poster nr	Voornaam	Achternaam	Title of my poster	
1	Joanna	von Berg	What's in a name? Defining ischemic stroke phenotypes for Genome Wide Association Studies	
2	Adrien	Melquiond	Sorting out the Heroes from the Zeroes: the added value of short MD simulations to molecular docking	
3	Josephine	Daub	Genetic interactions in childhood cancer	
4	Li	Xue	Bridging artificial intelligence and physics-based docking for better modelling of biomolecular complexes	
5	Cunliang	Geng	tbd	
6	Armel	Lefebvre	Reconstructing Experiments: A Scientific Forensics Approach to Evaluate Reproducibility	
7	Anna-Lena	Lamprecht	Automated Composition of Scientific Workflows in Mass Spectrometry Based Proteomics	
8	Martijn	Molenaar	LION/web: a novel web-based tool for lipidomic data analysis	
9	David	Nieuwenhuijse	ViRNN: Short-read metagenomic sequence classification using deep learning	
10	Niels	Aerts	Elucidating the gene regulatory network that integrates plant abscisic acid and jasmonic acid signaling	
11	Christina	Papastolopoulou	Genetic interactions underlying mutational signatures in paediatric cancer	
12	Bas	Dutilh	Global phylogeography and ancient evolution of the widespread human gut virus crAssphage	
13	Jorge	Roel	Less is more: Coarse-grained protein-protein docking in HADDOCK	
14	Ksenia	Arkhipova	Expanding knowledge of gene order conservation in viral genomes	
15	Roy	Straver	Consensus calling using raw nanopore data for Cyclomics	
16	Abhinandan	Devaprasad	Profiling regulatory networks of 40 distinct immune cells reveal progenitor plasticity and complex regulation of human hematopoietic cell lineages	
17	Weiyang	Tao	Integration of transcriptomic and epigenetic profiling shows that CXCL4 induces novel gene regulatory networks and molecular pathways in dendritic cells	
18	Terry	Huisman	Shifts in microbial community composition along the wastewater pipeline used in the bioremediation of the Griftpark	
19	Nila	Servaas	Identitification of key pathways and genes dysregulated in myeloid dendritic cells from systemic sclerosis patients using co-expression network analysis	
20	Yaron	Caspi	Age-related intracranial volume — A longitudinal study	
21	Marek	Prachar	From Self-Assembling to Regenerative Medicine	
22	Sergio	Arredondo Alonso	mlplasmids: a user-friendly tool to predict plasmid- and chromosome- derived sequences for single species	
23	Jose	Espejo Valle-Inclan	Generation of a reference set of cancer structural genomic variation using an integrated long-read sequencing approach	
24	Li	Xue	iScore: a novel graph-kernel based scoring functions for computational docking	
25	Hina naz	Khan	RNA splicing of PBMCs of patients with sepsis	
26	Saheli	De	MiRNA target prediction using Deep Learning frameworks	
27	Marleen	Nieboer	The prioritization of structural variation	
28	Duong	Vu	fMLC: Fast Multi-Level Clustering and Visualization of Large Molecular Datasets	
29	Hanneke	van Deutekom	Extrapolating incomplete HLA allele database entries improves identification of matching alleles using NGS	
30	Marieke	Klein	Investigating the genetic overlap between ADHD, aggressive and antisocial behavior, and brain volumes	
31	Charlotte	van Noort	Modeling Membrane Protein Complexes with HADDOCK	
32	Joep	de Ligt	Campus wide compute infrastructure and services	
33	Brian	Jimenez-Garcia	Data-Driven Structural Bioinformatics: The HADDOCK Ecosystem	
34	Joanna	Wolthuis	MetaboShiny – identify each mass, en masse	
35	Mao	Peng	Comparative analysis of basidiomycete transcriptomes reveals a core set of expressed genes encoding plant biomass degrading enzymes	
36	Sarah	Mehrem	On what level do phages and hosts co-evolve?	