Karsten Suhre

List of Publications by Citations

265 papers	16,295 citations	66 h-index	121 g-index
311 ext. papers	20,014 ext. citations	8 avg, IF	6.39 L-index

#	Paper	IF	Citations
265	Human metabolic individuality in biomedical and pharmaceutical research. *Nature*, **2011**, 477, 54-60	50.4	728
264	An atlas of genetic influences on human blood metabolites. *Nature Genetics*, **2014**, 46, 543-550	36.3	695
263	ElNemo: a normal mode web server for protein movement analysis and the generation of templates for molecular replacement. *Nucleic Acids Research*, **2004**, 32, W610-4	20.1	544
262	Genetics meets metabolomics: a genome-wide association study of metabolite profiles in human serum. *PLoS Genetics*, **2008**, 4, e1000282	6	538
261	Genomic atlas of the human plasma proteome. *Nature*, **2018**, 558, 73-79	50.4	529
260	A genome-wide perspective of genetic variation in human metabolism. *Nature Genetics*, **2010**, 42, 137-41	36.3	515
259	Novel biomarkers for pre-diabetes identified by metabolomics. *Molecular Systems Biology*, **2012**, 8, 615	12.2	468
258	Metabolic footprint of diabetes: a multiplatform metabolomics study in an epidemiological setting. *PLoS ONE*, **2010**, 5, e13953	3.7	425
257	Metabolomics enables precision medicine: "A White Paper, Community Perspective". *Metabolomics*, **2016**, 12, 149	4.7	327
256	Differences between human plasma and serum metabolite profiles. *PLoS ONE*, **2011**, 6, e21230	3.7	271
255	3DCoffee: combining protein sequences and structures within multiple sequence alignments. *Journal of Molecular Biology*, **2004**, 340, 385-95	6.5	271
254	Biomarkers for type 2 diabetes and impaired fasting glucose using a nontargeted metabolomics approach. *Diabetes*, **2013**, 62, 4270-6	0.9	268
253	Discovery of sexual dimorphisms in metabolic and genetic biomarkers. *PLoS Genetics*, **2011**, 7, e10022156		256
252	Connecting genetic risk to disease end points through the human blood plasma proteome. *Nature Communications*, **2017**, 8, 14357	17.4	249
251	The dynamic range of the human metabolome revealed by challenges. *FASEB Journal*, **2012**, 26, 2607-19	0.9	226
250	Procedure for tissue sample preparation and metabolite extraction for high-throughput targeted metabolomics. *Metabolomics*, **2012**, 8, 133-142	4.7	217
249	Gaussian graphical modeling reconstructs pathway reactions from high-throughput metabolomics data. *BMC Systems Biology*, **2011**, 5, 21	3.5	207

248	Metabolic network failures in Alzheimer's disease: A biochemical roadmap. *Alzheimerjs and Dementia*, **2017**, 13, 965-984	1.2	201
247	Human serum metabolic profiles are age dependent. *Aging Cell*, **2012**, 11, 960-7	9.9	193
246	A genome-wide association study of metabolic traits in human urine. *Nature Genetics*, **2011**, 43, 565-9	36.3	188
245	FusionDB: a database for in-depth analysis of prokaryotic gene fusion events. *Nucleic Acids Research*, **2004**, 32, D273-6	20.1	187
244	MassTRIX: mass translator into pathways. *Nucleic Acids Research*, **2008**, 36, W481-4	20.1	171
243	Metabolomic markers reveal novel pathways of ageing and early development in human populations. *International Journal of Epidemiology*, **2013**, 42, 1111-9	7.8	166
242	SNiPA: an interactive, genetic variant-centered annotation browser. *Bioinformatics*, **2015**, 31, 1334-6	7.2	160
241	Epigenetics meets metabolomics: an epigenome-wide association study with blood serum metabolic traits. *Human Molecular Genetics*, **2014**, 23, 534-45	5.6	147
240	Reductive genome evolution from the mother of Rickettsia. *PLoS Genetics*, **2007**, 3, e14	6	142
239	Mimivirus and the emerging concept of "giant" virus. *Virus Research*, **2006**, 117, 133-44	6.4	141
238	Genetic variation in metabolic phenotypes: study designs and applications. *Nature Reviews Genetics*, **2012**, 13, 759-69	30.1	140
237	3DCoffee@igs: a web server for combining sequences and structures into a multiple sequence alignment. *Nucleic Acids Research*, **2004**, 32, W37-40	20.1	139
236	Tropheryma whipplei Twist: a human pathogenic Actinobacteria with a reduced genome. *Genome Research*, **2003**, 13, 1800-9	9.7	132
235	Genomic correlates of hyperthermostability, an update. *Journal of Biological Chemistry*, **2003**, 278, 17198-202	5.4	131
234	Gender-specific pathway differences in the human serum metabolome. *Metabolomics*, **2015**, 11, 1815-1823	4.7	130
233	Genome-wide association study identifies novel genetic variants contributing to variation in blood metabolite levels. *Nature Communications*, **2015**, 6, 7208	17.4	126
232	Mining the unknown: a systems approach to metabolite identification combining genetic and metabolic information. *PLoS Genetics*, **2012**, 8, e1003005	6	126
231	A Metabolome-Wide Association Study of Kidney Function and Disease in the General Population. *Journal of the American Society of Nephrology: JASN*, **2016**, 27, 1175-88	12.7	119

230	Overview of the Meso-NH model version 5.4 and its applications. *Geoscientific Model Development*, **2018**, 11, 1929-1969	6.3	114
229	Genome-wide mapping of plasma protein QTLs identifies putatively causal genes and pathways for cardiovascular disease. *Nature Communications*, **2018**, 9, 3268	17.4	111
228	A genome-wide metabolic QTL analysis in Europeans implicates two loci shaped by recent positive selection. *PLoS Genetics*, **2011**, 7, e1002270	6	109
227	Childhood obesity is associated with changes in the serum metabolite profile. *Obesity Facts*, **2012**, 5, 660-70	5.1	106
226	Associations of circulating plasma microRNAs with age, body mass index and sex in a population-based study. *BMC Medical Genomics*, **2015**, 8, 61	3.7	105
225	Bioinformatics analysis of targeted metabolomics--uncovering old and new tales of diabetic mice under medication. *Endocrinology*, **2008**, 149, 3478-89	4.8	104
224	Mouse phenotyping. *Methods*, **2011**, 53, 120-35	4.6	103
223	Set1 is required for meiotic S-phase onset, double-strand break formation and middle gene expression. *EMBO Journal*, **2004**, 23, 1957-67	13	103
222	NORMA: a tool for flexible fitting of high-resolution protein structures into low-resolution electron-microscopy-derived density maps. *Acta Crystallographica Section D: Biological Crystallography*, **2006**, 62, 1098-100		99
221	Leveraging cross-species transcription factor binding site patterns: from diabetes risk loci to disease mechanisms. *Cell*, **2014**, 156, 343-58	56.2	96
220	Targeted metabolomics profiles are strongly correlated with nutritional patterns in women. *Metabolomics*, **2013**, 9, 506-514	4.7	93
219	Metabolic profiling reveals distinct variations linked to nicotine consumption in humans--first results from the KORA study. *PLoS ONE*, **2008**, 3, e3863	3.7	92
218	On the potential of normal-mode analysis for solving difficult molecular-replacement problems. *Acta Crystallographica Section D: Biological Crystallography*, **2004**, 60, 796-9		85
217	Serum metabolite concentrations and decreased GFR in the general population. *American Journal of Kidney Diseases*, **2012**, 60, 197-206	7.4	84
216	Metabolites associate with kidney function decline and incident chronic kidney disease in the general population. *Nephrology Dialysis Transplantation*, **2013**, 28, 2131-8	4.3	84
215	CaspR: a web server for automated molecular replacement using homology modelling. *Nucleic Acids Research*, **2004**, 32, W606-9	20.1	81
214	Genetics of human metabolism: an update. *Human Molecular Genetics*, **2015**, 24, R93-R101	5.6	79
213	Comprehensive transcriptomic and proteomic characterization of human mesenchymal stem cells reveals source specific cellular markers. *Scientific Reports*, **2016**, 6, 21507	4.9	79

212	Gene and Genome Duplication in Acanthamoeba polyphaga Mimivirus. *Journal of Virology*, **2005**, 79, 15561-15598	6.6	78
211	Effects of smoking and smoking cessation on human serum metabolite profile: results from the KORA cohort study. *BMC Medicine*, **2013**, 11, 60	11.4	77
210	Metabolomics platforms for genome wide association studies--linking the genome to the metabolome. *Current Opinion in Biotechnology*, **2013**, 24, 39-47	11.4	77
209	Body fat free mass is associated with the serum metabolite profile in a population-based study. *PLoS ONE*, **2012**, 7, e40009	3.7	77
208	Effects of metformin on metabolite profiles and LDL cholesterol in patients with type 2 diabetes. *Diabetes Care*, **2015**, 38, 1858-67	14.6	76
207	Epigenetic associations of type 2 diabetes and BMI in an Arab population. *Clinical Epigenetics*, **2016**, 8, 13	7.7	76
206	Metabolic profiling in diabetes. *Journal of Endocrinology*, **2014**, 221, R75-85	4.7	73
205	High TCR diversity ensures optimal function and homeostasis of Foxp3+ regulatory T cells. *European Journal of Immunology*, **2011**, 41, 3101-13	6.1	71
204	MassTRIX reloaded: combined analysis and visualization of transcriptome and metabolome data. *PLoS ONE*, **2012**, 7, e39860	3.7	71
203	On the hypothesis-free testing of metabolite ratios in genome-wide and metabolome-wide association studies. *BMC Bioinformatics*, **2012**, 13, 120	3.6	70
202	A first genetic map of date palm (Phoenix dactylifera) reveals long-range genome structure conservation in the palms. *BMC Genomics*, **2014**, 15, 285	4.5	68
201	ORILAM, a three-moment lognormal aerosol scheme for mesoscale atmospheric model: Online coupling into the Meso-NH-C model and validation on the Escompte campaign. *Journal of Geophysical Research*, **2005**, 110,		68
200	Genome-Wide Association Study with Targeted and Non-targeted NMR Metabolomics Identifies 15 Novel Loci of Urinary Human Metabolic Individuality. *PLoS Genetics*, **2015**, 11, e1005487	6	66
199	Association of DNA methylation with age, gender, and smoking in an Arab population. *Clinical Epigenetics*, **2015**, 7, 6	7.7	65
198	MIPS: curated databases and comprehensive secondary data resources in 2010. *Nucleic Acids Research*, **2011**, 39, D220-4	20.1	65
197	The Human Blood Metabolome-Transcriptome Interface. *PLoS Genetics*, **2015**, 11, e1005274	6	65
196	Questionnaire-based self-reported nutrition habits associate with serum metabolism as revealed by quantitative targeted metabolomics. *European Journal of Epidemiology*, **2011**, 26, 145-56	12.1	64
195	Genome-wide association study of caffeine metabolites provides new insights to caffeine metabolism and dietary caffeine-consumption behavior. *Human Molecular Genetics*, **2016**, 25, 5472-5482	5.6	64

194	A systems view of type 2 diabetes-associated metabolic perturbations in saliva, blood and urine at different timescales of glycaemic control. *Diabetologia*, **2015**, 58, 1855-67	10.3	63
193	Metabolomic identification of a novel pathway of blood pressure regulation involving hexadecanedioate. *Hypertension*, **2015**, 66, 422-9	8.5	63
192	Mimivirus gene promoters exhibit an unprecedented conservation among all eukaryotes. *Proceedings of the National Academy of Sciences of the United States of America*, **2005**, 102, 14689-93	11.5	63
191	Gene and genome duplication in Acanthamoeba polyphaga Mimivirus. *Journal of Virology*, **2005**, 79, 14095-101	8.6	63
190	Characterization of missing values in untargeted MS-based metabolomics data and evaluation of missing data handling strategies. *Metabolomics*, **2018**, 14, 128	4.7	63
189	Genetics meets proteomics: perspectives for large population-based studies. *Nature Reviews Genetics*, **2021**, 22, 19-37	30.1	62
188	Alcohol-induced metabolomic differences in humans. *Translational Psychiatry*, **2013**, 3, e276	8.6	61
187	Indigenous Arabs are descendants of the earliest split from ancient Eurasian populations. *Genome Research*, **2016**, 26, 151-62	9.7	60
186	1,5-Anhydroglucitol in saliva is a noninvasive marker of short-term glycemic control. *Journal of Clinical Endocrinology and Metabolism*, **2014**, 99, E479-83	5.6	58
185	Integrative genetic and metabolite profiling analysis suggests altered phosphatidylcholine metabolism in asthma. *Allergy: European Journal of Allergy and Clinical Immunology*, **2013**, 68, 629-36	9.3	57
184	Development of a reduced chemical scheme for use in mesoscale meteorological models. *Atmospheric Environment*, **2000**, 34, 2633-2644	5.3	57
183	Genus-wide sequencing supports a two-locus model for sex-determination in Phoenix. *Nature Communications*, **2018**, 9, 3969	17.4	57
182	Internal and external mixing in atmospheric aerosols by coagulation: Impact on the optical and hygroscopic properties of the sulphate-soot system. *Atmospheric Environment*, **1997**, 31, 1393-1402	5.3	54
181	Metabolomic profiles indicate distinct physiological pathways affected by two loci with major divergent effect on Bos taurus growth and lipid deposition. *Physiological Genomics*, **2010**, 42A, 79-88	3.6	53
180	metaP-server: a web-based metabolomics data analysis tool. *Journal of Biomedicine and Biotechnology*, **2011**, 2011,		52
179	Structural genomics of highly conserved microbial genes of unknown function in search of new antibacterial targets. *Journal of Structural and Functional Genomics*, **2003**, 4, 141-57		52
178	Multi-omic signature of body weight change: results from a population-based cohort study. *BMC Medicine*, **2015**, 13, 48	11.4	51
177	Metabolomics approach reveals effects of antihypertensives and lipid-lowering drugs on the human metabolism. *European Journal of Epidemiology*, **2014**, 29, 325-36	12.1	51

176	Long term conservation of human metabolic phenotypes and link to heritability. *Metabolomics*, **2014**, 10, 1005-1017	4.7	50
175	GFscore: a general nonlinear consensus scoring function for high-throughput docking. *Journal of Chemical Information and Modeling*, **2006**, 46, 1704-12	6.1	49
174	Phydbac "Gene Function Predictor": a gene annotation tool based on genomic context analysis. *BMC Bioinformatics*, **2005**, 6, 247	3.6	49
173	Variation in the human lipidome associated with coffee consumption as revealed by quantitative targeted metabolomics. *Molecular Nutrition and Food Research*, **2009**, 53, 1357-65	5.9	48
172	Genetic Influences on Metabolite Levels: A Comparison across Metabolomic Platforms. *PLoS ONE*, **2016**, 11, e0153672	3.7	48
171	Physico-chemical modeling of the First Aerosol Characterization Experiment (ACE 1) Lagrangian B: 1. A moving column approach. *Journal of Geophysical Research*, **1998**, 103, 16433-16455		47
170	Annotation of bacterial genomes using improved phylogenomic profiles. *Bioinformatics*, **2003**, 19 Suppl 1, i105-7	7.2	45
169	ProGeM: a framework for the prioritization of candidate causal genes at molecular quantitative trait loci. *Nucleic Acids Research*, **2019**, 47, e3	20.1	45
168	A pilot study comparing the metabolic profiles of elite-level athletes from different sporting disciplines. *Sports Medicine - Open*, **2018**, 4, 2	6.1	44
167	Ozone-rich transients in the upper equatorial Atlantic troposphere. *Nature*, **1997**, 388, 661-663	50.4	44
166	A systems biology approach using metabolomic data reveals genes and pathways interacting to modulate divergent growth in cattle. *BMC Genomics*, **2013**, 14, 798	4.5	42
165	Urine Metabolite Profiles Predictive of Human Kidney Allograft Status. *Journal of the American Society of Nephrology: JASN*, **2016**, 27, 626-36	12.7	41
164	Estimation of prokaryote genomic DNA G+C content by sequencing universally conserved genes. *International Journal of Systematic and Evolutionary Microbiology*, **2006**, 56, 1025-1029	2.2	41
163	Mesenchymal cell interaction with ovarian cancer cells triggers pro-metastatic properties. *PLoS ONE*, **2012**, 7, e38340	3.7	41
162	Metabolic switch during adipogenesis: From branched chain amino acid catabolism to lipid synthesis. *Archives of Biochemistry and Biophysics*, **2016**, 589, 93-107	4.1	40
161	Changing metabolic signatures of amino acids and lipids during the prediabetic period in a pig model with impaired incretin function and reduced □cell mass. *Diabetes*, **2012**, 61, 2166-75	0.9	40
160	The Saliva Metabolome in Association to Oral Health Status. *Journal of Dental Research*, **2019**, 98, 642-65[illegible]1	11	39
159	A Genome-Wide Survey of Date Palm Cultivars Supports Two Major Subpopulations in Phoenix dactylifera. *G3: Genes, Genomes, Genetics*, **2015**, 5, 1429-38	3.2	39

158	Accelerated lipid catabolism and autophagy are cancer survival mechanisms under inhibited glutaminolysis. *Cancer Letters*, **2018**, 430, 133-147	9.9	38
157	Cohort profile: Greifswald approach to individualized medicine (GANI_MED). *Journal of Translational Medicine*, **2014**, 12, 144	8.5	37
156	Metabolite profiling reveals new insights into the regulation of serum urate in humans. *Metabolomics*, **2014**, 10, 141-151	4.7	36
155	Biochemical insights from population studies with genetics and metabolomics. *Archives of Biochemistry and Biophysics*, **2016**, 589, 168-76	4.1	34
154	Epigenetics meets proteomics in an epigenome-wide association study with circulating blood plasma protein traits. *Nature Communications*, **2020**, 11, 15	17.4	34
153	Network-based approach for analyzing intra- and interfluid metabolite associations in human blood, urine, and saliva. *Journal of Proteome Research*, **2015**, 14, 1183-94	5.6	33
152	Whole-exome sequencing identifies common and rare variant metabolic QTLs in a Middle Eastern population. *Nature Communications*, **2018**, 9, 333	17.4	33
151	Genetic studies of urinary metabolites illuminate mechanisms of detoxification and excretion in humans. *Nature Genetics*, **2020**, 52, 167-176	36.3	32
150	Metabolomics of dates (Phoenix dactylifera) reveals a highly dynamic ripening process accounting for major variation in fruit composition. *BMC Plant Biology*, **2015**, 15, 291	5.3	32
149	Conformational flexibility of Mycobacterium tuberculosis thioredoxin reductase: crystal structure and normal-mode analysis. *Acta Crystallographica Section D: Biological Crystallography*, **2005**, 61, 1603-11		31
148	Effect of induced hypoglycemia on inflammation and oxidative stress in type 2 diabetes and control subjects. *Scientific Reports*, **2020**, 10, 4750	4.9	30
147	Metformin Effect on Nontargeted Metabolite Profiles in Patients With Type 2 Diabetes and in Multiple Murine Tissues. *Diabetes*, **2016**, 65, 3776-3785	0.9	30
146	Increased amino acids levels and the risk of developing of hypertriglyceridemia in a 7-year follow-up. *Journal of Endocrinological Investigation*, **2014**, 37, 369-74	5.2	30
145	Metformin supports the antidiabetic effect of a sodium glucose cotransporter 2 inhibitor by suppressing endogenous glucose production in diabetic mice. *Diabetes*, **2015**, 64, 284-90	0.9	29
144	Defining the genetic control of human blood plasma N-glycome using genome-wide association study. *Human Molecular Genetics*, **2019**, 28, 2062-2077	5.6	28
143	Complementarity of SOMAscan to LC-MS/MS and RNA-seq for quantitative profiling of human embryonic and mesenchymal stem cells. *Journal of Proteomics*, **2017**, 150, 86-97	3.9	28
142	Role of medium- and short-chain L-3-hydroxyacyl-CoA dehydrogenase in the regulation of body weight and thermogenesis. *Endocrinology*, **2011**, 152, 4641-51	4.8	28
141	Improvement of myocardial infarction risk prediction via inflammation-associated metabolite biomarkers. *Heart*, **2017**, 103, 1278-1285	5.1	27

140	Circulating Protein Signatures and Causal Candidates for Type 2 Diabetes. *Diabetes*, **2020**, 69, 1843-1853	10.9	27
139	Mapping the genetic architecture of gene regulation in whole blood. *PLoS ONE*, **2014**, 9, e93844	3.7	27
138	Phydbac2: improved inference of gene function using interactive phylogenomic profiling and chromosomal location analysis. *Nucleic Acids Research*, **2004**, 32, W336-9	20.1	26
137	Systems biology analysis merging phenotype, metabolomic and genomic data identifies Non-SMC Condensin I Complex, Subunit G (NCAPG) and cellular maintenance processes as major contributors to genetic variability in bovine feed efficiency. *PLoS ONE*, **2015**, 10, e0124574	3.7	26
136	Metabolic signatures differentiate ovarian from colon cancer cell lines. *Journal of Translational Medicine*, **2015**, 13, 223	8.5	25
135	Mesenchymal cell interaction with ovarian cancer cells induces a background dependent pro-metastatic transcriptomic profile. *Journal of Translational Medicine*, **2014**, 12, 59	8.5	25
134	Genetic associations with lipoprotein subfractions provide information on their biological nature. *Human Molecular Genetics*, **2012**, 21, 1433-43	5.6	25
133	Identification of a potential biomarker for FABP4 inhibition: the power of lipidomics in preclinical drug testing. *Journal of Biomolecular Screening*, **2011**, 16, 467-75		25
132	Evidence for Stress-like Alterations in the HPA-Axis in Women Taking Oral Contraceptives. *Scientific Reports*, **2017**, 7, 14111	4.9	24
131	Ethnic and gender differences in advanced glycation end products measured by skin auto-fluorescence. *Dermato-Endocrinology*, **2013**, 5, 325-30		24
130	DI-ICR-FT-MS-based high-throughput deep metabotyping: a case study of the Caenorhabditis elegans-Pseudomonas aeruginosa infection model. *Analytical and Bioanalytical Chemistry*, **2015**, 407, 1059-73	4.4	23
129	Metabolomic profiles in individuals with negative affectivity and social inhibition: a population-based study of Type D personality. *Psychoneuroendocrinology*, **2013**, 38, 1299-309	5	23
128	Novel genetic associations with serum level metabolites identified by phenotype set enrichment analyses. *Human Molecular Genetics*, **2014**, 23, 5847-57	5.6	23
127	Determination of strongly overlapping signaling activity from microarray data. *BMC Bioinformatics*, **2006**, 7, 99	3.6	23
126	Inference of gene function based on gene fusion events: the rosetta-stone method. *Methods in Molecular Biology*, **2007**, 396, 31-41	1.4	22
125	Alterations in Lipid and Inositol Metabolisms in Two Dopaminergic Disorders. *PLoS ONE*, **2016**, 11, e0147129	3.7	22
124	Deep molecular phenotypes link complex disorders and physiological insult to CpG methylation. *Human Molecular Genetics*, **2018**, 27, 1106-1121	5.6	21
123	Metabolite ratios as potential biomarkers for type 2 diabetes: a DIRECT study. *Diabetologia*, **2018**, 61, 117-129	10.3	21

122	From Discovery to Translation: Characterization of C-Mannosyltryptophan and Pseudouridine as Markers of Kidney Function. *Scientific Reports*, **2017**, 7, 17400	4.9	21
121	Metabolomics profiling reveals novel markers for leukocyte telomere length. *Aging*, **2016**, 8, 77-94	5.6	21
120	Nesting of colon and ovarian cancer cells in the endothelial niche is associated with alterations in glycan and lipid metabolism. *Scientific Reports*, **2017**, 7, 39999	4.9	20
119	Identification and MS-assisted interpretation of genetically influenced NMR signals in human plasma. *Genome Medicine*, **2013**, 5, 13	14.4	20
118	The association between various smoking behaviors, cotinine biomarkers and skin autofluorescence, a marker for advanced glycation end product accumulation. *PLoS ONE*, **2017**, 12, e0179330	2.7	20
117	Comparison of metabolic profiles of acutely ill and short-term weight recovered patients with anorexia nervosa reveals alterations of 33 out of 163 metabolites. *Journal of Psychiatric Research*, **2012**, 46, 1600-9	5.2	19
116	Characteristics of mutants designed to incorporate a new ion pair into the structure of a cold adapted subtilisin-like serine proteinase. *Biochimica Et Biophysica Acta - Proteins and Proteomics*, **2009**, 1794, 512-8	4	19
115	Associations between thyroid hormones and serum metabolite profiles in an euthyroid population. *Metabolomics*, **2014**, 10, 152-164	4.7	18
114	Bayesian independent component analysis recovers pathway signatures from blood metabolomics data. *Journal of Proteome Research*, **2012**, 11, 4120-31	5.6	18
113	Mimivirus TyrRS: preliminary structural and functional characterization of the first amino-acyl tRNA synthetase found in a virus. *Acta Crystallographica Section F: Structural Biology Communications*, **2005**, 61, 212-5		18
112	Atlantic subtropical potential vorticity barrier as seen by Measurements of Ozone by Airbus In-Service Aircraft (MOZAIC) flights. *Journal of Geophysical Research*, **1998**, 103, 25681-25693		18
111	Type 2 diabetes is associated with postprandial amino acid measures. *Archives of Biochemistry and Biophysics*, **2016**, 589, 138-44	4.1	17
110	Novel subpopulations in date palm (Phoenix dactylifera) identified by population-wide organellar genome sequencing. *BMC Genomics*, **2019**, 20, 498	4.5	17
109	Metabolomics of Dynamic Changes in Insulin Resistance Before and After Exercise in PCOS. *Frontiers in Endocrinology*, **2019**, 10, 116	5.7	17
108	Measurement of 1,5-anhydroglucitol in blood and saliva: from non-targeted metabolomics to biochemical assay. *Journal of Translational Medicine*, **2016**, 14, 140	8.5	17
107	Metabolomics of Ramadan fasting: an opportunity for the controlled study of physiological responses to food intake. *Journal of Translational Medicine*, **2014**, 12, 161	8.5	17
106	PSEA: Phenotype Set Enrichment Analysis--a new method for analysis of multiple phenotypes. *Genetic Epidemiology*, **2012**, 36, 244-52	2.6	17
105	Discovery of protein-coding palindromic repeats in Wolbachia. *Trends in Microbiology*, **2005**, 13, 253-5	12.4	17

104	Metabolomics profiling of xenobiotics in elite athletes: relevance to supplement consumption. *Journal of the International Society of Sports Nutrition*, **2018**, 15, 48	4.5	17
103	Genome-Wide Association Study Reveals a Novel Association Between MYBPC3 Gene Polymorphism, Endurance Athlete Status, Aerobic Capacity and Steroid Metabolism. *Frontiers in Genetics*, **2020**, 11, 595	4.5	16
102	Metabolic Fingerprints of Circulating IGF-1 and the IGF-1/IGFBP-3 Ratio: A Multifluid Metabolomics Study. *Journal of Clinical Endocrinology and Metabolism*, **2016**, 101, 4730-4742	5.6	16
101	Large Scale Metabolic Profiling identifies Novel Steroids linked to Rheumatoid Arthritis. *Scientific Reports*, **2017**, 7, 9137	4.9	16
100	Deciphering the Plasma Proteome of Type 2 Diabetes. *Diabetes*, **2020**, 69, 2766-2778	0.9	15
99	Identification of putative biomarkers for type 2 diabetes using metabolomics in the Korea Association REsource (KARE) cohort. *Metabolomics*, **2016**, 12, 1	4.7	15
98	Unraveling the functional role of the orphan solute carrier, SLC22A24 in the transport of steroid conjugates through metabolomic and genome-wide association studies. *PLoS Genetics*, **2019**, 15, e1008208	6	14
97	Metabolomics Identifies Novel Blood Biomarkers of Pulmonary Function and COPD in the General Population. *Metabolites*, **2019**, 9,	5.6	14
96	Sex differences in urine metabolites related with risk of diabetes using NMR spectroscopy: results of the study of health in pomerania. *Metabolomics*, **2015**, 11, 1405-1415	4.7	14
95	A comprehensive metabolomic data set of date palm fruit. *Data in Brief*, **2018**, 18, 1313-1321	1.2	14
94	Characterization of the metabolic profile associated with serum 25-hydroxyvitamin D: a cross-sectional analysis in population-based data. *International Journal of Epidemiology*, **2016**, 45, 1469-1481	7.8	14
93	Comparative analysis of plasma metabolomics response to metabolic challenge tests in healthy subjects and influence of the FTO obesity risk allele. *Metabolomics*, **2014**, 10, 386-401	4.7	14
92	Interrogating causal pathways linking genetic variants, small molecule metabolites, and circulating lipids. *Genome Medicine*, **2014**, 6, 25	14.4	14
91	Whole genome sequencing in the Middle Eastern Qatari population identifies genetic associations with 45 clinically relevant traits. *Nature Communications*, **2021**, 12, 1250	17.4	14
90	Revealing the role of the human blood plasma proteome in obesity using genetic drivers. *Nature Communications*, **2021**, 12, 1279	17.4	14
89	Automated workflow-based exploitation of pathway databases provides new insights into genetic associations of metabolite profiles. *BMC Genomics*, **2013**, 14, 865	4.5	13
88	Putative gene promoter sequences in the chlorella viruses. *Virology*, **2008**, 380, 388-93	3.6	13
87	Isolation, characterization, and bioinformatic analysis of calmodulin-binding protein cmbB reveals a novel tandem IP22 repeat common to many Dictyostelium and Mimivirus proteins. *Biochemical and Biophysical Research Communications*, **2006**, 346, 879-88	3.4	13

86	Where cancer genomics should go next: a clinician© perspective. *Human Molecular Genetics*, **2014**, 23, R69-75	5.6	12
85	Phenotype-driven identification of modules in a hierarchical map of multifluid metabolic correlations. *Npj Systems Biology and Applications*, **2017**, 3, 28	5	12
84	Mixing of boundary layer and upper tropospheric ozone during a deep convective event over Western Europe. *Atmospheric Environment*, **2002**, 36, 4491-4501	5.3	12
83	Alterations in long noncoding RNAs in women with and without polycystic ovarian syndrome. *Clinical Endocrinology*, **2019**, 91, 793-797	3.4	11
82	On the potential of models for location and scale for genome-wide DNA methylation data. *BMC Bioinformatics*, **2014**, 15, 232	3.6	11
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