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| **NAME** | **AFFILIATION** | **REASON FOR ATTENDING/INTEREST IN BIOCOMPUTE STANDARDS** |
| Jonathon Keeney (Secretary) | George Washington University | I'm spearheading a public private partnership effort around BioCompute, and would like to formalize the standard through the IEEE process. |
| Carole Goble | The University of Manchester, UK  ELIXIR, BioExcel and ResearchObject.org | Co-lead the Research Object initiative, responsible for driving CWL and BCO related activities for ELIXIR (the EU Research Infrastructure for Life Science Data).  ExCo lead of ELIXIR Interoperability Platform and Head of Node for ELIXIR-UK Node.  Been involved in BioCompute since 2016 and one of the authors of the BCO paper/arxiv paper.  (I’ll also be presenting BCO at the Cancer Early Detection meeting in Portland Oregon in October). Also using BCO in the EU BioExcel Centre of Excellence. ROs are also used in the NIH Data Commons projects, and Manchester are to be undertaking (subcontracted) that work. |
| Raja Mazumder | GW | BioCompute co-founder and chair of this meeting |
| Jonas Almeida | Stony Brook Univ (SUNY), Prof and CTO | Development of consumer-facing biomedical informatics applications, co-author of BCO paper with an eye on Web/Cloud orchestration. |
| Ryan Connor | NCBI [contractor] | Representing NCBI Virus group - general interest in bio-data standards |
| Theresa Wohlever (Attending as a stand-in for Jonathan Jacobs) | QIAGEN | Interested in aligning QIAGEN Bioinformatic software to the BCO collaboration standards. |

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| Hiroki Morizono | Childrens National Medical Center | Interested in application of BCO for precision medicine and reproducible data processing |
| Vahan Simonyan | CRISPR TX | BioCompute Co-founder and co-chair |
| Charles Hadley King | George Washington University | GW BioCompute technical lead |
| Jason Travis | TGen North | Biocompute for research organizations and developing standards compliant bioinformatics systems |
| Jonathan LoTempio | Children’s National Medical Center | Scripted BCO, executable BCO |
| Janisha Patel | George Washington University | BCO paper contributor, technical researcher |
| David Michaels | DNAnexus | Heavily involved in regulatory ecosystem |
| Stian Soiland-Reyes | The University of Manchester | Research Object technical lead, Common Workflow Language leadership team, W3C PROV co-author. Interested in integrations with existing community standards. |
| Jonathan Pryke | AstraZeneca / MedImmune | Application of BCO for precision medicine & data provenance |
| Paul Duncan | Merck and Co., Inc | BCO for regulated applications |
| Ruth Bandler | FDA | Initial FDA BCO COR |
| Michael R. Crusoe | Common Workflow Language project | CWL co-founder & CWL project lead, ResearchObject.org participant. To see BCO build upon and extend existing standards. |
| Carole Carey | IEEE Engineering in Medicine and Biology Society & Liaison to IEEE-SA (Standards Association) | Sponsoring Society for IEEE P2791 |
| Lee Black | Amazon Web Services | Interested in standards for genomics and precision medicine |
| Eric Donaldson | FDA |  |
| Kristy Cloyd-Warwick | DNAnexus | Interested in standards for genomics |
| Josep L. Gelpi | Barcelona Supercomputing Centre. Univ. of Barcelona. BioExcel. ELIXIR-ES | Interested in use opportunities for BCO. Representing Spanish ELIXIR Node. |
| Ben Busby | NCBI | Interested in building reproducible tools for biological data science |
| Jianchao Yao | Merck & Co | Representing Merck in the BCO consortium. Working with the FDA on a mock submission of BCO. |
| Ogan Abaan | Seven Bridges Genomics | Making the biocompute object a standard for exchanging information |
| Asa Oudes | Genedata | Interested in ensuring compatibility of Genedata solutions with IEEE standards. |
| Anton Golikov | FDA | BioCompute Co-founder and FDA technical lead |
| Amanda Bell | GWU | BCO technician and contributor |
| Soo Kim | IEEE-SA | IEEE Staff |
| Alexis Norris | FDA | Interested in standards for genomics datasets, from a regulatory perspective. |
| Megan Potterbusch | GWU | Interested in BCOs for reproducibility of scientific research |
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| Payal Banerjee | Children’s National Medical Center | BCO as reference for reproducible data processing for next generation sequencing pipelines and implementation of BCO as an executable |
| Surajit Bhattacharya | Children’s National Medical Center | BCO as reference for reproducible data processing for next generation sequencing pipelines and implementation of BCO as an executable |
| Jizu Zhi | FDA | Interested in BCO and standardizations |
| Gil Alterovitz | Harvard | To get standardized ngs workflow instances |
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Total voting members: 29

Total nonvoting members: 6