# Pacific Symposium on Biocomputing 2016 January 4-8, 2016 Big Island of Hawaii

Program Book



#### PACIFIC SYMPOSIUM ON BIOCOMPUTING 2016

Big Island of Hawaii, January 4-8, 2016

Welcome to PSB 2016!

We have prepared this program book to give you quick access to information you need for PSB 2016.

Enclosed you will find

- Logistics information
- Menus for PSB hosted meals
- Full conference schedule
- Call for Session and Workshop Proposals for PSB 2017
- Poster/abstract titles and authors
- Participant List

Conference materials are also available online at http://psb.stanford.edu/conference-materials/.

PSB 2016 gratefully acknowledges the support the Institute for Computational Biology, a collaborative effort of Case Western Reserve University, the Cleveland Clinic Foundation, and University Hospitals; the National Institutes of Health (NIH), the National Science Foundation (NSF); and the International Society for Computational Biology (ISCB). If you or your institution are interested in sponsoring, PSB, please contact Tiffany Murray at psb.hawaii@gmail.com

If you have any questions, the PSB registration staff (Tiffany Murray, Georgia Hansen, Brant Hansen, Kasey Miller, and BJ Morrison-McKay) are happy to help you.

Aloha!

Russ Altman Keith Dunker Larry Hunter Teri Klein Maryln Ritchie

The PSB 2016 Organizers

#### PACIFIC SYMPOSIUM ON BIOCOMPUTING 2016

Big Island of Hawaii, January 4-8, 2016

#### SPEAKER INFORMATION

Oral presentations of accepted proceedings papers will take place in Salon 2 & 3. Speakers are allotted 10 minutes for presentation and 5 minutes for questions for a total of 15 minutes. Instructions for uploading talks were sent to authors with oral presentations. If you need assistance with this, please see Tiffany Murray or another PSB staff member.

#### POSTER INFORMATION

Please set up your poster upon arrival. Please use the easel marked with the same number that your poster abstract has been assigned in abstract book (e.g., if your abstract is on page 50 of the book, use the easel marked #50). Once your poster is displayed, please make sure that the number is still visible. *Please remove your poster by 9:00pm, Thursday, January 7.* 

#### The poster session is on Wednesday, January 6 12:30-2:30.

12:30-1:30: Posters on ODD numbered boards presented. 1:30-2:30: Posters on EVEN numbered boards presented.

The poster session is centered around lunch. Please eat, talk, and network.

#### INTERNET CONNECTIVITY

PSB is connected to the Internet via a secure wireless network for the conference. For transferring your data/presentations or technical difficulties with the network, please see PSB IT/AV staff members Al Conde, Paul Murray, or Ryan Whaley. PSB's wireless Internet access (802.11b/g/n) is available in the Salons 1, 2, 3, the Plaza Ballroom, and the Ballroom Courtyard of the Fairmont Orchid.

Specific instructions to connect to the PSB wireless network will be posted onsite.

#### PACIFIC SYMPOSIUM ON BIOCOMPUTING 2016

Big Island of Hawaii, January 4-8, 2016

#### PSB 2016 HOSTED RECEPTIONS & MEALS

#### Monday, January 4

7:30-8:30pm 21<sup>st</sup> Anniversary Reception, Kilohana Room (near Concierge)

Sponsored by Case Western Reserve University, the Cleveland

Clinic Foundation, and University Hospitals

#### Tuesday, January 5

7:30-8:30am Continental Breakfast, Ballroom Courtyard 12:00-1:15pm PSB Hosted Lunch, Ballroom Courtyard

#### Wednesday, January 6

7:30-8:30am Continental Breakfast, Ballroom Courtyard

12:30-2:30pm Poster Session with Lunch, Salon 1 and Ballroom Courtyard

#### Thursday, January 7

7:30-8:30am Continental Breakfast, Ballroom Courtyard

5:30-7:30pm Dinner Party, Coconut Grove

#### Friday, January 8

7:30-8:30am Continental Breakfast, Ballroom Courtyard

Non-participant meal tickets may be purchased at the PSB registration desk:

- All Meals: \$550 adults; \$275 children 6-12 years
- Reception Ticket Only: \$35 adults; \$15 children 6-12 years
- Banquet Ticket Only: \$150 adults; \$75 children 6-12 years
- Reception & Banquet Only: \$180 adults; \$85 children 6-12 years

Please remember to wear your PSB nametag & lanyard at all PSB functions.

Enjoy the conference!

#### Pacific Symposium on Biocomputing 2016 Hosted Meal Menus

#### Monday, January 4

#### 21<sup>st</sup> Anniversary Reception (7:30-8:30pm)

Wine, beer, champagne, soft drinks, coffee & tea service

Triple Chocolate Mousse Cake decorated with the gold strawberries, passion fruit sauce & raspberries

#### Tuesday, January 5

#### Breakfast (7:30am-8:30am)

Orange Juice, Guava Juice & Pineapple Juice

Sliced Fresh Seasonal Fruit Tray

Island Fruit Breads, Freshly Baked Croissants, Tropical Danishes & Muffins, Sweet Butter & Tropical Fruit Preserves

Puako Muffin - Diced Portuguese Sausage, Scrambled Eggs & Monterey Jack Cheese {vegetarian option included}

Selection of Dry Cereals & Muesli

Royal Kona Blend Coffee, Decaffeinated Coffee & Fairmont Signature Teas

#### Lunch (12:00pm-1:15pm)

Freshly Tossed Kekela Farms Baby Greens with Garden Vegetables, Spiced Mac Nuts, Caramelized Hilo Papaya & Ka'u Orange Vinaigrette

German-style Potato Salad

Strawberry Papaya with Fresh Lime

Taro & Maui Potato Chips

Toy Dogs: Wagyu Mini Dogs {& Vegetarian Option} Pretzel Buns, Pickle Relish, Hot Peppers, Onions, Mustard, Ketchup, Salsa, Shredded Cheese, Paniolo Cowboy Chili

Chocolate Chip Macadamia Nut Pie

Mango Ginger Crisp with Raw Sugar Streusel

Freshly Brewed Iced Tea & Lemonade

#### Wednesday, January 6

#### Breakfast (7:30am-8:30am)

Orange Juice, Guava Juice & Pineapple Juice

Sliced Fresh Seasonal Fruit Tray

Island Fruit Breads, Freshly Baked Croissants, Tropical Inspired Danishes & Muffins, Sweet Butter & Tropical Fruit Preserves

Selection of Dry Cereals & Muesli

South Point Vegetable Burrito: Spinach, Sweet Peppers, Onions, Cheddar Cheese, Black Beans & Brown Rice, served with Tomato Salsa

Royal Kona Blend Coffee, Decaffeinated Coffee & Fairmont Signature Teas

#### Lunch (12:30pm-2:30pm)

Island Baby Romaine, Focaccia Croutons, Traditional Caesar Dressing, Shaved Parmesan & Julienne Prosciutto

Tropical Fruit Display

Cole Slaw

Pickled Vegetable Salad

Taro Chips and Maui Chips

Sliders on Hawaiian Sweet Rolls: Kahua Ranch Wagyu Beef Slider, Kalua Pork Slider, Truffle Mushroom & Tofu Slider

Assorted Ice Cream Bars

Freshly Brewed Iced Tea, Lemonade, Beer, Wine, and Soft Drinks

#### Pacific Symposium on Biocomputing 2016 Hosted Meals (con't)

#### Thursday, January 7

#### Breakfast (7:30am-8:30am)

Orange Juice, Guava Juice & Pineapple Juice

Sliced Fresh Seasonal Fruit Tray

Island Fruit Breads, Freshly Baked Croissants, Tropical Inspired Danishes & Muffins, Sweet Butter & Tropical Fruit Preserves

Selection of Dry Cereals & Muesli

Hilo Sunrise Wrap: Egg, Chorizo, Cheddar Cheese & Cilantro with Mango-Pineapple Salsa {vegetarian option included}

Royal Kona Blend Coffee, Decaffeinated Coffee & Fairmont Signature Teas

#### Dinner Reception (5:30pm-7:30pm)

Fresh Baby Greens, Tomato Wedges, Cucumbers, Shaved Onions, Sliced Mushrooms, White Balsamic Vinaigrette & Ranch Dressing

Waimea Tomato & Marinated Artichoke Salad

Big Island Style Ahi Poke with Crushed Kukui Nuts, Onions, Ogo, Shoyu, Sesame Oil & Sambal Dim Sum Station - Char Siu Bao, Island Style Pot Stickers, Pork & Shrimp Siu Mai, Shrimp Har Gau, Soy-Mustard Ponzu Sauce & Sweet Chili Sauce

Micro Steam Buns & Hoisin Kalua Pork

Furikake Seared Mahi Mahi, Lemongrass Emulsion

Imu-Style Cooked Kalua Pork

Mango Huli Chicken, Papaya & Edamame Lau'ai

Crispy Curry Tofu, Japanese Eggplant, Asparagus Shiitake Mushrooms, Sautéed Spinach with Thai Curry Sauce

Chef's Seasonal Vegetable

Kim Chee & Vegetable Fried Rice

Saffron Passion Fruit Panna Cotta, Popping Boba

Choice of Cupcakes: MacNut, Red Velvet, Chocolate & Pistachio

Royal Kona Blend Regular & Decaffeinated Coffee & Fairmont Signature Teas

Complete open bar including cocktails, blended drinks, wine, beer, non-alcoholic drinks

#### Friday, January 8

#### Breakfast (7:30am-8:30am)

Orange Juice, Guava Juice & Pineapple Juice

Sliced Fresh Seasonal Fruit Tray

Island Fruit Breads, Freshly Baked Croissants, Tropical Inspired Danishes & Muffins, Sweet Butter & Tropical Fruit Preserves

Selection of Dry Cereals & Muesli, Individual Greek Yogurt, Plain & Assorted Fruit Flavored Island Quiche: Maui Onions & Pipikaula (vegetarian option included)

Royal Kona Blend Coffee, Decaffeinated Coffee & Fairmont Signature Teas

Sunday, January 3, 2016	
Registration	
4:00-5:30pm Registration Ballroom P	refunction
Monday, January 4, 2016	
Registration	
7:30-9:30am Registration Ballroom P	refunction
12:00-2:00pm Registration Ballroom P	
6:00-7:30pm Registration Ballroom P	refunction
Workshops	
9:00-12:00 Translational Bioinformatics 101	Salon 2&3
Speakers: Jessica Tenenbaum, Lewis Frey, Robert Freimuth, Su Madhavan, Joshua Denny	ıbha
9:00-12:00 Computational Approaches to Study Microbes and Microbiomes	Plaza
Speakers: Susan Holmes, Igor Zhulin Jouline, Tandy Warnow, Cl	
Myers, Yana Bromberg, Casey Greene, Iddo Friedberg, Fiona Br	
Anne Hoen, Dan Knights, Craig Lessard, Serghei Mangul, James	
Nam-phuong Nguyen	,
Break	
12:00-1:30 Lunch on own	
Workshop	
·	Salon 2&3
Life-Long Precision Medicine	
1:30 Introduction and welcome	
1:40 Robert Nussbaum, Newborn screening and genome sequencing	
2:10 Stephen Kingsmore, Integrating deep phenotyping and genome sequen	ncing to
enable precision medicine in neonatal intensive care units	· ·
3:00 Jennifer Puck, Using newborn dried blood spots to obtain deep sequen enhanced, early diagnosis	nce data for
3:30 Steven E. Brenner, <i>Analysis challenges of newborn genome sequence</i>	10
4:00 Sean Mooney, <i>Understanding the complex genetics of simple Mendelia</i>	
1:30-4:30 Biomedical Applications of Topology and Abstract Algebras	Plaza
Speakers: Eric Neumann, Svetlana Lockwood	Гіада
Break	
4:30-7:30 Dinner on own	
	Kilohana
7:30-8:30pm 21 <sup>st</sup> Anniversary Reception with Drinks and Desserts	

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Tuesuay, j	anuary 3, 2010			
Registration				
7:30-9:00	Registration	Ballroom	Prefunction	
10:30-11:30	Registration	Ballroom	Prefunction	
1:00-2:00	Registration	Ballroom	Prefunction	
Breakfast		Ballroor	n Courtyard	
7:30-8:30	PSB Breakfast			
Welcome			Salon 2&3	
8:30-8:40	Welcome			
Session 1:	Innovative Approaches to Combining Genotype, Phe	notype,	Salon 2&3	
	Epigenetic, and Exposure Data for Precision Diagnos	stics		
8:40-8:50	Introduction			
	Melissa Haendel, Nicole Washington, Maricel Kann			
8:50-9:10	Invited Talk: Elissa Chesler			
	GeneWeaver: Finding relations among genes, pheno	otypes, and	l diseases	
	from heterogeneous functional genomics data		doodoo	
	Functional genomic studies of disease now encompa			
	of research resulting in a large volume of diverse ass and their relations to the biological concepts reflected			
	paradigms, diagnostic criteria and other disease rela			
	GeneWeaver system is a platform for harmonizing a	•		
	resulting gene lists, gene networks and descriptive m	•	•	
	data-driven discovery of relations among experiment			
	disease through user guided interrogation backed by			
	graph algorithms and dynamic visualization of results			
	system, users can find convergent evidence from mu			
	experimental paradigms of a role for genes in diseas			
	align individual gene centric data sets to the databas	•		
	data, identify model organism phenotypes and specif		•	
	onto various facets of human disease, and classify d		•	
	based on their intrinsic biological similarity rather tha		•	
	characteristics.			
9:10-9:25	Discovering Patient Phenotypes Using Generalized L	Low Rank I	Models	
	Alejandro Schuler, Vincent Liu, Joe Wan, Alison Callahan, Madeleine			
	Udell, David E. Stark, Nigam H. Shah			
9:25-9:40	Diagnosis-Guided Method for Identifying Multi-Modal	lity Neuroir	naging	
	Biomarkers Associated with Genetic Risk Factors in	Alzheimer	s Disease	
	Xiaoke Hao, Jingwen Yan, Xiaohui Yao, Shannon L.	Risacher,	Andrew J.	
	Saykin, Daoqiang Zhang, Li Shen, for the ADNI			
9:40-9:55	Metabolomics Differential Correlation Network Analysis	sis of Oste	oarthritis	
	Ting Hu, Weidong Zhang, Zhaozhi Fan, Guang Sun,	Sergei Lik	hodi,	
	Edward Randell, Guangju Zhai			
9:55-10:10	Integrating Clinical Laboratory Measures and ICD-9	Code Diag	noses in	
	Phenome-Wide Association Studies			
	Anurag Verma, Joseph B. Leader, Shefali S. Verma,	Alex Frase	e, John	

Please wear your PSB name tag and lanyard for admission to all events and hosted meals.

#### Tuesday, January 5, 2016 (con't)

Tuesday	
10:10-10:25	Wallace, Scott Dudek, Daniel R. Lavage, Cristopher V. Van Hout, Frederick E. Dewey, John Penn, Alex Lopez, John D. Overton, David J. Carey, David H. Ledbetter, H. Lester Kirchner, Marylyn D. Ritchie, Sarah A. Pendergrass  Investigating the Importance of Anatomical Homology for Cross-Species Phenotype Comparisons Using Semantic Similarity  Prashanti Manda, Christopher J. Mungall, James P. Balhoff, Hilmar Lapp, Todd J. Vision
Break 10:25-10:50	
Keynote 10:50-12:00	Salon 2&3 Creating a Gene X Phenome Catalog through Integration of Genome, Transcriptome, and EHR Data Nancy Cox, with introduction by Russ Altman
Lunch	Ballroom Courtyard
12:00-1:15	PSB Hosted Lunch
Session 2:	Discovery of Molecularly Targeted Therapies Salon 2&3
1:30-1:40	Introduction
1:40-2:00	Philip R.O. Payne, Kun Huang, Nigam Shah Invited Talk: Scott Kahn
	Balancing Short Term Dx with Longer Term Knowledge Creation Paradoxically it seems as though the immediate needs to diagnose and treat a patient using genomic information can be at the expense of any longer term effort to aggregate such patient data for subsequent analysis and knowledge creation. This presentation will offer a different perspective that relies upon an open architecture in which patient reporting, cohort aggregation and analysis, and an extensible knowledgebase are all integrated with one another, and ultimately integrated within a healthcare organization's information infrastructure.
2:00-2:15	Collective Pairwise Classification for Multi-Way Analysis of Disease and Drug Data Marinka Zitnik, Blaz Zupan
2:15-2:30	Computing Therapy for Precision Medicine: Collaborative Filtering Integrates and Predicts Multi-entity Interactions Sam Regenbogen, Angela D. Wilkins, Olivier Lichtarge
2:30-2:45	An Integrated Network Approach to Identifying Biological Pathways and Environmental Exposure Interactions in Complex Diseases Christian Darabos, Jingya Qiu, Jason H. Moore
2:45-3:00	Integrating Genetic and Structural Data on Human Protein Kinome in Network-Based Modeling of Kinase Sensitivities and Resistance to Targeted and Personalized Anticancer Drugs Gennady M. Verkhivker

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#### Tuesday, January 5, 2016 (con't)

3:00-3:15	A Framework for Attribute-Based Community Detection with Applications to Integrated Functional Genomics Han Yu, Rachael Hageman Blair
Discussion S	Sessions
3:15-4:15	Discovery of Molecularly Targeted Therapies
4:15-5:15	Innovative Approaches to Combining Genotype, Phenotype, Epigenetic, and Exposure Data for Precision Diagnostics
Break	
5:30	Dinner on own

	Wednesda	ay,	anuary	6	<b>2016</b>
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Wednesda	y, january 0, 2010		
Registration 7:30-8:30	Registration	Ballroom Prefunctio	n
Breakfast	Registration	Ballroom Courtyan	_
7:30-8:30	PSB Breakfast	Balliooni Courtyan	u
Announceme		Salon 2&	<u>~</u>
8:30-8:40	Morning Announcements –	Salon 20	J
Session 3:	Precision Medicine: Data and Discovery for Improved	d Salon 2&	3
00000011 0.	Health and Therapy	u Gaion Za	Ŭ
8:40-8:45	Introduction		
	Bruce Aronow, Steven Brenner, Sean Mooney, Alexa	ander Morgan	
8:45-9:20	Invited Talk: Jennifer Doudna	-	
	CRISPR Biology: A New Era in Genome Engineering		
	The advent of facile genome engineering using the b		
	CRISPR-Cas9 system in animals and plants is trans-		
	present a brief history of CRISPR biology from its init	•	
	the elucidation of the CRISPR-Cas9 enzyme mechan	• •	
	foundation for remarkable developments using this to		
	regulate or visualize genomic loci in a wide variety of		•
	These results highlight a new era in which genomic r	•	
	longer a bottleneck to experiments, paving the way to discoveries in biology, with applications in all branches		
	and strategies for human therapeutics. Recent result		
	molecular mechanism of Cas9 and its use for targete	•	
	therapies will be discussed.	od com bacca	
9:20-9:35	Multitask Feature Selection with Task Descriptors		
	Victor Bellon, Veronique Stoven, Chloe-Agathe Azer	ncott	
9:35-9:50	Knowledge Driven Binning and PheWAS Analysis in		
	Personalized Medicine Research Project Using BioB	Bin	
	Anna O. Basile, John R. Wallace, Peggy Peissig, Ca	therine A. McCarty,	
	Murray Brilliant, Marylyn D. Ritchie		
9:50-10:05	Biofilter as a Functional Annotation Pipeline for Com	mon and Rare Copy	
	Number Burden		
	Dokyoon Kim, Anastasia Lucas, Joseph Glessner, S		
	Bradford, Ruowang Li, Alex T. Frase, Hakon Hakona Murray Brilliant, Marylyn D. Ritchie	arson, Peggy Peissig,	
10:05 10:20	Separating the Causes and Consequences in Disease	so Transcriptomo	
10.03-10.20	Yong Fuga Li, Fuxiao Xin, and Russ B. Altman	se Transcriptome	
10:20-10:35	The Challenges in Using Electronic Health Records	for	
10.20 10.00	Pharmacogenomics and Precision Medicine Research		
	Sarah M. Laper, Nicole A. Restrepo, Dana C. Crawfo		
10:35-10:50	Kidney Disease Genetics and the Importance of Dive		
	Medicine	-	
	Jessica N. Cooke Bailey, Sarah Wilson, Kristin Brow	n-Gentry, Robert	
	Goodloe, Dana C. Crawford		

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#### Wednesday, January 6, 2016 (con't)

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Personalized Hypothesis Tests for Detecting Medication Response in
Parkinson Disease Patients Using iPhone Sensor Data
Elias Chaibub Neto, Brian M. Bot, Thanneer Perumal, Larsson Omberg,
Justin Guinney, Mike Kellen, Arno Klein, Stephen H. Friend, Andrew D.
Trister
PRISM: A Data-Driven Platform for Monitoring Mental Health
Maulik R. Kamdar, Michelle J. Wu
Patient-Specific Data Fusion for Cancer Stratification and Personalised
Treatment
Vladimir Gligorijevic, <u>Noel Malod-Dognin</u> , Natasa Przulj
Bayesian Biclustering for Patient Stratification
Sahand Khakabimamaghani, Martin Ester
Predicting Significance of Unknown Variants in Glial Tumors Through
Sub-Class Enrichment
Alex M. Fichtenholtz, Nicholas D. Camarda, Eric K. Neumann
One-Class Detection of Cell States in Tumor Subtypes
Artem Sokolov, Evan O. Paull, Joshua M. Stuart
on Lunch Salon 1 and Ballroom Courtyard
Poster Session with Lunch
(Posters on ODD numbered boards presented from 12:30-1:30)
(Posters on EVEN numbered boards presented from 1:30-2:30)
ession Salon 2&3
Precision Medicine: Data and Discovery for Improved Health and Therapy
Meeting Salon 2&3
ISCB Open Meeting
Dinner and evening on own

Thursday, Janua	iry 7	, Z	U16	)
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	January 7, 2010	Dellas and Occurt cond
Breakfast 7:30-8:30	PSB Breakfast	Ballroom Courtyard
		Colon 202
Announceme 8:30-8:40	Morning Announcements	Salon 2&3
Session 4:	Social Media Mining for Public Health Monitoring and	Salon 2&3
36881011 4.	Surveillance	3al011 203
8:40-8:50	Introduction	
0.40-0.00	Graciela Gonzalez, Matthew Scotch, Karen Smith, Jol	nn Brownstein
	Abeed Sarker, Michael Paul, Azadeh Nikfarjam	in Brownstoin,
8:50-9:10	Invited Speaker: Abeed Sarker	
	Social media mining for pharmacovigilance challeng	es and progress
	Social media is increasingly being used for public hea	Ith monitoring tasks
	and the trend is likely to continue because of the rapid	•
	this domain. This talk will focus on the DIEGO lab's ef	
	mining for pharmacovigilance, and how the technique	
	learned can be used for performing related public hea	Ith monitoring tasks
0.40 0.25	from social media.	Casial Madia
9:10-9:25	Towards Early Discovery of Salient Health Threats: A Emotion Classification Technique	Social Media
	Bahadorreza Ofoghi, Meghan Mann, <u>Karin Verspoor</u>	
9:25-9:40	Monitoring Potential Drug Interactions and Reactions	via Network Analysis
0.20 0.10	of Instagram User Timelines	via rvotivorit rinaryolo
	Rion Brattig Correia, Lang Li, Luis M. Rocha	
9:40-9:55	Text Classification for Automatic Detection of E-Cigare	ette Use and Use for
	Smoking Cessation from Twitter: A Feasibility Pilot	
	Yin Aphinyanaphongs, Armine Lulejian, Duncan Penfo	old Brown, Richard
	Bonneau, Paul Krebs	
9:55-10:10	Insights from Machine-Learned Diet Success Prediction	on
10.10 10.05	Ingmar Weber, Palakorn Achananuparp	va of Cooled Madia
10:10-10:25	Predicting Individual Well-Being Through the Languag H. Andrew Schwartz, Maarten Sap, Margaret L. Kern,	
	Eichstaedt, Adam Kapelner, Megha Agrawal, Eduardo	
	Dziurzynski, Gregory Park, David Stillwell, Michal Kos	
	Seligman, Lyle H. Ungar	mora, maran E.i .
10:25-10:40	Finding Potentially Unsafe Nutritional Supplements fro	om User Reviews
	with Topic Modeling	
	Ryan Sullivan, Abeed Sarker, Karen O'Connor, Aman	da Goodin, Mark
	Karlsrud, Graciela Gonzalez	
Discussion S		Salon 2&3
	Social Media Mining for Public Health Monitoring and	Surveillance
	Regulatory RNA	
Break		
12:45-2:00	Lunch on Own	

#### Thursday, January 7, 2016 (con't)

NIH Open N	Meeting Salon 2&3
2:00-3:30	
	A SWOT Analysis of Big Data @ the NIH
	Philip E. Bourne, Associate Director for Data Science, National Institutes of Health
	Funding Computational Research at NIGMS, and the New MIRA Program Rochelle Long, Acting Director, PPBC Division, NIGMS, NIH Introduction to NLM Career Development Awards and Research Grants in Biomedical Informatics
	Jane Ye, Program Officer, Division of Extramural Programs, NLM, NIH
Break 3:30-5:30	
Dinner	Coconut Grov
5:30-7:30	Dinner Party
Keynote:	Salon 2&
7:30-8:45	Public Health Information in Social Networks Winter Mason, with introduction by Marylyn Ritchie

Friday, ]	anuary	8	, 20	116
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Breakfast   7:30-8:30   Breakfast   Ballroom Courtyard   7:30-8:30   Breakfast   Salon 2&3   Salon 2&3   Salon 2&3   Salon 2&3   Closing Announcements   Methods to Enhance the Reproducibility of Precision Medicine   Salon 2&3   Introduction   Arjun Manrai, Chirag Patel, Nils Gehlenborg, Nicholas Tatonetti, John loannidis, Isaac Kohane   Dynamically Evolving Clinical Practices and Implications for Predicting   Medical Decisions   Jonathan H. Chen, Mary K. Goldstein, Steven M. Asch, Russ B. Altman   Identification of Questionable Exclusion Criteria in Mental Disorder Clinical Trials Using a Medical Encyclopedia   Handong Ma, Chunhua Weng   Repurposing Germline Exomes of the Cancer Genome Atlas Demands a Cautious Approach and Sample-Specific Variant Filtering   Amanda Koire, Panagiotis Katsonis, Olivier Lichtarge   9:35-9:50   Reproducible Research Workflow in R for the Analysis of Personalized Human Microbiome Data   Benjamin Callahan, Diana Proctor, David Relman, Julia Fukuyama, Susan   Holmes   Reproducible and Shareable Quantifications of Pathogenicity   Arjun K. Manrai, Brice L. Wang, Chirag J. Patel, Isaac S. Kohane   Discussion   Salon 2&3   Reproducible and Shareable Quantifications of Pathogenicity   Arjun K. Manrai, Brice L. Wang, Chirag J. Patel, Isaac S. Kohane   Discussion   Social Media Mining Shared Task Workshop   Salon 2&3   Social Media Mining Shared Task Workshop   This workshop was designed to act as a platform for the application of state-of-the-art natural language processing systems on social media data. Using the DIEGO Lab Adverse Drug Reaction Twitter corpus, we prepared a shared task focusing on text classification and information extraction.   Eleven teams participated in the shared task and several teams will be presenting their systems at the workshop. This workshop complements the Social Media Mining for Public Health Monitoring and Surveillance   PSB oversee vou next vear!	riiuay, jai	luary 8, 2010	
8:30-8:40 Closing Announcements  Session 7: Methods to Enhance the Reproducibility of Precision Medicine Salon 2&3 Introduction Arjun Manrai, Chirag Patel, Nils Gehlenborg, Nicholas Tatonetti, John loannidis, Isaac Kohane Dynamically Evolving Clinical Practices and Implications for Predicting Medical Decisions Jonathan H. Chen, Mary K. Goldstein, Steven M. Asch, Russ B. Altman Identification of Questionable Exclusion Criteria in Mental Disorder Clinical Trials Using a Medical Encyclopedia Handong Ma, Chunhua Weng Repurposing Germline Exomes of the Cancer Genome Atlas Demands a Cautious Approach and Sample-Specific Variant Filtering Amanda Koire, Panagiotis Katsonis, Olivier Lichtarge 9:35-9:50 Reproducible Research Workflow in R for the Analysis of Personalized Human Microbiome Data Benjamin Callahan, Diana Proctor, David Relman, Julia Fukuyama, Susan Holmes  9:50-10:05 Reproducible and Shareable Quantifications of Pathogenicity Arjun K. Manrai, Brice L. Wang, Chirag J. Patel, Isaac S. Kohane  Discussion Session Salon 2&3 10:05-11:05 Methods to Enhance the Reproducibility of Precision Medicine  Break 11:05-12:30 Lunch on own  Workshop 12:30-3:30 Social Media Mining Shared Task Workshop This workshop was designed to act as a platform for the application of state-of-the-art natural language processing systems on social media data. Using the DIEGO Lab Adverse Drug Reaction Twitter corpus, we prepared a shared task focusing on text classification and information extraction. Eleven teams participated in the shared task and several teams will be presenting their systems at the workshop. This workshop complements the Social Media Mining for Public Health Monitoring and Surveillance session.			Courtyard
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### Call for Session Proposals Pacific Symposium on Biocomputing

The Big Island of Hawaii, January 4-8, 2017

This is an invitation to submit proposals for sessions in the 22<sup>nd</sup> Pacific Symposium on Biocomputing (PSB), to be held January 4-8, 2017 on the Kohala Coast, the Big Island of Hawaii. PSB will bring together top researchers from the US, the Asian Pacific nations and around the world to exchange research for the presentation of work in databases, algorithms, interfaces, natural language processing, modeling and other computational methods, as applied to biological problems, with emphasis on applications in data-rich areas of molecular biology.

PSB has been designed to be responsive to the need for critical mass in sub-disciplines within biocomputing. For that reason, it is the only meeting whose sessions are defined dynamically each year in response to specific proposals. PSB sessions are organized by leaders in the emerging areas, and targeted to provide a forum for publication and discussion of research in biocomputing's "hot topics." In this way, PSB provides an early forum for serious examination of emerging methods and approaches in this rapidly changing field.

Proposals for such sessions are hereby solicited. A proposal should identify a coherent topic that can be addressed by 3 to 12 papers. For example, a session might bring together papers on alternative approaches to a particular biological question or it might examine the applications of a particular technology in a variety of biological areas. A sample of more than 20 different decisions at previous PSB meetings have included:

- Discovery of Molecularly Targeted Therapies
- Innovative Approaches to Combining Genotype, Phenotype, Epigenetic, and Exposure Data for Precision Diagnostics
- Methods to Enhance the Reproducibility of Precision Medicine
- Precision Medicine: Data and Discovery for Improved Health and Therapy
- Regulatory RNA
- Social Media Mining for Public Health Monitoring and Surveillance

A major goal of PSB is to create productive interaction among the rather different research cultures of computer science and biology. Recognizing that many in the computational community believe the value of a meeting depends on the quality of its review process, every full manuscript submitted to PSB is reviewed by at least two anonymous reviewers, as well as the session organizers and meeting chairs. Accepted papers are distributed in a proceedings volume at the meeting, and are indexed in Medline and other bibliographic databases. However, we also recognize that many potential participants, particularly from the biological community, do not choose to submit full papers to conferences, even with assurance of availability and broad indexing. Therefore, PSB also includes a separate open submission abstract book and open poster presentations in which any registrant can present research.

#### Responsibilities of a session chair

As a proposer of an accepted session, you will become the session chair. The chair's primary responsibility is to solicit high quality papers for the session, oversee the review, and write a tutorial introduction to your session. You are to solicit manuscripts, have them refereed, collaborate with the conference chairs in determining which manuscripts are to be accepted and structure presentation time in your session. Please contact us at the address below if you have any questions about the obligations entailed in becoming a session organizer.

### Call for Session Proposals Pacific Symposium on Biocomputing

The Big Island of Hawaii, January 4-8, 2017

#### Procedure for submitting proposals

Session proposals can be as short as a single page and should in no case be more than 6 pages. The proposal should:

- Define a specific technical area to be covered.
- Justify why the proposed area is appropriate for PSB. Discuss why the topic is timely and important, and how the topic has been addressed in other conferences or recent publications.
- Argue that there is likely to be sufficient high quality, unpublished material to fill the
  session, e.g., a list of researchers you intend to solicit for papers. If there are unlikely to
  be adequate paper submissions, consider applying for a PSB workshop instead.
- Provide a short autobiographical sketch and an explicit statement that your organization endorses your involvement.
- Proposals must be submitted online at <a href="http://psb.wufoo.com/forms/psb-session-proposal-submission-form/">http://psb.wufoo.com/forms/psb-session-proposal-submission-form/</a>.

#### **Important Dates**

February, 15, 2016 Proposals Due

March 15, 2016 Notification Regarding the Proposals

Each proposal will be evaluated by the organizing committee. We look forward to your submissions.

#### Questions? Please send email to

#### Lawrence Hunter, Ph.D.

Director, Center for Computational Pharmacology & Computational Bioscience Program Professor of Pharmacology (Denver) & Computer Science (Boulder)

E-mail: larry.hunter@ucdenver.edu

### Call for Workshop Proposals Pacific Symposium on Biocomputing

The Big Island of Hawaii, January 4-8, 2017

This is an invitation to submit proposals for workshops at the 22nd Pacific Symposium on Biocomputing (PSB), to be held January 4-8, 2017 on the Kohala Coast, the Big Island of Hawaii. PSB will bring together top researchers from the US, the Asian Pacific nations and around the world to exchange research for the presentation of work in databases, algorithms, interfaces, natural language processing, modeling and other computational methods, as applied to biological problems, with emphasis on applications in data-rich areas of molecular biology.

We are soliciting proposals for three hour workshops appropriate to the diverse PSB conference community. Workshops can include: panels of experts on emerging areas of computational biology that are not likely to attract the number of paper submissions required for a session; open discussions of topics important to the computational biology community; or other topics and formats. Workshops at previous PSB meetings have included:

- Computational Biology in the Cloud: Methods and New Insights from Computing at Scale
- Computational Challenges of Mass Phenotyping
- Computational Identification and Functional Analysis of Non-Coding RNAs
- The Future of Genome-Based Medicine
- Modeling Cell Heterogeneity: from Single-Cell Variations to Mixed Cells Populations
- Training the Next Generation of Quantitative Biologists
- Uncovering the Etiology of Autism Spectrum Disorders: Genomics, Bioinformatics, Environment, Data Collection and Exploration, and Future Possibilities

#### Responsibilities of a workshop chair

As a proposer of an accepted workshop, you will become the workshop chair. The chair's primary responsibility is to organize and preside over the presentations related to your workshop.

#### **Procedure for submitting proposals**

Proposals can be as short as a single page and should in no case be more than 6 pages. The proposal should:

- Define a specific technical area to be covered, and a format for the workshop.
- Justify why the proposed area is appropriate for PSB. Discuss why the topic is timely and important, and how the topic has been addressed in other conferences or recent publications. Describe why the proposed format is appropriate for the workshop.

### Call for Workshop Proposals Pacific Symposium on Biocomputing

The Big Island of Hawaii, January 4-8, 2017

#### **Procedure for submitting proposals (continued)**

- If the workshop proposes invited experts, provide evidence that the experts will attend the workshop. Participants are required to register for the conference, and, unlike sessions, no financial support from the conference is offered to workshop organizers. Emails from potential participants are sufficient.
- Provide a short autobiographical sketch and an explicit statement that your organization endorses your involvement.
- Proposals must be submitted online at <a href="https://psb.wufoo.com/forms/psb-workshop-proposal-submission-form/">https://psb.wufoo.com/forms/psb-workshop-proposal-submission-form/</a>.

#### **Important Dates**

February 13, 2016 Proposals Due March 13, 2016 Notification Regarding the Proposals

Each proposal will be evaluated by the organizing committee. We look forward to your submissions.

#### Questions? Please send email to

#### Lawrence Hunter, Ph.D.

Director, Center for Computational Pharmacology & Computational Bioscience Program Professor of Pharmacology (Denver) & Computer Science (Boulder)

E-mail: larry.hunter@ucdenver.edu

#### **Pacific Symposium on Biocomputing 2016**

# Abstract Book Table of Contents and Index Only

The full abstract book is available at <a href="http://psb.stanford.edu/conference-materials/">http://psb.stanford.edu/conference-materials/</a> and in print at the PSB Registration Desk.

**Poster Presenters:** Poster space is assigned by abstract page number. Please find the page that your abstract is on and put your poster on the poster board with the corresponding number (e.g., if your abstract is on page 50, put your poster on board #50).

Proceedings papers with oral presentations #11-48 are not assigned poster space.

Papers are organized by session and then the last name of the first author.

Presenting authors' names are underlined in the Table of Contents and in the list of authors on the abstracts.

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