GTPases in Keratinocyte Differentiation”
Katie Gu1, Eon Rios1, Paul Khavari1
Department of Epithelial Biology1, Stanford University

“Identification of the Molecular Partners of Toll-6 and Toll-7 in the Developing Antennal Lobe”
Ricardo Guajardo1, Jiefu Li1, Liqun Luo1
Department of Biology1, Stanford University

“Engineering Screening Approaches to Validate Cancer Driver Genes”
Daniel Hart1, Sean de la O1, Antonia Dominguez2, Ameen Salahudeen1, Stanley Qi2, Calvin Kuo1
Departments of Medicine (Division of Hematology)1 and Bioengineering2, Stanford University

2012 Undergraduate Summer Research Program Participants

19
“Utilizing Haptic Interfaces to Understand Motor Control”
Valerie Hau¹, Alok Subbarao³, Jananan Mithrakumar², Samir Menon¹, Oussama Khatib¹
Departments of Computer Science¹ and Electrical Engineering², Stanford University; Department of Biomedical Engineering³, San Jose State University

“Steps Toward Huntington’s Disease Therapeutics: ProBDNF Treatment and the Microfluidic Co-Culture System”
Nicolas Herrera¹, Wei Wang¹, Yanmin Yang¹
Department of Neurology & Neurological Sciences¹, Stanford University

“Myb-RFP Expression and Regulation of a Polo-GFP Transgene in Larval Wing Disc Development of Drosophila”
Kathleen Howell¹,², Joseph Lipsick¹,²
Departments of Pathology¹ and Genetics², Stanford University

“Replacing Blood Culture: Combined Broad-Range Microbial ID and AST Directly from Whole Blood”
Annie Hu¹, Nadya Andini¹, Samuel Yang¹
Department of Emergency Medicine¹, Stanford University

“Variable Topography Antenna for Single-Element Beam-Steering”
Karen Huynh¹, Chris Vassos¹, Yuji Tanabe¹, Ada Poon¹
Department of Electrical Engineering¹, Stanford University

“Characterization of Single-Guide RNAs in CRISPR/Cas9 Knockin Mice for Understanding Disease Phenotypes in Genome-Wide Association Studies”
Anna Jaffe¹, Priya Rajasethupathy¹, Karl Deisseroth¹,²
Departments of Bioengineering¹ and Psychiatry & Behavioral Sciences², Stanford University

“Behavioral Assessment of Frequency Discrimination in Mice”
Zina Jawadi¹, Jinkyung Kim¹, Homer Abaya¹, John Oghalai¹
Department of Otolaryngology – Head & Neck Surgery¹, Stanford University

“Dynamic Functional Connectivity Using Resting-State fMRI in Children with Autism”
Ariana Johnson¹, Shaozheng Qin², Tiawen Chen², Rachel Rehert², Vinod Menon¹,²,³,⁴
Symbolic Systems Program¹, Departments of Psychiatry & Behavioral Sciences² and Neurology & Neurological Sciences³, and Stanford Neurosciences Institute⁴, Stanford University

“Mimicking the Human Bone Marrow: Developing a 3D Co-Culture System to Increase Hematopoietic Stem Cell Proliferation”
Sharon Kam¹, Minyoung Youn², Anupama Narla², Joy Y. Wu², Fan Yang⁴, Kathleen M. Sakamoto²
Departments of Biology¹, Pediatrics², Endocrinology³, and Bioengineering⁴, Stanford University

“Identifying Microbiome Signatures of Steroid-Refractory Graft-vs-Host Disease in HSCT Patients”
Joyce Kang¹, Tessa Andermann², Jessica Ribado¹, Katia Tkachenko¹,³, Eli Moss¹, Ami Bhatt¹,³
Departments of Genetics¹ and Medicine (Divisions of Infectious Diseases² and Hematology³), Stanford University

“Hunting for New Therapeutic Approaches: Using CRISPR Systems to Treat Huntington’s Chorea”
Aris John Kare¹, Dehua Zhao¹, Lei Stanley Qi¹,²,³
Departments of Bioengineering¹ and Chemical & Systems Biology² and ChEM-H³, Stanford University

“Selecting Otic Sensory Lineage Cells from Mouse ECS with the Fbxo2 Marker and Optimizing Conditions for Sensory Epithelia”
Sawa Keymeuen¹, Byron Hartman¹, Stefan Heller¹
Department of Otolaryngology¹, Stanford University
“Using Cell-Free RNA to Monitor BCR-ABL1 Mutations in Brain-Metastatic Leukemia”
Lina Khoeur¹, Yingmei Li¹, Melanie Hayden-Gephart¹
Department of Neurosurgery¹, Stanford University

“Optogenetic Stimulation of iPSC-Derived Corticospinal Motor Neurons for Spinal Cord Injury”
Joseph Kirollos¹, James Weiman², Giles Plant²
Departments of Biology¹ and Neurosurgery², Stanford University

“Development of Cellular Models to Assess the Effects of Mutant Huntingtin Protein Aggregation on Global Proteome Stability”
Nira Krasnow¹, Airlia Thompson¹, Ron Kopito¹
Department of Biology¹, Stanford University

“Lattice Systems of Vapor-Mediated Droplets”
Anna Lai¹, Stefan Karpitschka², Manu Prakash²
Departments of Mechanical Engineering¹ and Bioengineering², Stanford University

“Developing a Novel Bipolar Catheter Ablation System for Treating Ventricular Tachycardia”
Andrew Lee¹, Meghedi Babakhanian², Paul J. Wang¹,²
Departments of Bioengineering¹ and Cardiovascular Medicine², Stanford University

“Development of a Pipeline for Determining Allele Specific Translation”
Angela Li¹,², Can Cenik¹, Jason Reuter¹, Maheetha Bharadwaj¹, Michael Snyder¹
Department of Genetics¹ and Program in Biomedical Computation², Stanford University

“Characterizing and Maintaining Immune Cell Populations within Tumor Organoid Cultures”
Lillian Liao¹, James T. Neal¹, Iris Liu¹, Calvin Kuo¹
Department of Medicine (Division of Hematology)¹, Stanford University

“The Combinatorial Effect of Genetic Risk Factors and Maternal Immune Challenges on Embryonic Brain Development”
Alexander Lopez¹, Kristina Correa¹, Aditi Narayan¹, Brooke Babineau¹, Theo Palmer¹
Department of Neurosurgery¹, Stanford University

“Sex Differences in White Matter Correlates of Suicidal Ideation in Adolescents: A Diffusion Tensor Imaging Study”
Daniel Lowet¹, Tiffany Ho¹, Sarah Ordaz², Ian Gotlib¹
Departments of Psychology¹ and Psychiatry & Behavioral Sciences², Stanford University

“A Novel Bioluminescence Reporter-Based Sensor for Interrogating Drug-Induced Autophagy”
Eric Marceau¹, Ian Chen¹, Joseph Wu¹
Stanford Cardiovascular Institute¹, Stanford University
“Electroresponsive Polypyrrole Nanoparticles for Controlled Drug Delivery”
Aidan McCarty1, Devleena Samanta1, Niloufar Hosseini-Nassab1, Richard N. Zare1
Department of Chemistry1, Stanford University

“Embryonic Brain Vascular Phenotypes of Endothelial-Specific Gpr124 KO Mice Versus Reck KO Mice”
Thi Nguyen1, Mario Vallon1, Junlei Chang1, Calvin Kuo1
Department of Medicine (Division of Hematology)1, Stanford University

“How to Get the Girl: Quantifying Courtship Behavior in Male Cichlids Using Automated Behavior Tracking”
Luladay Price1, Scott Juntti1, Quentin Gaudry2, Russ Fernald1
Department of Biology1, Stanford University; Department of Biology2, University of Maryland

“Age- and Brain Region-Dependent Dysregulation of Oligodendrocyte Precursor Cell Population Dynamics in a Mouse Model of Neurofibromatosis Type 1”
Preethi Raghavan1, James Lennon2,3, Michelle Monje2,3
Departments of Bioengineering2 and Neurology3 and Institute for Stem Cell Biology & Regenerative Medicine2, Stanford University

“Mingling Microbes: Assessing Fungi-Bacteria Interactions in the Human Gut Microbiota”
Christina Ren1, Les Dethlefsen1, Arati Patankar1, Amy Lorber1, David Relman1
Department of Microbiology & Immunology1, Stanford University

“Transcription Factors, COUP-TFII and Nkx2-3, Act in Concert to Modulate MAdCAM1 Expression”
Walter Roper1, Thanh Theresa Dihn1, Nicole Salazar1, Julian Pan1, Milladur Rahman1, Eugene C. Butcher1,2
Laboratory of Immunology & Vascular Biology (Department of Pathology)1, Stanford University; Center for Molecular Biology & Medicine2, VA Palo Alto Health Care System

“Mechanistic Insights into a New Gli-Dependent Cancer Therapeutic”
Zach Rosenthal1, Alison Ondrus1, Marisa Hom1, James Chen1,2,3
Departments of Chemical & Systems Biology1, Developmental Biology2, and Chemistry3, Stanford University

“Man’s Best Model: A Genomic Comparison of Ameloblastoma in Humans and Dogs”
Persiana Saffari1, Boaz Arzi1, Robert West1, Frank Verstraete2, Jonathan Pollack1
Department of Pathology1, Stanford University; Department of Surgical & Radiological Sciences2, University of California, Davis School of Veterinary Medicine

“Teaching a Computer to Recognize Faces: Impact of Convolutional Neural Network Architecture and Image Variations”
Megha Srivastava1, Kalanit Grill-Spector1
Department of Psychology1, Stanford University

“In-situ Two-Photon Imaging of the Gerbil Organ of Corti”
Gabriela M. Steiner1,2, Sunil Puria1,2, Anthony J. Ricci1,2
Department of Otolaryngology – Head & Neck Surgery1 and Otobiomechanics Research Group2, Stanford University

“Effects of Posture on Resting-State Functional Magnetic Resonance Imaging (rsfMRI)”
Grace Tam1,2, Hadi Hosseini1, Allan Reiss3,4
Departments of Biology1, Psychiatry & Behavioral Sciences3, and Radiology4 and Center for Interdisciplinary Brain Sciences Research2, Stanford University

“Hello Operator: Human Neuronal Population Activity During Mathematical Cognition”
Sonia Targ1, Yuqing Zhu1, Amy Daitch1, Pedro Pinheiro-Chagas1, Josef Parvizi1
Department of Neurology & Neurological Sciences1, Stanford University
“Using Non-Canonical Amino Acids to Probe Short Hydrogen Bonds in Photoactive Yellow Protein”
Ben Thomson¹, Steven Boxer¹
Department of Chemistry¹, Stanford University

“Investigating Female Mate Choice in Malawi Cichlids”
Paul Tran¹, Alina Nguyen², Allie Byrne¹, Ryan York¹, Russell Fernald¹
Department of Biology¹, Stanford University; University of Notre Dame²

“Systematic Identification of Synaptic Ligands that Bind Astrocytic Neurexin-1”
Rebecca Triplett¹, Justin Trotter¹, Zhang Bo¹, Shane Antony Liddelow², Alexandra Munch², Ben A. Barres², Tom Südhof¹
Departments of Molecular & Cellular Physiology¹ and Neurobiology², Stanford University

“Sleep and Learning in Mouse Model of Smith Magenis Syndrome”
Mashbayar Tugsbayar¹, Bayarsaikhan Chuluun¹, H. Craig Heller¹
Department of Biology¹, Stanford University

“Optimizing Dosage of Small Animal Radiotherapy Using the PXi X-Rad SmART System”
Aileen Wang¹, Stavros Melemenidis¹, Edward Graves¹
Department of Radiation Oncology¹, Stanford University

“Toxoplasma gondii MAF1b Binds the Host Cell MIB Complex to Mediate Mitochondrial Association”
Felice D. Kelly¹, Brian M. Wei¹, Michelle L. Parker³, Martin J. Boulanger³, John C. Boothroyd¹
Department of Microbiology & Immunology¹, Stanford University; Department of Biochemistry & Microbiology², University of Victoria

“Neural Correlates of Suicidality in Depressed Adolescents”
Eileen Williams¹, Natalie Colich¹, Ian Gotlib¹
Department of Psychology¹, Stanford University

“Dynamic Binding of Novel Transcription Factors NF45 and NF90 to PD-1 Promoter upon T Cell Activation”
Timothy Ting-Hsuan Wu¹, LingFang Shi², Peter N. Kao²
Departments of Biology¹ and Medicine (Division of Pulmonary & Critical Care Medicine)², Stanford University

“Profiling the Transcriptional Response to Activity in the Drosophila Brain”
David Zimmerman¹, Thomas Clandinin¹
Department of Neurobiology¹, Stanford University

To view all Stanford Bio-X USRP poster titles and faculty talks from previous years, please visit: http://biox.stanford.edu/research/undergraduate-research
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