

2011 Sage Bionetworks Commons Congress Participants and Community of Interest Workgroups (13 April 2011)

A. Engaged Public	B. Map Building	C. Federation	D. Enabling Sharing	E. Compute Platform
How can the public become more actively engaged in building better maps of disease? New roles for citizens and patients in building disease models and encouraging sharing Leaders: Lara Mangravite & Sharon Terry	Where are the best disease models? Who is building them? What are the next directions? Leaders: Jonathan Derry & Eric Schadt	What really is the Federation? How has it worked? How do you plan to expand it? Leaders: Justin Guinney & Andrea Califano	Moving away from current practices of collaboration: Draft key documents that can allow unlocking of currently inaccessible data sources Leaders: John Wilbanks, Robert Cook-Deegan, Jonathan Izant	What is the computational platform you are building? What is needed next? How can I help build it and how can I use it? What can I do to help the platform serve its best role for the community? Leaders: Mike Kellen & Martin Morgan
Samuel Aparicio UBC/BC Cancer Agency Linda Avey Brainstorm Research Kristin Baxter Genetic Alliance Greg Biggers Genomera	Beth Anderson Arkitek Studios Tatsuya Andou Takeda Doug Bassett Ingenuity Tim Behrens Genentech, Inc.	Pankaj Agarwal Glaxosmithkline Mariano Alvarez Columbia U. Mukesh Bansal Columbia U. Adam Bly Seed	Craig Alexander Howard Hughes Medical Institute Nick Anderson U. Washington Misha Angrist Duke Rachel Ankeny U. Adelaide	Saeid Akhtari NextBio Brandon Allgood Numerate, Inc Anca Bucur Philips Research Kenneth H. Buetow NIH Barry Bunin Collaborative Drug Discovery (CDD), Inc. Richard Cave Public Library of Science
Hans Bitter F. Hoffmann-La Roche Ltd Robi Blumenstein CHDI Foundation	Jason Bobe PersonalGenomes.org Robert Bruggner Stanford	Chas Bountra Oxford / SGC Philip Bourne UCSF	Orli Bahcall Nature Genetics Joffe Baker Genomic Health	Jim Davies University of Oxford Nicole Deflaux Sage Bionetworks
Sylvie Bove Lund University Russell Bromley Translational Research Acceleration Consulting Ben Carollo National Cancer Institute Adam Clark FasterCures	David Burdick Sage Bionetworks Nathan Caffo Presage Biosciences John Castle Novartis Vivien Chan Novartis	Atul Butte Stanford University Carlos Caldas Cambridge Research Institute Andrea Califano Columbia U. Patricia Cooper California Biotechnology Foundation Xudong Dai Sage bionetworks	Jeroen Belien VUmc, dept of Pathology Adam Berger Institute of Medicine Chris Cain Biocentury Bob Cook-Deegan Duke Monica Duke UKOLN, University of Bath, UK Aled Edwards SGC	Aris Floratos Columbia U. David Fore Lyba Matt Furia Sage Bionetworks
Maureen Cronin Foundation Medicine, Inc David Duncan Fortune/NY Times/UC Berkeley	Paul Chapman Takeda Tim Clarke Harvard	Janusz Dutkowskj UC San Diego	Kelly Edwards U. Washington	Robert Gentleman Genentech
Debra Fumagalli	Jonathan Derry Sage Bionetworks	Stephen Edwards U.S. EPA	Lorraine Egan Damon Runyon Cancer Research Foundation Michael Eisen PLoS	David Heckerman Microsoft Research John Hill Sage Bionetworks Colin Hill Via Science
Monohar Furtado Life Technologies Farah Gerdes Wilson Sonsini Goodrich & Rosati Lisa Green Creative Commons	Joel Dudley Stanford University Keith Elliston CHDI Foundation Tomoyuki Fujisawa Takeda David Haussler UCSC Yudong He Amgen	Dan Galloway NCI-ICBP Chris Gaterj Sage Bionetworks	Thomas Freeman boehringer-ingenheim Thomas Goetz Wired Mark Hertle Life Science Discovery Fund Liz Horn Genetic Alliance	Alex Hodgson 1DegreeBio Liang Hoe Apple Bruce Hoff Sage Bionetworks
James Gusella Massachusetts General Hospital Jeff Cloudera Hammerbacher Marc Hurlbert Avon Foundation for Women Peter Kapitein Inspire2Live	Tianxiao Huan Sage bionetworks Erich Huang Sage Bionetworks Jos Jonkers Netherlands Cancer Institute Peter Kang Stanford University	Josh Greenberg Alfred P. Sloan Foundation Justin Guinney Sage Bionetworks Gregory Hannum UC San Diego Trey Ideker UCSF Daphne Koller Stanford University Teri Melese UCSF	Iain Hrynaskiewicz BioMed Central Joseph Jackson Open Science Summit Vitalina Komashko Sage Bionetworks Kelly LeMarco AAAS/Science Translational Medicine Clifford Lynch Coalition for Networked Info.	Michael Kamenick Recombinant data Michael Kellen Sage Bionetworks Tom Krohn Eli Lilly Allan Kuchinsky Agilent Technologies
Lara Mangravite Sage Bionetworks Adam Margolin Broad Institute Brad Miller Helices Simon Noble CHDI Foundation Thea Norman Arch2POCM	Andrew Kasarskis Pac Bio Hiroaki Kitano SBI & OIST Ron Krauss CHORI John Lamb Pfizer Paul Lasko Canadian Institutes of Health Research Jill Loftiss GSK	Quingying Meng Sage Bionetworks Antonina Mitrofanova Columbia U. David Nicholson Merck Research Laboratories Vern Norvick Wilson Sonsini Goodrich & Rosati Keiichiro Ono UC San Diego Bengt Persson	Brig Mecham Sage Bionetworks Toshiyuki Mori Takeda Bernard Munos InnoThink Andy Oram O'Reilly Media Mette Peters Sage Bionetworks Nancy Shepard Medco Leming Shi US FDA Sandra Silberman Quintiles, Inc.	Ilya Kupersmidt NextBio Greg Leibfreid J&J PRD Peter Li Univ. Birmingham Michael Linderman Stanford University Kurt Messersmith Amazon
Jane Reese-Coulbourne Reagan-Udall Foundation Scott Sacane Catalytic Data Streams Laura Shawver The Clarity Foundation Robert Shelton Private Access, Inc.	Marcy MacDonald MGH/HMS Minoru Masago Takeda Gerrit Meijer VU university medical center Sandy Miller Kauffman Foundation	Karen Sachs Stanford University Vicki Seyfert-Margolis FDA William Shin Columbia U. MacKenzie Smith MIT Libraries	Peter Speyer IHME Marty Tenenbaum CollanRx Paul Uhliir National Academy of Sciences Laura Van t Veer UCSF John Wilbanks Creative Commons	Martin Morgan Fred Hutchinson Cancer Research Center Kimberly Myers NCI Eric Neumann Clinical Semantics
Michael Stocum Quintiles Christine Suver Sage Bionetworks Sharon Terry Genetic Alliance	Barend Mons Netherlands Bioinformatics Centre Spyro Mousseis TGEN Amanda Myers University of Miami	Michael Snyder Stanford University Rob Tirrell Stanford Lex Vander Ploeg Abraxis Zhi Wang Sage Bionetworks Xia Yang Sage Bionetworks	Bin Zhang Sage Bionetworks	Tom Pollard The British Library & DataCite Mohammed Rahman Novartis Michael Reich Broad Institute
Luke Timmerman Xconomy Erik van Veenendaal Inspire2Live	Gaurav Pandey UC Berkeley Dexter Pratt Selventa Paul Rejto Pfizer	Judy Zhong Roche GROUP F Stephen Friend Sage Bionetworks Alyssa Goodman Harvard Jonathan Izant Sage Bionetworks Peter Klein Griffith Television Christian Perry Abby Schwartz Harvard		Philippe Rocca-Serra University of Oxford Susanna-Assunta Sansone BioSharing
Keith Yamamoto UCSF Jun Zhu Sage Bionetworks	Michel Roberge Centre for Drug R&D Chris Roberts Merck Anders Kamal Saini Sage Bionetworks Eric Schadt Pac Bio David Shaywitz Theravance Solveig Sieberts Sage Bionetworks Taylor Sittler UCSF Andreas Sundquist OnaNexus Hiro Yoshi Takeda Toyoshi Andrew Trister Univ of Washington Fred W. Turek Northwestern Univ	Regina Warmoth Sage Bionetworks		Xavier Schildwacher Sage Bionetworks Nigam Shah Stanford University Todd Smith Geospiza Hilary Srere Apple Jessie Tenenbaum Duke Univ. Kaitlin Thaney Digital Science Jeremy Thompson Novartis Jeff Trent TGEN Nicholas Tsinoremas University fo Miami Martijn van Iersel Maastricht University, the Netherlands Lalit Verma Eli Lilly and Company
	Michael Wellems Northwestern Univ			Hans-Martin Will Microsoft Bryn Williams-Jones Pfizer Worldwide R&D