

**Genome Sequencing Program Meeting March 2017**
**Wednesday, March 29, 2017**

<b>Start</b>	<b>ID</b>	<b>Agenda Item</b>	<b>Moderator/Speaker(s)</b>
8:30	D1S1	Introduction	Lu Wang
8:35	D1S2	University of Washington CMG	Mike Bamshad
9:05	D1S3	Yale CMG	Richard Lifton
9:35	D1S4	Broad Institute CMG	Daniel MacArthur
10:05	D1S5	Baylor-Hopkins CMG	Jim Lupski
10:35		<b>Break</b>	
10:55	D1S6	Invited Talk: Undiagnosed Diseases Network (UDN)	David Adams
11:40	D1S7	Challenges and Opportunities Baylor-Hopkins: When are LOF Mutations Not LOF Mutations Broad: CMG Usage of ClinVar and the Beacon Network Yale: Control-free analysis of Transmitted Mutation Burden UW: Analysis of de novo Variants	Baylor-Hopkins: Zeynep Hande Coban Broad: Heidi Rehm Yale: Richard Lifton UW: Tychele Turner
12:40		<b>Lunch (NHGRI Staff meeting with ESP)</b>	
2:10	D1S8	Matchmaker Exchange Experiences	Overview: Heidi Rehm Baylor-Hopkins: Nara Sobreira UW: Jessica Chong Broad: Harindra Arachchi Yale: Francesc Lopez GeneMatcher Survey Results: Ada Hamosh All: Discussion
2:55	D1S9	Data Sharing Activities	Debbie Nickerson
3:10	D1S10	Data Analysis WG Update (heterogeneous, complex phenotypes; combining phenotypes of common interests pre-pipeline; new tools; other topics)	Eric Boerwinkle
3:35	D1S11	Unsolved Cases WG Update	David Valle/ Liz Blue
4:00		<b>Break</b>	
4:15	D1S12	Status of Mendelian Disease Gene Discovery	David Valle
4:45	D1S13	ESP Feedback	ESP
5:00	D1S14	Wrap-up and Action Items	GSPCC/ Lu Wang
5:15		<b>Wednesday Meeting Adjourns</b>	
6:30		<b>Group Dinner at Redwood Restaurant &amp; Bar 7121 Bethesda Ln, Bethesda</b>	