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1	Xesus Abalo	Sequencing at the National Genomics Infrastructure
2	Inci Aksoylu	Plasma profiling for protein-based stratification in ALS
3	Leon Bergner	Molecular mechanisms of membrane sculpting by the human Caveolin-1 complex
4	Hans Blom	Advanced Light Microscopy support
5	Adriana Carvajal Jimenez	High-Throughput Profiling of Transcription Factor Binding Dynamics in Response to Epigenetic Modulation
6	Kenia Chavez Ramos	Elasto-Internal Spiral Microfluids for Size-Based Particle Separation Applied in Sepsis Diagnostic
7	Arash Darzian	Selective Translation: When Length, Structure, and Nucleotides Decide
8	Sulagna Dasgupta	Context-Specific Regulatory Network Response to Transcription Factor Dosage
9	Diana Daumiller	AI design of Peptide Binders
10	Sofia Dimou	Development of a translational arrest peptide-based system to probe co-translational interactions of an ER-integral chaperone in yeas
11	Christopher Dirks	Systematic profiling of SAMHD1 nucleoside analogue interactions
12	Jelle Dylus	Examining cellular response to purine antimetabolite nelarabine to gain insight into the mechanism-of-action in T-ALL
13	Robin Ebbestad	Nanoscale Morphometrics for Stratification of Patients with Glomerulonephritis
14	Lisa Eichhorn	High-throughput Evolutionary Selection of DNA Nanostructures for Biomedical Applications
15	Alper Eroglu	Detecting cell phenotype interactions that drive tumor growth
16	José Eduardo Ferreira Araújo	Clinical Proteomics Mass Spectrometry and Global Proteomics and Proteogenomics Facility
17	Wojciech Fijalkowski	Influence of formulation buffer on structure of clinically relevant lipid nanoparticles
18	Giovanni Giglio	Cell cycle-based Replication Timing (2C-RT) to detect local and genome-wide DNA replication perturbations
19	Pauline Granit	Metal promiscuity in ancestral and modern-day terpene cyclases
20	Louise Gsell	Inferring cell interaction history from snapshot tumor data
21	Susanna Hellberg	Modeling the colon tumor microenvironment: A 3D silk-scaffold in vitro network to investigate the role of estrogen and hypoxia
22	Reba Howard	Macromolecular Dynamics, Structure, Pharmacology & AI in the Lindahl Research Team

23	Emil Johansson	The Evidence Proteome
24	Marie Lavin	IT Support Services
25	Qjuzhen Li	The Structural History of Eukarya
26	Karin Lindman	Site Support
27	Honglian Liu	Nanopore based diagnosis of phenotypic antimicrobial resistance
28	Dania Machlab	SciLifeLab Bioinformatics Platform – NBIS
29	Oliver Morusewicz	Loss of SMUG1 Sensitizes Cells to PARP Inhibitors via Enhanced PARP1 Trapping and Accelerated Replication
30	Riccardo Mosca	Decipher riboregulation in healthy and diseased cells
31	Maryam Nejati	The Hsp90 co-chaperone Sti1 is the cytoplasmic retention determinant of Stp1 latency
32	Jakub Palacka	VOLUMINEX: A new route towards 3D Spatial Transcriptomics for Infection Studies based on DNA microscopy
33	Elin Palm	Beyond pH – Ion concentration impact on lipid nanoparticle (LNP) structure along the endosomal pathway
34	Jasper Panten	Studying gene dosage response at single cell resolution using CRISPR perturbation screens
35	Philipp Rentzsch	The Developmental Genotype-Tissue Expression (dGTEx) Project
36	Leah Rosen	Studying gene dosage response at single cell resolution using CRISPR perturbation screens
37	Nora Ross	Quantitative Microscopy Reveals Differential NK Cell Activation Dynamics Across DNAM-1 Ligand
38	Johanna Sandling Johansson	KI External Engagement Office
39	Anna Schuppe	Optimizing Whole-cell Catalysis of 2,5-Furandicarboxylic acid (FDCA)
40	Doreen Schwochow	A dog is not a wolf- how the interactome has changed man's best friend
41	Maximilian Senftleben	Bioimage analysis support on multiplexed image data by SciLifeLab BIIF
42	Noora Sissala	Diagnostic plasma protein biomarkers for lung adenocarcinoma using in-depth proteomics and machine learning
43	Ronald Sjöberg	High-throughput Antigen Screening Using Protein and Peptide Arrays
44	Ana Spencer	Next-generation neurotherapeutics: integrating microfluidics and targeted nanoparticles
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50	Jen-Hung Wang	<i>Single cell image-based profiling of micro/nano plastic particle concentrations on various cell types</i>
51	Nina Warnke	<i>Towards high-throughput biophysics in protein sequence space</i>
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53	Yuqi Zheng	<i>Isotonic regression for deriving Posterior Error Probabilities for peptide identifications</i>
54	Rebecca Howard	<i>SciLifeLab Cellular & Molecular Imaging (CMI) Platform</i>
55	Rebecca Howard	<i>Diversity, Equity & Inclusion at SciLifeLab (DEIS)</i>
56	Eric Sonnhammer	<i>FunCoup 6: advancing functional association networks across species with directed links and improved user experience</i>