Program

08:00	Bus from BMC Uppsala to Aula Magna
08:45	Registration and coffee, hanging posters
09:30	Welcome Words
	Ylva Engström, Chair of the SciLifeLab Board
	Introduction to the Conference
	Kerstin Lindblad-Toh, Chair of the Scientific Committee
09:45	Investigating mammalian evolution and human disease through comparative genomics in hundreds of species
	Elinor Karlsson, UMass Chan Medical School and Broad Institute, USA
10:15	A Million-Year-Old Journey: Exploring Mammoth Speciation and Adaptive Evolution through time
	Tom van der Valk, Swedish Museum of Natural History
10:40	Coffee break and Poster session I
11:20	Insights from high quality genome sequencing across the tree of life
	Richard Durbin, University of Cambridge, UK
11:55	Conservation genomics of the Scandinavian Arctic fox
	Karin Norén, Stockholm University
12:20	Lunch break and mingle, exhibitions
13:40	Domestication hologenomics – what are we missing without taking this approach?
	Tom Gilbert, University of Copenhagen, DK
14:15	Sequencing the supergene that governs Darwin's different forms of flowers
	Tanja Slotte, Stockholm University
14:40	Flash Talks from Junior researchers (selected from submitted abstracts)
	5-minute talk/researcher
15:05	Coffee break and Poster session II
15:45	The remarkable population structure of Atlantic and Baltic herring – selection, selection, selection
	Leif Andersson, Uppsala University

16:20 Why cyanobacterial blooms produce toxins – unravelling the underlying genetic diversity in the microcystin gene cluster

Karin Rengefors, Lund University

- 16:45 Best Poster Award ceremony Closing words
- 17:00 End of Conference

Meet the speakers and mingle; snacks and free beverages

Bus to Uppsala departs at 17:00 and at 18:30