ECA Conference, Leuven, June 29, 2025. "Hands-on", interactive workshops						Room	*Supervisor*
	Speaker	**Topic**	**Sessions**	**Attendees per session**	**What you will need**		
UCSC	**Robert Kuhn** (Former Associate Director, UCSC Genome Browser)	The use of online resources to annotate/map clinical CNVs	**1) 9:30-11:00** **2) 11:15-12:45**	30	Laptop computer	**ALO 07.100**	*Van Den Bogaert*
Illumina	**Eithan Zand Marco Russmann**	Bridging cytogenetics workflows: Hands-on analysis of arrays to NGS	**1) 9:30-11:00** **2) 11:15-12:45**	30	Laptop computer	**ALO 05.200**	*Pat Heslop-Harrison*
Bionano	**Dana Jaber**	Analysis of clinical and haemotological samples, in group of three (the computer connects directly to the Bionano US server in US).	**1) 9:30-11:00** **2) 11:15-12:45**	15	Laptop computer	**ALO 05.100**	*Jean-Michel Dupont & Amber Verhasselt*
Nanopore	**Erwin Reiling Erika Souche**	Nanopore based sequencing for structural variant detection, from sample to report. **1) wet lab** session with a maximum of 12 participants. 2) dry lab: (i) how to use Epi2me; (ii) real examples; (iii) hands on and support for troubleshooting.	**1) wet lab*** **9:30-11:00** *Participants should gather at the reception. Joris Vermesch will escort them to his lab. **2) dry lab** **11:15-12:4**	**1) 12** **2) 24**	Laptop computer	**ALO 07.200**	*Joris Vermeesch*