

EUPA2018 - POSTERS NUMBER

POSTER NR.	ID	NAME	SURNAME	TITLE	TOPICS
1	16	ENRIQUE	CALVO	A NOVEL DATA-INDEPENDENT PROTEOMICS APPROACH (DIS) REVEALS THE MECHANISM OF SUPER-ASSEMBLY OF RESPIRATORY COMPLEXES III AND IV	CELLULAR PROTEOMICS
2	62	MARIE	LOCARD-PAULET	PROTEOMIC ANALYSIS OF REGULATORY T-CELLS REVEALS THE IMPORTANCE OF THEMIS1 IN CONTROLLING THEIR SUPPRESSIVE FUNCTION	CELLULAR PROTEOMICS
3	97	MARIA	JASSINSKAJA	IN-DEPTH PROTEOMIC CHARACTERISATION OF RARE HAEMATOPOIETIC PROGENITORS IN THE FOETUS AND ADULT	CELLULAR PROTEOMICS
4	105	BARLIER	CÉLINE	UNTARGETED "LABEL FREE" PROTEOMICS STUDY TO UNDERSTAND PLATELET TRANSFUSION ADVERSE REACTION	CELLULAR PROTEOMICS
5	112	LARADI	SANDRINE	PLATELET PROTEOME CHANGES DURING PLATELET TRANSFUSION ADVERSE REACTION	CELLULAR PROTEOMICS
6	119	EDUARD	SABIDÓ	DIA+: A NOVEL DATA-INDEPENDENT ACQUISITION METHOD COMBINES MULTIPLE PRECURSOR CHARGES TO BOOST PEPTIDE SIGNAL.	CELLULAR PROTEOMICS
7	122	EVELYN	RAMBERGER	THE C/EBPA INTERACTOME UNRAVELLED BY A NOVEL MASS SPECTROMETRIC SCREENING METHOD	CELLULAR PROTEOMICS
8	123	JOSE ANGEL	ALONSO PIÑEIRO	DETERMINATION OF THE SIGNALING PATHWAYS INVOLVED ON HEME-OXYGENASE 1 OVER-EXPRESSION IN EPCS IN ATHEROSCLEROSIS.	CELLULAR PROTEOMICS
9	154	MANUEL	FUENTES	GLOBAL PROTEOME LANDSCAPE DURING ANTIGEN-DEPENDENT B-CELL DIFFERENTIATION REVEALS DYNAMIC CELL SIGNALING PROFILES ACROSS DISTINCT B-CELL SUBPOPULATIONS	CELLULAR PROTEOMICS
10	174	SCHNEIDER	JUSTINE	N-TERMINOMICS AS A TOOL FOR IDENTIFYING PROTEOLYTIC CLEAVAGE OF TOXOPLASMA GONDII	CELLULAR PROTEOMICS
11	182	MAURIZIO	RONCI	SHOTGUN PROTEOMICS OF HUMAN PANCREATIC ISLETS: EFFECT OF CYTOKINES EXPOSURE	CELLULAR PROTEOMICS

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12	185	IRENE	IZQUIERDO	EXTENSIVE PROTEOMIC ANALYSIS OF LIPID RAFTS COMPOSITION ON RESTING AND ACTIVATED PLATELETS.	CELLULAR PROTEOMICS
13	207	MARCELLO	MANFREDI	PROTEOMIC SIGNATURE OF PDAC CELLS WITH MUTANT P53	CELLULAR PROTEOMICS
14	209	ELETTRA	BARBERIS	UNVEILING THE HUMAN ANCIENT PROTEOME	CELLULAR PROTEOMICS
15	210	ELETTRA	BARBERIS	STUDY OF MITOCHONDRIAL PROTEOME CHANGES DUE TO EIF6 DEPLETION WITH SWATH-MS ACQUISITION	CELLULAR PROTEOMICS
16	221	VIVIANA	GRECO	QUANTITATIVE PERSULFIDE SITE IDENTIFICATION HIGHLIGHTS A DIRECT INVOLVEMENT OF H2S IN AMYOTROPHIC LATERAL SCLEROSIS	CELLULAR PROTEOMICS AND PTMS
17	226	BERNHARD	BLANK-LANDESHAMMER	REFINED GENE ANNOTATION AND IDENTIFICATION OF RECODING RNA-EDITING EVENTS IN A FUNGAL MODEL ORGANISM BY PROTEOGENOMICS	CELLULAR PROTEOMICS AND PTMS
18	56	FRANCESC	MÁRQUEZ	TIMSTOF PRO AND PASEF FOR HIGH SENSITIVITY PROTEOMICS AND PHOSPHOPROTEOMICS	PTMS
19	58	HUGO	OSORIO	INNOVATIVE GLYCOPROTEOMIC APPROACHES FOR CANCER BIOMARKER DISCOVERY	PTMS
20	113	MÓNICA	CARRERA	RAPID PHOSPHOPROTEOMICS BY HIFU-TIO2-SCX-LC-MS/MS	PTMS
21	150	HUGO	SANTOS	DIFFERENTIAL PHOSPHORYLATION IN HUMAN FETUIN-A MAY PROVIDE A SWITCH TO PREVENT SOFT TISSUE CALCIFICATION IN PATIENTS IN PERITONEAL DIALYSIS	PTMS
22	153	MANUEL	FUENTES	INTEGRATIVE PROTEOMICS AND PHOSPHOPROTEOMICS PROFILING OF CHRONIC LYMPHOCYTIC LEUKEMIA TUMOR B-CELLS AND THE SIGNALING PATHWAYS INVOLVED	PTMS

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23	183	IRENE	IZQUIERDO	DEPICTING THE PLATELET CLEC-2 SIGNALLING CASCADE BY A PHOSPHOTYROSINE PROTEOMIC APPROACH: IMPACT OF SECONDARY MEDIATORS.	PTMS
24	211	REBECCA	HERZOG	PROTEOME-WIDE REDOX PROFILING IN MESOTHELIAL CELLS DURING EXPERIMENTAL PERITONEAL DIALYSIS BY STRATEGIC THIOL REACTIVE LABELING	PTMS
25	216	SHUAI	WU	AN INTACT WORKFLOW FOR HUMAN CANCER CELL LINE PHOSPHOPEPTIDE ANALYSIS (10/15)	PTMS
26	217	KRISTYNA	PIMKOVA	QUANTITATIVE ANALYSIS OF PROTEIN REDOX POSTTRANSLATIONAL MODIFICATIONS IN HAEMATOPOIETIC STEM AND PROGENITOR CELLS	PTMS
27	227	OLIVER	PAGEL	APPLIED PHOSPHOPROTEOMICS: SELECTIVE VIEW ON CGMP/PKG STIMULATION BY RIOCIGUAT VERSUS CAMP/PKA STIMULATION IN HUMAN PLATELETS	PTMS
28	26	HUGO	LÓPEZ-FERNÁNDEZ	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
29	49	ELENA	PONOMARENKO	ACTION PLAN TO DECIPHER FUNCTIONS OF CHR18 ENCODED PROTEINS IN THE FRAME OF NEXT-CP50 CHALLENGE	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
30	50	JULIAN	USZKOREIT	COMPARISON OF DIFFERENT APPROACHES FOR THE CREATION OF SPECTRAL LIBRARIES FOR DATA INDEPENDENT ACQUISITION ANALYSES	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
31	65	BOUYSSIÉ	DAVID	EFFICIENT AND USER FRIENDLY LABEL-FREE QUANTIFICATION WITH THE PROLINE SOFTWARE SUITE	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
32	85	KSENIA	KUZNETSOVA	ARTIFACTUAL METHIONINE-TO-ISOTHREONINE CONVERSION DURING SAMPLE PREPARATION IN PROTEOGENOMICS	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
33	132	EKATERINA	POVERENNAYA	ESPECIAL CASES OF PROTEOMIC STUDIES: WHICH SPLICE-FORM SHOULD BE TREATED AS CANONICAL?	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION

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POSTER NR.	ID	NAME	SURNAME	TITLE	TOPICS
34	144	OLGA	KISELEVA	CHESS IN PROTEOMICS: WINNING COMBINATION OF 2DE AND MS/MS	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
35	151	CHIARA	MONTI	THE FUNCTIONAL MITOCHONDRIAL HUMAN PROTEOME NETWORK	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
36	155	MOUTON BARBOSA	EMMANUELLE	IN-DEPTH PROTEOMIC CHARACTERIZATION OF COMPLEX CELLULAR PROTEOMES USING THE Q-EXACTIVE HF-X AND PROLINE SOFTWARE	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
37	167	MONICA	FRANCH	PROTEOGENOMICS: COMBINING NEXT GENERATION SEQUENCING AND MASS SPECTROMETRY IN CANDIDA ALBICANS	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
38	213	WOJCIECH	MICHALAK	SOFTWARE FOR QUANTITATIVE ANALYSIS OF PROTEIN COMPLEXES	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
39	219	NAVRATAN	BAGWAN	COMPREHENSIVE HYPOTHESIS-FREE IDENTIFICATION AND QUANTIFICATION OF THE MODIFIED PROTEOME.	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
40	223	SURYA	GUPTA	THE ONLINE TABLOID PROTEOME: AN ANNOTATED DATABASE OF PROTEIN ASSOCIATIONS	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
41	236	REBEKAH	SAYERS	SIMPLIFYING THE USE OF ION LIBRARIES DURING DATA PROCESSING OF DATA INDEPENDENT ACQUISITION PROTEOMICS DATA	BIOINFORMATICS: DATA MANAGEMENT AND INTERPRETATION
42	146	MERCEDES	PARDO	LANDSCAPE OF THE MALARIA INTERACTOME	MICROORGANISMS AND MICROBIOTA
43	156	VANESSA	CASAS	POTENTIAL ROLE OF ACETYLATION ON BACTERIAL METABOLIC PATHWAYS	MICROORGANISMS AND MICROBIOTA
44	170	DINA	REŠETAR	EVALUATION OF PROTEIN EXTRACTION PROTOCOLS FOR SHOTGUN METAPROTEOME CHARACTERIZATION	MICROORGANISMS AND MICROBIOTA

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45	179	GLORIA	MOLERO	STUDY OF THE SECRETOME OF TWO UROPATHOGENIC ESCHERICHIA COLI STRAINS WITH DIFFERENT ANTIBIOTIC SUSCEPTIBILITY.	MICROORGANISMS AND MICROBIOTA
46	193	AIDA	PITARCH	THE ANTIBODY-MEDIATED IMMUNE RESPONSE TO INVASIVE CANDIDIASIS DISCRIMINATES BETWEEN CANDIDA ALBICANS YEASTS AND HYPHAE	MICROORGANISMS AND MICROBIOTA
47	194	GLORIA	MOLERO	PSEUDOMONAS AERUGINOSA PAO1 PEPTIDEATLAS	MICROORGANISMS AND MICROBIOTA
48	234	ALESSIO	SOGGIU	IDENTIFICATION OF NEW CANDIDATE ANTIGENS TO SUPPORT DIVA STRATEGIES IN BRUCELLOSIS	MICROORGANISMS AND MICROBIOTA
49	248	ANA	GONZÁLEZ ABRIL	CHARACTERIZATION OF BACTERIOPHAGE PEPTIDES FOR PATHOGEN IDENTIFICATION BY LC-ESI-MS/MS.	MICROORGANISMS AND MICROBIOTA
50	249	ANA	GONZÁLEZ ABRIL	CHARACTERIZATION OF PEPTIDES FOR HISTAMINE-PRODUCING BACTERIA IDENTIFICATION BY LC-ESI-MS/MS.	MICROORGANISMS AND MICROBIOTA
51	9	EN-CHI	LIAO	SECRETOMIC ANALYSIS OF METASTATIC BIOMARKERS IN ORAL CANCER	DISEASE PROTEOMICS I
52	67	SUSANA	JORGE	OVERCOMING OCT DRAWBACKS FOR RENAL TISSUE PROTEOMICS	DISEASE PROTEOMICS I
53	68	KEVIN	PEREIRA	DIAGNOSING CHROMOPHOBE RENAL CELL CARCINOMA AND RENAL CELL ONCOCYTOMA BY MALDI-MASS SPECTROMETRY-BASED PROFILING	DISEASE PROTEOMICS I
54	78	LUÍS ANDRÉ	CARVALHO	ULTRASONIC-ASSISTED PROTEIN EXTRACTION FROM FFPE NASOPHARYNGEAL CARCINOMA SAMPLES	DISEASE PROTEOMICS I
55	81	JOANA	FLORES	PINPOINTING NEW BIOMARKERS OF BLADDER CANCER IN LIQUID BIOPSIES	DISEASE PROTEOMICS I

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56	87	BEATRIZ	YLLERA	EXOSOMES AS NEW TOOLS FOR SEARCHING RELEVANT PROTEINS IN COLORECTAL CANCER METASTASIS	DISEASE PROTEOMICS I
57	92	LUCIA	SANTORELLI	URINARY EXOSOMES FROM PATIENTS WITH CC-RENAL CELL CARCINOMA AT DIFFERENT STAGES: A PROTEOMIC APPROACH	DISEASE PROTEOMICS I
58	102	RODRIGO	BARDERAS	IDENTIFICATION OF TUMOR ASSOCIATED ANTIGENS IN COLORECTAL CANCER PATIENTS BY MASS-SPECTROMETRY BASED PROTEOMICS	DISEASE PROTEOMICS I
59	133	SITEK	BARBARA	PROTEOME ANALYSIS OF MICRODISSECTED TUMOR CELLS REVEALS ANNEXIN A10 AS BIOMARKER CANDIDATE FOR DIFFERENTIATION OF INTRAHEPATIC CHOLANGIOCARCINOMA AND LIVER METASTASES OF PANCREATIC DUCTAL ADENOCARCINOMA	DISEASE PROTEOMICS I
60	180	ANNA	ILLIANO	DETECTION OF HCT116 MUTATED PROTEINS BY MRM MASS SPECTROMETRY FOR PRECLINICAL DIAGNOSIS OF COLORECTAL CANCER	DISEASE PROTEOMICS I
61	186	FERNANDO	CORRALES	QUANTITATIVE TARGETED PROTEOMIC ANALYSIS OF ONE-CARBON METABOLISM PROTEINS IN HUMAN LIVER CANCER	DISEASE PROTEOMICS I
62	222	FRANCESC	CANALS SURÍS	IMMUNO-MS ASSAY TO MONITOR LIF PROTEIN FOR TUMOR STRATIFICATION AND MONITORING OF LIF-TARGETED THERAPY	DISEASE PROTEOMICS I
63	229	DIANA LASHIDUA	FERNÁNDEZ COTO	QUANTITATIVE PROTEOMICS REVEALS PROTEINS AND PATHWAYS INVOLVED IN THE PROGRESSION FROM NON-CANCEROUS LESIONS TO GASTRIC CANCER.	DISEASE PROTEOMICS I
64	13	FROMENT	CARINE	ANCIENT TOOTH PROTEOMES	PLANTS AND ANIMAL PROTEOMICS
65	41	CHIARA	GUGLIELMETTI	PARAOXONASE 1 (PON1) AS PLASMA MARKER TO DETECT ILLICIT TREATMENT WITH DEXAMETHASONE IN VEAL CALVES	PLANTS AND ANIMAL PROTEOMICS
66	66	JAVIER	BERNAL PAMPÍN	CHANGES IN THE PHOSPHORYLATION STATUS OF STORAGE PROTEINS LINKED TO POTATO SPROUTING	PLANTS AND ANIMAL PROTEOMICS

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67	73	ANGELA	MORENO LÓPZ	PHOSPHOPROTEOMIC ANALYSIS REVEALS THAT SALMONELLA TYPHIMURIUM INCREASES NEUTROPHIL SURVIVAL AFTER INFECTION AND DECREASES THEIR IMMUNE RESPONSE CAPACITY	PLANTS AND ANIMAL PROTEOMICS
68	74	SCHAEFFER-REISS	CHRISTINE	DEVELOPMENT OF MASS SPECTROMETRY-BASED ANALYSIS OF BLOOD PROTEIN GLYCATION IN BIRDS	PLANTS AND ANIMAL PROTEOMICS
69	79	JOSÉ LUIS	CAPELO	PROTEOMIC STUDY IN MUS MUSCULUS UNDER ARSENIC, CADMIUM AND MERCURY EXPOSURE TO EVALUATE METAL TOXICITY	PLANTS AND ANIMAL PROTEOMICS
70	91	LORENA	FRANCO MARTÍNEZ	HAPTOGLOBIN, OVOSTATIN, AND ALBUMIN AS POSSIBLE SALIVARY BIOMARKERS OF CANINE PARVOVIROSIS.	PLANTS AND ANIMAL PROTEOMICS
71	103	ANNA	BASSOLS	A PROTEOMIC APPROACH TO UNDERSTAND THE EFFECTS OF ENVIRONMENTAL ENRICHMENT IN THE PIG HIPPOCAMPUS	PLANTS AND ANIMAL PROTEOMICS
72	124	JESUS V	JORRIN NOVO	PROTEOMICS OF HOLM OAK (QUERCUS ILEX): AN ORPHAN AND RECALCITRANT PLANT SPECIES. HOW TO GET A CONFIDENT PROTEIN IDENTIFICATION AND QUANTIFICATION IN BOTTOM-UP/SHOTGUN APPROACH	PLANTS AND ANIMAL PROTEOMICS
73	125	JESUS V	JORRIN NOVO	PROTEOMICS, AND ITS INTEGRATION WITH TRANSCRIPTOMICS AND METABOLOMICS, ALLOWED THE RECONSTRUCTION OF THE METABOLISM AS IT OCCURS IN QUERCUS ILEX	PLANTS AND ANIMAL PROTEOMICS
74	139	RODRIGO	BARDERAS	THE RECOMBINANT ALLERGEN OLE E 7 ASSEMBLED BY DE NOVO MASS SPECTROMETRY AS A TOOL FOR DIAGNOSIS OF OLIVE POLLEN ALLERGY.	PLANTS AND ANIMAL PROTEOMICS
75	148	YOUNG-HO	CHUNG	PROTEOMIC STUDY ON ADVANCED GLYCATION END-PRODUCTS TREATMENT IN KIDNEY OF MICE	PLANTS AND ANIMAL PROTEOMICS
76	171	LAUTARO	BRACCO	PROTEOMIC CHARACTERIZATION OF SOYBEAN SEED HULL FOR BIOTECHNOLOGICAL PURPOSES	PLANTS AND ANIMAL PROTEOMICS
77	173	LAUTARO	BRACCO	PHYSICOCHEMICAL CHARACTERIZATION OF PROTEIC CONTAMINANTS FROM INSECT LARVAE INFECTED WITH RECOMBINANT BACULOVIRUS FOR PROTEIN EXPRESSION	PLANTS AND ANIMAL PROTEOMICS

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78	231	GERMÁN	ROSANO	PROTEOME VARIATION OF THE RAT LIVER AFTER SIMPLE COLD STORAGE USING AN EX VIVO MODEL	PLANTS AND ANIMAL PROTEOMICS
79	39	MARÍA	GARCÍA VENCE	OPTIMIZATION OF A DEPARAFFINIZATION/EXTRACTION PROTEIN PROTOCOL FROM FFPE THYROID SAMPLES TO APPLY PROTEOMIC TECHNIQUES	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
80	48	ALEXANDER	ARCHAKOV	SENSITIVITY AND SPECIFICITY OF MASS-SPECTROMETRIC PROTEOMIC TECHNOLOGIES AS FACTORS DETERMINING THE PROSPECTS FOR THE CHPP PROJECT	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
81	51	SUSANA BELÉN	BRAVO LÓPEZ	NEW PROTEIN EXTRACTION PROTOCOL FROM FORMALIN-FIXED PARAFFIN EMBEDDED KIDNEY TISSUE SAMPLES	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
82	76	SIMON	DALED	PILLAR BASED HPLC: THE NEXT BIG THING?	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
83	88	GONÇALO	MARTINS	ULTRA-FAST PROTEIN DIGESTION FOR LABEL-FREE PROTEIN QUANTIFICATION	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
84	126	FELIX	ELORTZA	PROTEOMIC ANALYSIS OF SILICA HYBRID SOL-GEL COATINGS: A POTENTIAL TOOL FOR PREDICTING THE BIOCOMPATIBILITY OF IMPLANTS IN VIVO	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
85	135	ALBERTO	PARADELA	OPTIMIZATION OF SERUM EXOSOME ISOLATION METHODS AND PROTEIN CARGO ANALYSIS BY SHOTGUN PROTEOMICS	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
86	168	MARÍA DEL PILAR	CHANTADA VÁZQUEZ	FAST PROTOCOLS FOR THE ENRICHMENT OF LOW-ABUNDANCE PROTEINTS IN PLASMA SAMPLES	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
87	203	PAUL	HUMPHREY	DEVELOP OF A ROBUST AND REPRODUCIBLE GLOBAL PLASMA PROTEOME PROFILING WORKFLOW	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
88	205	PAUL	HUMPHREY	HANDS-FREE SAMPLE PREPARATION FOR PROTEOMICS USING UNIVERSAL CHEMISTRY AND A MICROFLUIDIC BENCHTOP INSTRUMENT	STANDARDIZATION AND TECHNOLOGICAL INNOVATION

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89	214	NDIAYE	SEGA	DEVELOPMENT OF QUALITY CONTROL STANDARD FOR TANDEM MASS TAGS (TMT) WORKFLOW	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
90	218	OXANA	TRIFONOVA	DRIED BLOOD SPOT SAMPLING FOR CLINICAL METABOLOMICS: EFFECTS OF DIFFERENT PAPERS AND SAMPLE STORAGE STABILITY	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
91	220	VIALARET	JEROME	LC/MS MABS QUANTIFICATION IN HUMAN SERUM AFTER PROTEIN-A TIPS OR NSMOL SAMPLE PREPARATIONS	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
92	233	ANTONIO	SERNA SANZ	MULTI-LABORATORY ASSESSMENT OF PERFORMANCE OF SWATH@ ACQUISITION MASS SPECTROMETRY	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
93	235	ANTONIO	SERNA SANZ	STANDARDIZING AND HARMONIZING MULTIPLE TRIPLETOF@ SYSTEMS FOR DDA AND DIA USING A DEDICATED PERFORMANCE KIT	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
94	237	MARÍA ÁNGELES	EXPÓSITO CASTRO	IMPLEMENTATION OF PROTEOMICS AS A QUALITY CONTROL TOOL IN THE FASHION AND FOOD INDUSTRIES	STANDARDIZATION AND TECHNOLOGICAL INNOVATION
95	2	ENRIQUE	SANTAMARÍA	ELUCIDATION OF EARLY-ONSET MOLECULAR DERANGEMENTS IN ALZHEIMER'S DISEASE BY OLFACTORY PROTEOMICS: FROM MICE TO HUMANS	DISEASE PROTEOMICS II
96	27	JOAQUIN	FERNANDEZ-IRIGOYEN	NEUROANATOMICAL QUANTITATIVE PROTEOMICS REVEALS COMMON PATHOGENIC BIOLOGICAL ROUTES BETWEEN AMYOTROPHIC LATERAL SCLEROSIS (ALS) AND FRONTOTEMPORAL DEMENTIA (FTD)	DISEASE PROTEOMICS II
97	34	IRENE	ZUBIRI	UNRAVELLING PHENOTYPIC HETEROGENEITY IN ALS USING QUANTITATIVE PROTEOMICS: FROM ANIMAL MODELS OF THE DISEASE TO HUMAN PATHOLOGY.	DISEASE PROTEOMICS II
98	59	JOSÉ EDUARDO	ARAÚJO	PROTEOMICS ANALYSIS OF THE PERITONEAL DIALYSIS EFFLUENT BY HIRIEF AND TMT: A LONGITUDINAL STUDY	DISEASE PROTEOMICS II
99	61	LAURA	MOURINO-ALVAREZ	IMPROVING DIAGNOSTIC OF DEGENERATIVE AORTIC STENOSIS: FROM RABBIT TO HUMAN SAMPLES	DISEASE PROTEOMICS II

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100	94	CRISTINA	VALENTIM COELHO	PROTEOMIC EFFECTS OF PAP THERAPY ON RED BLOOD CELLS IN MALES WITH OBSTRUCTIVE SLEEP APNEA	DISEASE PROTEOMICS II
101	141	LUCÍA	BELTRÁN CAMACHO	MOLUCULAR CHANGES IN RESPONSE TO CELL THERAPY IN CRITICAL LIMB ISCHEMIA A PROTEOMIC APPROACH	DISEASE PROTEOMICS II
102	142	PABLO	SAN SEGUNDO ACOSTA	IDENTIFICATION OF NEW DIAGNOSTICS BIOMARKERS OF ALZHEIMER DISEASE'S PATIENTS BY PHAGE DISPLAY AND PROTEIN MICROARRAYS	DISEASE PROTEOMICS II
103	145	LIDIA	HERMIDA NOGUEIRA	SECRETOME ANALYSIS OF LEUKOCYTE-PLATELET RICH FIBRIN MEMBRANES USED IN WOUND HEALING	DISEASE PROTEOMICS II
104	147	M ^o CARMEN	DURAN-RUIZ	ASSESSING EPCS POTENTIAL BIOMARKERS FOR CARDIOVASCULAR REGENERATIVE MEDICINE	DISEASE PROTEOMICS II
105	152	ELIANDRE	DE OLIVEIRA	A PROTEOMICS STUDY ON SEPTIC SHOCK PATIENTS: AN OVERVIEW ON SEPSIS-INDUCED MYOCARDIAL DYSFUNCTION	DISEASE PROTEOMICS II
106	162	TIZIANA	ALBERIO	EXPLORING THE PROTEOME ALTERATIONS IN PARK2-MUTATED PATIENTS	DISEASE PROTEOMICS II
107	197	AIDA	PITARCH	COMPUTATIONAL IMMUNOMICS REVEALS AN ANTI-CANDIDA SURFACE ANTIBODY-RECTIVITY SIGNATURE FOR INVASIVE CANDIDIASIS	DISEASE PROTEOMICS II
108	200	FLORIAN	WIESENHOFER	PROTEOMIC AND METABOLOMIC ANALYSIS OF MOLECULAR PROCESSES INVOLVED IN CLINICAL PERITONEAL DIALYSIS	DISEASE PROTEOMICS II
109	201	REBECCA	HERZOG	CROSS-OMICS ANALYSIS OF PROTEOME AND TRANSCRIPTOME DYNAMICS DURING CLINICAL PERITONEAL DIALYSIS THERAPY	DISEASE PROTEOMICS II
110	80	MAR	VILANOVA	TOP-DOWN PROTEOMIC APPROACH FOR THE STUDY OF BASIC PROTEINS IN THE HUMAN SPERM NUCLEUS	TOP-DOWN AND CHEMICAL PROTEOMICS

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111	215	NDIAYE	SEGA	ADVANCES IN ORBITRAP INSTRUMENTATION FOR NATIVE TOP-DOWN ANALYSIS OF NON-COVALENT PROTEIN COMPLEXES	TOP-DOWN AND CHEMICAL PROTEOMICS
112	225	JOAO	RODRIGUES	DEVELOPMENT OF IMMOBILISED AFFINITY BASED PROBES FOR SELECTIVE ENRICHMENT OF GLYCOSYLTRANSFERASES	TOP-DOWN AND CHEMICAL PROTEOMICS
113	6	YI-TING	TSAI	THE ANTRODIA CINNAMOMEA REDUCE OBESITY BY REGULATING LIPID METABOLISM AND BIOSYNTHESIS IN OB/OB MICE	FOOD & NUTRITION PROTEOMICS
114	164	MANUEL	PAZOS	INTEGRATION OF HEPATIC SWATH-BASED PROTEOMICS AND METABOLOMICS TO BETTER UNDERSTAND DIET-INDUCED METABOLIC DISORDERS	FOOD & NUTRITION PROTEOMICS
115	166	SERENA	LACERENZA	A PROTEOMIC APPROACH TO STUDY THE NEUROPROTECTIVE EFFECT OF OLEOCANTHAL IN SH-SY5Y CELLS	FOOD & NUTRITION PROTEOMICS
116	191	VALERIA	MARZANO	PROTEOMICS TO OVERCOME PRECAUTIONARY LABELING: A TRANSLATIONAL APPROACH TO FOOD	FOOD & NUTRITION PROTEOMICS
117	23	DEBORAH	PENQUE	PPROTEOMICS OF OBSTRUCTIVE SLEEP APNEA : NEW INSIGHTS IN PATIENTS WITH DIABETES MELLITUS	DISEASE PROTEOMICS III
118	29	IRAIDE	ESCOBES CORCUERA	DIFFERENTIAL PROTEOMIC ANALYSIS OF ENDOMETRIAL FLUID REFLECTS INCREASED INFLAMMATION AND EXTRACELLULAR MATRIX REMODELING IN NON-IMPLANTATIVE IVF CYCLES	DISEASE PROTEOMICS III
119	35	SOFIA	NEVES	OCCUPATIONAL SECONDHAND SMOKE EXPOSURE: A PROTEOMICS APPROACH	DISEASE PROTEOMICS III
120	71	ANA	GUTIÉRREZ	SEX BIAS ON THE SALIVARY PROTEIN BAND DISTRIBUTION AFTER A MAXIMUM AEROBIC EXERCISE TEST	DISEASE PROTEOMICS III
121	95	NINI JOHANNA	PEDROZA DIAZ	PROTEOMIC ANALYSIS OF CEREBROSPINAL FLUID OF PATIENTS WITH NEUROPSYCHIATRIC LUPUS: LOOKING FOR BIOMARKERS	DISEASE PROTEOMICS III

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122	137	ALBERTO	PARADELA	QUALITATIVE AND QUANTITATIVE PROTEOMICS ANALYSIS IN PLACENTA FOR PREECLAMPSIA-ASSOCIATED BIOMARKER DISCOVERY	DISEASE PROTEOMICS III
123	159	CRISTINA	RUIZ-ROMERO	IDENTIFICATION OF A CIRCULATING PROTEIN SIGNATURE THAT CORRELATES WITH SYNOVIAL PATHOTYPES IN RHEUMATOID ARTHRITIS PATIENTS	DISEASE PROTEOMICS III
124	163	VALENTINA	CALAMIA	COMBINING CLINICAL VARIABLES AND OMICS DATA FOR OSTEOARTHRITIS PATIENT STRATIFICATION	DISEASE PROTEOMICS III
125	188	JAIME	SANCHO	SEMI-QUANTITATIVE ANALYSIS OF PROTEIN ABUNDANCE IN EXOSOMES FROM PERITONEAL EXUDATES IN AN EXPERIMENTAL LUPUS MODEL	DISEASE PROTEOMICS III
126	192	LUCIA	GONZALEZ RODRIGUEZ	TARGETED PROTEOMICS TO DETECT ENDOGENOUS PEPTIDES DIFFERENTIALLY RELEASED FROM HUMAN KNEE AND HIP CARTILAGES	DISEASE PROTEOMICS III
127	212	NATALIA	RIVERA	SERCHING BIOMARKERS IN LUPUS NEPHRITIS BY PROTEOMICS	DISEASE PROTEOMICS III
128	238	CLAUDIA	DI CARLO	MULTI-OMICS PROFILING OF PANCREATIC CANCER STEM-LIKE CELLS	