

Hours	Monday	Tuesday	Wednesday	Thursday	Friday
07:30 -- 08:45	Breakfast	Breakfast	Breakfast	Breakfast	Breakfast
08:45 -- 09:30	Welcome words + Intro talks (3 min per person): Part 1 of 2	Plenary 2 (Nimesh Pinnamaneni, Helixworks)	Plenary 3 (Zohar Yakhini, Technion & RUNI)	Plenary 4 (Robert Grass, ETH)	Plenary 5 (Benno Schwikowski, Institut Pasteur)
09:30 -- 10:15					Potential pop-up talk
10:15 -- 10:45	Coffee	Coffee	Coffee	Coffee	Coffee
10:45 -- 11:30	Intro talks: Part 2 of 2	Breakout session (Roe Amit)	Breakout session (Roni Rak)	Group photo	Short talks3: Anina Grucia / Frederik Walter
11:30 -- 12:15		Breakout session (Eitan Yaakobi)	Breakout session (Mark Somoza)	Breakout session (Olgica Milenkovic)	
12:15 -- 13:45	Lunch	Lunch	Lunch	Lunch	Lunch
13:45 -- 15:15	Plenary 1 (Tzachi Pilpel, Weizmann Insitute)	Short talks 1: Roman Sokolovskii / Roni Rak	Excursion to Trier	Short talks 2: Zhiying Wang / Belaid Hamoum	Departure
15:15 -- 15:45	Coffee + cake	Coffee + cake		Coffee + cake	
15:45 -- 16:30	Lightning talk (advertising session chair topics: Roe, Eitan)	Active time: summary of earlier breakout sessions, lightning talk: Mark Somoza, pop-up talk, or breakout in smaller groups		Breakout (Lior Nissim)	
16:30 -- 17:15	Icebreaking activity			Active time: summary of earlier breakout sessions, pop-up talk, breakout in smaller groups	
17:15 -- 18:00	Free time	Free time		Dinner	
18:00 -- 19:30	Dinner	Dinner		Get together	
19:30 --	Get together	Get together			

Plenaries

Tzachi Pilpel: "An introduction into the genetic code"

Nimesh Pinnamaneni: "From Concept to Automation: Realising a Composite Motif-Based DNA Data Storage System"

Zohar Yakhini

Robert Grass

Benno Schwikowski: "Bivariate Monotonic Functions as Novel Models for Complex Biological Phenotypes"

Breakout sessions

Roe Amit: "Establishing a set of communal grand-challenges"

Eitan Yaakobi

Roni Rak: CRISPR Library Design

Mark Somoza (and Zohar Yakhini): CRISPR

Lior Nissim: "Synthetic promoters and synthetic gene circuits"

Olgica Milenkovic: "PacBio sequencing and the information content in the kinetics data"

Short talks							
Roman Sokolovskii: Motif-Based DNA Storage							
Roni Rak: CRISPR screen							
Zhiying Wang: "Secret Sharing for DNA Probability Vectors"							
Belaid Hamoum							
Anina Grucia: "Random access coverage depth problem"							
Frederik Walter							