Day 1: Sunday, December 3, 2017 - Washington DC / Rockville Hilton Hotel

Mentorship Activity

Minimal Standards Working Group Convenes (Twinbrook Room)
Common Repository Working Group Convenes (Montrose Room)
Tools and Resources Working Group Convenes (Democracy Room)
Germline Gene Working Group Convenes (TBA)

Opening of AIRR Community Meeting & Presentation of Working Group Progress

Time	Activities
	Working Groups meet concurrently
9:00 - 12:00	Mentorship Activity for New Investigators: Victor Greiff, University of Oslo Steps in AIRR Data Capture and Analysis (Jackson Room)
12:00 - 12:45	Lunch (on own)
12:45 - 1:00	MEETING BEGINS: Introduction of Conference Organizers, Working Group Leaders, and Review of the Agenda (Washington Theater)
1:00 - 3:00	All Working Groups convene with conference attendees to discuss recommendations including:
1:00 - 2:00 2:00 - 3:00	Implementation Details from Minimal Standards Working Group Implementation Details from Common Repository Working Group
3:00 - 3:30	Break
3:30 - 3:50 3:50 - 4:00 4:00 - 4:15 4:15 - 4:30	Tools and Resources – Introduction and Software WG update Tools and Resources – File Formats WG update Tools and Resources – Biological Standards WG update Towards standards for TCR repertoire studies – Encarnita Marrioti- Ferrandiz
4:30 - 5:30	Presentation of Proposed Governance Structure: Discussion & Vote on Joining TAbS (The Antibody Society)
5:30 - 6:00	Reception ~ snacks provided ~ cash bar (Eisenhower Room) No government funds will be used for food and drinks

6:00 - 7:30	Recommendations from Germline Gene Working Group: AIRR Community's Role in IG Gene Naming and Its Relationship to the IUIS IG/TR/MH Nomenclature Committee
	Attendees are on their own for dinner

Day 2: Monday, December 4, 2017 – NIAID's Fishers Lane Facility Scientific Presentations: Sessions 1 – 3

Webinar Links for Virtual Attendance

Session 1: https://attendee.gotowebinar.com/register/7041796218829310979
Session 2: https://attendee.gotowebinar.com/register/7134530129027597827

Time	Activities
7:45 AM	Attendees gather in Rockville Hilton Hotel lobby to walk to NIH Fishers Lane Facility ~ arrive at facility by 8:00 to go through security
8:30 - 8:40	Introduction to the Day's Agenda (Room 1D13) Session 1: Germline Gene Inference Tools Moderator's Introduction: Andrew Collins, University of New South Wales
8:45 - 9:05	Presentation 1 - TIgGER for IGHV Gene Inference
9:05 - 9:25	Steve Kleinstein, Yale University Presentation 2 - IgDiscover for IGHV Gene Inference Martin Corcoran, Karolinska Institute
9:25 – 9:35	Shared Q&A
9:35 – 10:00	Presentation 3 – Evaluation and Validation of Germline Gene Inference Mats Ohlin, Lund University
10:00 - 10:10	Q&A
10:10 - 10:30	Panel Discussion: Validation & Comparison of Inference Tools
10:30 - 10:45	Break
10:45 - 10:55	Session 2: IGH Genomic Haplotypes Moderator's Introduction: Marie-Paule Lefranc, IMGT, University of Montpellier
10:55 - 11:20	Presentation 1 – New Human IG Haplotypes Corey Watson, University of Louisville Medical School
11:20 - 11:30	Q&A

11:30- 11:55	Presentation 2 – Role of Novel Regulatory Elements in the Formation of the Ig-κ Repertoire
11:55 - 12:05	Ann Feeney, The Scripps Research Institute Q&A
	Presentation 3 - Macaque IG Haplotypes
12:05 - 12:30	Akshaya Ramesh, University of California, San Francisco
12:30 - 12:40	Q&A
12:40 - 1:00	AIRR Community Discussion Discussion Leaders: Marie-Paule Lefranc & Corey Watson
	IGH Haplotype and Germline Gene Nomenclatures
	Boxed Lunches & Poster Session I (LD10 Lower Level)
1:00 - 1:45	
	Minimal Standards Implementation Demos
1:45 – 2:15	Steven Kleinstein, Nina Prak, Florian Rubelt, Christian Busse, & Syed Ahmad Chan Bukhari
	Session 3: IP Issues in Data Sharing across Shared Repositories
2:15 - 2:25	Moderator's Introduction:
	Tania Bubela, Simon Fraser University
	Presentation 1 - TBA
2:25 – 2:45	Harlan Robins, CEO, Adaptive Biotechnologies
	Presentation 2 - TBA on the Human Immunome Project
2:45 – 3:05	Jim Crowe, Vanderbilt University
	Break
3:05 – 3:25	
3:25 – 3:45	Presentation 3 - Intellectual Property Law
3:25 - 3:45	Jacob Sherkow, New York Law School
	Panel - Robins, Crowe and Sherkow - IP Issues in Data Sharing
3:45 - 4:30	Taner - Robins, Growe and Sherkow - It issues in Data Sharing
	Common Repository WG: Resolving IP and Protected Sequences
	Discussion and Community Vote
4:30 - 5:00	Lindsay Cowell & Corey Watson
5:00	End of Day at Fishers Lane Facility ~ Attendees return to Rockville Hilton Hotel
5:30 - 7:45	Reception and Tool Demos at the Washington DC/Rockville Hilton Hotel
0.00 7.10	Snacks provided ~ cash bar
	No government funds will be used for food and drinks

5:30 - 6:30 6:30 - 7:00	5 m Tool Demonstrations Panel on validation and improvement of AIRR analysis tools
7:00 - 7:45	Face-to-face tool demonstrations by teams
	Attendees are on their own for dinner

Day 3: Tuesday, December 5, 2017 – NIAID's Fishers Lane Facility Scientific Presentations: Sessions 4 and 5

Webinar Links for Virtual Attendance

Session 4: https://attendee.gotowebinar.com/register/2846600806957000963
Session 5: https://attendee.gotowebinar.com/register/2846600806957000963

Time	Activities
7:45 AM	Attendees gather in Washington DC / Rockville Hilton Hotel lobby to walk to NIAID's Fishers Lane Facility ~ arrive at facility by 8:00 to go through security
8:30 - 8:40	Introduction to Day 3 Agenda (Room 1D13) Session 4: AIRR Data in the Study of Disease Moderator's Introduction Steve Kleinstein, Yale University
8:40 - 9:05	Presentation 1 - Cancer Immunotherapy John O'Malley, Brigham and Women's Hospital, Boston
9:05 - 9:15	Q & A
9:15 - 9:40 9:40 - 9:50	Presentation 2 - Neoantigen-specific T-cell Receptor Repertoires from Healthy Donors Johanna Olweus, Oslo University Hospital Q & A
9:50 - 10:05	Break
10:05 - 11:30	Presentation 3 - BCR repertoire analysis in human SLE Iñaki Sanz, Emory University School of Medicine
11:30 - 11:40	Q & A
11:40 – 12:05	Presentation 4 - BCR repertoire sequencing in patients with primary immunodeficiency analysis in human SLE –
12:05 – 12:15	Johannes Truk, Oxford Vaccine Group, University of Oxford Q & A

12:15 - 2:00	Boxed Lunches and Poster Session II (LD10 Lower Level) concurrent with New Investigators' Lunch with Mentors
2:00 - 2:05	Session 5: Integration of AIRR Data with Complex Biological and Population Data Moderator's Introduction: Coorgo Coorgins University of Toyon Austin
2:05 - 2:30	George Georgiou, University of Texas – Austin Presentation 1 – Molecular Analysis of the Serum Antibody Repertoire
2:30 - 2:40	George Georgiou, University of Texas – Austin Q & A
2:40 – 3:05	Presentation 2 - Genetic Diversity of Non-Human Primate & Human Germline Ig Genes Gunilla Karlsson Hedestam, Karolinska Institute, Stockholm
3:05 - 3:15	Q & A
3:15 – 3:25	Break
3:25 – 3:50	Presentation 3 - The Antibody Response of Celiac Disease Ludvig Sollid, University of Oslo
3:50 - 4:00 4:00 - 5:00	Q & A AIRR Community Discussion Discussion Leaders: Felix Breden & Jamie Scott Proposed AIRR Community Governance Structure and Working Group Organization
5:00	End of Day at Fishers Lane Facility ~ Attendees return to Rockville Hilton Hotel
6:30 - 10:00	Attendees Convene for Group Dinner at Seasons52 in North Bethesda

Day 4: Wednesday, December 6, 2017 - NIAID's Fishers Lane Facility

Community Initiative Working Day

Time	Activities
7:45	Attendees gather in Washington DC / Rockville Hilton Hotel lobby to walk to NIAID Fishers Lane Facility ~ arrive at facility by 8:00 to go through security
8:40 - 9:30	Conclude discussion and vote on: AIRR Community Governance Structure
9:35 - 10:30	Conclude discussion and vote on: Working Group Organization
10:35 - 11:00	Conclude discussion and vote on: Germline Gene Working Group Recommendations
11:00 - 12:00	Elect AIRR Community Leadership
12:00	Full Meeting Adjourns
12:00 - 1:00	Lunch (on own)
1:00 - 4:00	Working Groups will convene at the Washington DC / Rockville Hilton Hotel to finalize membership & develop plans for the coming year (Montrose, Frederick and Twinbrook rooms)