Torben Hansen

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

81 208 43,992 340 h-index g-index citations papers 6.09 12 55,203 377 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
340	Four groups of type 2 diabetes contribute to the etiological and clinical heterogeneity in newly diagnosed individuals: An IMI DIRECT study <i>Cell Reports Medicine</i> , 2022 , 3, 100477	18	1
339	Association of milk intake with hay fever, asthma, and lung function: a Mendelian randomization analysis <i>European Journal of Epidemiology</i> , 2022 , 1	12.1	0
338	High incidence of amyotrophic lateral sclerosis in the Faroe Islands 2010-2020 <i>Annals of Clinical and Translational Neurology</i> , 2022 ,	5.3	2
337	Impairment of gut microbial biotin metabolism and host biotin status in severe obesity: effect of biotin and prebiotic supplementation on improved metabolism <i>Gut</i> , 2022 ,	19.2	5
336	Smoking during pregnancy is associated with child overweight independent of maternal pre-pregnancy BMI and genetic predisposition to adiposity <i>Scientific Reports</i> , 2022 , 12, 3135	4.9	O
335	Microbiome and metabolome features of the cardiometabolic disease spectrum <i>Nature Medicine</i> , 2022 ,	50.5	4
334	The Arg82Cys Polymorphism of the Protein Nepmucin Implies a Role in HDL Metabolism <i>Journal of the Endocrine Society</i> , 2022 , 6, bvac034	0.4	O
333	A novel nonsense variant in EXOC8 underlies a neurodevelopmental disorder Neurogenetics, 2022, 1	3	Ο
332	Multi-ancestry genetic study of type 2 diabetes highlights the power of diverse populations for discovery and translation <i>Nature Genetics</i> , 2022 ,	36.3	7
331	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021 ,	50.4	24
330	Combinatorial, additive and dose-dependent drug-microbiome associations. <i>Nature</i> , 2021 ,	50.4	11
329	Recessive Genome-wide Meta-analysis Illuminates Genetic Architecture of Type 2 Diabetes. <i>Diabetes</i> , 2021 ,	0.9	0
328	Processes Underlying Glycemic Deterioration in Type 2 Diabetes: An IMI DIRECT Study. <i>Diabetes Care</i> , 2021 , 44, 511-518	14.6	6
327	A Previously Undescribed Highly Prevalent Phage Identified in a Danish Enteric Virome Catalog. <i>MSystems</i> , 2021 , 6, e0038221	7.6	0
326	Non-linear interaction between physical activity and polygenic risk score of body mass index in Danish and Russian populations. <i>PLoS ONE</i> , 2021 , 16, e0258748	3.7	
325	Evidence for shared genetics between physical activity, sedentary behaviour and adiposity-related traits. <i>Obesity Reviews</i> , 2021 , 22, e13182	10.6	2
324	Amyotrophic lateral sclerosis in the Faroe Islands - a genealogical study. <i>Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration</i> , 2021 , 22, 571-575	3.6	1

(2021-2021)

323	Successful Use of a GLP-1 Receptor Agonist as Add-on Therapy to Sulfonylurea in the Treatment of Neonatal Diabetes. <i>European Journal of Case Reports in Internal Medicine</i> , 2021 , 8, 002352	1.2		
322	The role of a traditional and western diet on glucose homeostasis in Greenlandic Inuit carriers and non-carriers of type 2 diabetes variant in the TBC1D4 gene: A protocol for a randomized clinical trial. <i>Contemporary Clinical Trials Communications</i> , 2021 , 21, 100734	1.8	О	
321	Physical activity attenuates postprandial hyperglycaemia in homozygous TBC1D4 loss-of-function mutation carriers. <i>Diabetologia</i> , 2021 , 64, 1795-1804	10.3	3	
320	FGL1 as a modulator of plasma D-dimer levels: Exome-wide marker analysis of plasma tPA, PAI-1, and D-dimer. <i>Journal of Thrombosis and Haemostasis</i> , 2021 , 19, 2019-2028	15.4	1	
319	Do genetic risk scores for childhood adiposity operate independent of BMI of their mothers?. <i>International Journal of Obesity</i> , 2021 , 45, 2006-2015	5.5		
318	Genetic markers of abdominal obesity and weight loss after gastric bypass surgery. <i>PLoS ONE</i> , 2021 , 16, e0252525	3.7	0	
317	Human and preclinical studies of the host-gut microbiome co-metabolite hippurate as a marker and mediator of metabolic health. <i>Gut</i> , 2021 , 70, 2105-2114	19.2	13	
316	The trans-ancestral genomic architecture of glycemic traits. <i>Nature Genetics</i> , 2021 , 53, 840-860	36.3	44	
315	Conjugated C-6 hydroxylated bile acids in serum relate to human metabolic health and gut Clostridia species. <i>Scientific Reports</i> , 2021 , 11, 13252	4.9	0	
314	Insulin resistance genetic risk score and burden of coronary artery disease in patients referred for coronary angiography. <i>PLoS ONE</i> , 2021 , 16, e0252855	3.7	Ο	
313	Determinants of penetrance and variable expressivity in monogenic metabolic conditions across 77,184 exomes. <i>Nature Communications</i> , 2021 , 12, 3505	17.4	5	
312	Profiles of Glucose Metabolism in Different Prediabetes Phenotypes, Classified by Fasting Glycemia, 2-Hour OGTT, Glycated Hemoglobin, and 1-Hour OGTT: An IMI DIRECT Study. <i>Diabetes</i> , 2021 , 70, 2092-2106	0.9	4	
311	Abdominal and gluteofemoral fat depots show opposing associations with postprandial lipemia. <i>American Journal of Clinical Nutrition</i> , 2021 , 114, 1467-1475	7	2	
310	The Effect of Melatonin on Incretin Hormones: Results From Experimental and Randomized Clinical Studies. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021 , 106, e5109-e5123	5.6	0	
309	Obesity treatment effect in Danish children and adolescents carrying Melanocortin-4 Receptor mutations. <i>International Journal of Obesity</i> , 2021 , 45, 66-76	5.5	2	
308	Acute metabolic effects of melatonin-A randomized crossover study in healthy young men. <i>Journal of Pineal Research</i> , 2021 , 70, e12706	10.4	7	
307	Genome-Wide Association Analysis of Pancreatic Beta-Cell Glucose Sensitivity. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021 , 106, 80-90	5.6	2	
306	A diet-induced gut microbiota component and related plasma metabolites are associated with depressive-like behaviour in rats. <i>European Neuropsychopharmacology</i> , 2021 , 43, 10-21	1.2	7	

305	Large-scale association analyses identify host factors influencing human gut microbiome composition. <i>Nature Genetics</i> , 2021 , 53, 156-165	36.3	80
304	Genome-wide association study of circulating levels of glucagon during an oral glucose tolerance test. <i>BMC Medical Genomics</i> , 2021 , 14, 3	3.7	О
303	Fasting Plasma GLP-1 Is Associated With Overweight/Obesity and Cardiometabolic Risk Factors in Children and Adolescents. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021 , 106, 1718-1727	5.6	3
302	Progressive alcohol-related liver fibrosis is characterised by imbalanced collagen formation and degradation. <i>Alimentary Pharmacology and Therapeutics</i> , 2021 , 54, 1070-1080	6.1	5
301	Genomic and phenotypic insights from an atlas of genetic effects on DNA methylation. <i>Nature Genetics</i> , 2021 , 53, 1311-1321	36.3	27
300	Glucagon-Like Peptide-1 Is Associated With Systemic Inflammation in Pediatric Patients Treated With Hematopoietic Stem Cell Transplantation <i>Frontiers in Immunology</i> , 2021 , 12, 793588	8.4	O
299	Statin therapy is associated with lower prevalence of gut microbiota dysbiosis. <i>Nature</i> , 2020 , 581, 310-3	31550.4	100
298	Mendelian randomization analysis does not support causal associations of birth weight with hypertension risk and blood pressure in adulthood. <i>European Journal of Epidemiology</i> , 2020 , 35, 685-69	7 ^{12.1}	2
297	Skeletal muscle enhancer interactions identify genes controlling whole-body metabolism. <i>Nature Communications</i> , 2020 , 11, 2695	17.4	14
296	Atorvastatin for prevention of disease progression and hospitalisation in liver cirrhosis: protocol for a randomised, double-blind, placebo-controlled trial. <i>BMJ Open</i> , 2020 , 10, e035284	3	3
295	Genetic study of the Arctic CPT1A variant suggests that its effect on fatty acid levels is modulated by traditional Inuit diet. <i>European Journal of Human Genetics</i> , 2020 , 28, 1592-1601	5.3	6
294	GIP and GLP-1 Potentiate Sulfonylurea-Induced Insulin Secretion in Hepatocyte Nuclear Factor 1 Mutation Carriers. <i>Diabetes</i> , 2020 , 69, 1989-2002	0.9	7
293	Predicting and elucidating the etiology of fatty liver disease: A machine learning modeling and validation study in the IMI DIRECT cohorts. <i>PLoS Medicine</i> , 2020 , 17, e1003149	11.6	18
292	The influence of transmitted and non-transmitted parental BMI-associated alleles on the risk of overweight in childhood. <i>Scientific Reports</i> , 2020 , 10, 4806	4.9	7
291	Gain-of-function mutation in the voltage-gated potassium channel gene KCNQ1 and glucose-stimulated hypoinsulinemia - case report. <i>BMC Endocrine Disorders</i> , 2020 , 20, 38	3.3	4
290	Describing the fecal metabolome in cryogenically collected samples from healthy participants. <i>Scientific Reports</i> , 2020 , 10, 885	4.9	8
289	The derived allele of a novel intergenic variant at chromosome 11 associates with lower body mass index and a favorable metabolic phenotype in Greenlanders. <i>PLoS Genetics</i> , 2020 , 16, e1008544	6	1
288	Estimating narrow-sense heritability using family data from admixed populations. <i>Heredity</i> , 2020 , 124, 751-762	3.6	1

(2020-2020)

287	Predictors of weight loss after bariatric surgery-a cross-disciplinary approach combining physiological, social, and psychological measures. <i>International Journal of Obesity</i> , 2020 , 44, 2291-2302	5.5	11
286	Post-load glucose subgroups and associated metabolic traits in individuals with type 2 diabetes: An IMI-DIRECT study. <i>PLoS ONE</i> , 2020 , 15, e0242360	3.7	2
285	GLP-1 Receptor Agonist Treatment in Morbid Obesity and Type 2 Diabetes Due to Pathogenic Homozygous Melanocortin-4 Receptor Mutation: A Case Report. <i>Cell Reports Medicine</i> , 2020 , 1, 100006	18	8
284	FUT2-ABO epistasis increases the risk of early childhood asthma and Streptococcus pneumoniae respiratory illnesses. <i>Nature Communications</i> , 2020 , 11, 6398	17.4	4
283	The effect of diabetes and the common diabetogenic TBC1D4 p.Arg684Ter variant on cardiovascular risk in Inuit in Greenland. <i>Scientific Reports</i> , 2020 , 10, 22081	4.9	3
282	Disturbed eating behaviours do not impact treatment response in a paediatric obesity chronic care treatment programme. <i>Journal of Paediatrics and Child Health</i> , 2020 , 56, 542-549	1.3	4
281	The Effect of Overweight and Obesity on Liver Biochemical Markers in Children and Adolescents. Journal of Clinical Endocrinology and Metabolism, 2020 , 105,	5.6	12
280	Leptin, adiponectin, and their ratio as markers of insulin resistance and cardiometabolic risk in childhood obesity. <i>Pediatric Diabetes</i> , 2020 , 21, 194-202	3.6	30
279	The Polygenic and Monogenic Basis of Blood Traits and Diseases. <i>Cell</i> , 2020 , 182, 1214-1231.e11	56.2	96
278	Novel loci for childhood body mass index and shared heritability with adult cardiometabolic traits. <i>PLoS Genetics</i> , 2020 , 16, e1008718	6	25
277	Efficacy and Safety of Glimepiride With or Without Linagliptin Treatment in Patients With HNF1A Diabetes (Maturity-Onset Diabetes of the Young Type 3): A Randomized, Double-Blinded, Placebo-Controlled, Crossover Trial (GLIMLINA). <i>Diabetes Care</i> , 2020 , 43, 2025-2033	14.6	10
276	Comparative Studies of the Gut Microbiota in the Offspring of Mothers With and Without Gestational Diabetes. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020 , 10, 536282	5.9	4
275	Imidazole propionate is increased in diabetes and associated with dietary patterns and altered microbial ecology. <i>Nature Communications</i> , 2020 , 11, 5881	17.4	29
274	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. <i>Nature Genetics</i> , 2020 , 52, 1314-1332	36.3	26
273	Whole blood co-expression modules associate with metabolic traits and type 2 diabetes: an IMI-DIRECT study. <i>Genome Medicine</i> , 2020 , 12, 109	14.4	3
272	Metabolic and Genetic Risk Factors Are the Strongest Predictors of Severity of Alcohol-Related Liver Fibrosis. <i>Clinical Gastroenterology and Hepatology</i> , 2020 ,	6.9	9
271	Low-grade inflammation independently associates with cardiometabolic risk in children with overweight/obesity. <i>Nutrition, Metabolism and Cardiovascular Diseases</i> , 2020 , 30, 1544-1553	4.5	18
270	Gut microbiota profile and selected plasma metabolites in type 1 diabetes without and with stratification by albuminuria. <i>Diabetologia</i> , 2020 , 63, 2713-2724	10.3	8

269	The intestinal microbiome is a co-determinant of the postprandial plasma glucose response. <i>PLoS ONE</i> , 2020 , 15, e0238648	3.7	1
268	Trans-ethnic and Ancestry-Specific Blood-Cell Genetics in 746,667 Individuals from 5 Global Populations. <i>Cell</i> , 2020 , 182, 1198-1213.e14	56.2	88
267	Genetic Studies of Leptin Concentrations Implicate Leptin in the Regulation of Early Adiposity. <i>Diabetes</i> , 2020 , 69, 2806-2818	0.9	10
266	Genetic Determinants of Electrocardiographic P-Wave Duration and Relation to Atrial Fibrillation. <i>Circulation Genomic and Precision Medicine</i> , 2020 , 13, 387-395	5.2	4
265	Maturity-Onset Diabetes of the Young Identified Among Algerian Probands with Early-Onset Diabetes. <i>Diabetes, Metabolic Syndrome and Obesity: Targets and Therapy</i> , 2020 , 13, 4829-4837	3.4	
264	Data integration for prediction of weight loss in randomized controlled dietary trials. <i>Scientific Reports</i> , 2020 , 10, 20103	4.9	2
263	Structured exercise alters the gut microbiota in humans with overweight and obesity-A randomized controlled trial. <i>International Journal of Obesity</i> , 2020 , 44, 125-135	5.5	34
262	Adults with pathogenic MC4R mutations have increased final height and thereby increased bone mass. <i>Journal of Bone and Mineral Metabolism</i> , 2020 , 38, 117-125	2.9	3
261	Obesity, unfavourable lifestyle and genetic risk of type 2 diabetes: a case-cohort study. <i>Diabetologia</i> , 2020 , 63, 1324-1332	10.3	46
260	Predicting and elucidating the etiology of fatty liver disease: A machine learning modeling and validation study in the IMI DIRECT cohorts 2020 , 17, e1003149		
259	Predicting and elucidating the etiology of fatty liver disease: A machine learning modeling and validation study in the IMI DIRECT cohorts 2020 , 17, e1003149		
258	Predicting and elucidating the etiology of fatty liver disease: A machine learning modeling and validation study in the IMI DIRECT cohorts 2020 , 17, e1003149		
257	Predicting and elucidating the etiology of fatty liver disease: A machine learning modeling and validation study in the IMI DIRECT cohorts 2020 , 17, e1003149		
256	Predicting and elucidating the etiology of fatty liver disease: A machine learning modeling and validation study in the IMI DIRECT cohorts 2020 , 17, e1003149		
255	Abdominal adiposity and cardiometabolic risk factors in children and adolescents: a Mendelian randomization analysis. <i>American Journal of Clinical Nutrition</i> , 2019 , 110, 1079-1087	7	16
254	Screening of 31 genes involved in monogenic forms of obesity in 23 Pakistani probands with early-onset childhood obesity: a case report. <i>BMC Medical Genetics</i> , 2019 , 20, 152	2.1	О
253	Association of Birth Weight With Type 2 Diabetes and Glycemic Traits: A Mendelian Randomization Study. <i>JAMA Network Open</i> , 2019 , 2, e1910915	10.4	14
252	Genome-wide association meta-analysis of 30,000 samples identifies seven novel loci for quantitative ECG traits. <i>European Journal of Human Genetics</i> , 2019 , 27, 952-962	5.3	18

251	Polygenic predisposition to breast cancer and the risk of coronary artery disease. <i>International Journal of Cardiology</i> , 2019 , 291, 145-151	3.2	О
250	Discovery of biomarkers for glycaemic deterioration before and after the onset of type 2 diabetes: descriptive characteristics of the epidemiological studies within the IMI DIRECT Consortium. <i>Diabetologia</i> , 2019 , 62, 1601-1615	10.3	14
249	Exome-Derived Adiponectin-Associated Variants Implicate Obesity and Lipid Biology. <i>American Journal of Human Genetics</i> , 2019 , 105, 15-28	11	12
248	Exome sequencing of 20,791 cases of type 2 diabetes and 24,440 controls. <i>Nature</i> , 2019 , 570, 71-76	50.4	129
247	Glucose metabolism in children and adolescents: Population-based reference values and comparisons to children and adolescents enrolled in obesity treatment. <i>Pediatric Diabetes</i> , 2019 , 20, 538-548	3.6	10
246	Pro12Ala Ala carriers exhibit greater improvements in peripheral insulin sensitivity in response to 12 weeks of aerobic exercise training. <i>Physiological Genomics</i> , 2019 , 51, 254-260	3.6	2
245	Extracellular Vesicle Encapsulated MicroRNAs in Patients with Type 2 Diabetes Are Affected by Metformin Treatment. <i>Journal of Clinical Medicine</i> , 2019 , 8,	5.1	26
244	Genetic Determinants of Weight Loss After Bariatric Surgery. <i>Obesity Surgery</i> , 2019 , 29, 2554-2561	3.7	10
243	FADS and PPARG2 Single Nucleotide Polymorphisms are Associated with Plasma Lipids in 9-Mo-Old Infants. <i>Journal of Nutrition</i> , 2019 , 149, 708-715	4.1	3
242	Reappraisal of variants previously linked with sudden infant death syndrome: results from three population-based cohorts. <i>European Journal of Human Genetics</i> , 2019 , 27, 1427-1435	5.3	4
241	Maternal and fetal genetic effects on birth weight and their relevance to cardio-metabolic risk factors. <i>Nature Genetics</i> , 2019 , 51, 804-814	36.3	181
240	Haploinsufficiency of ARHGAP42 is associated with hypertension. <i>European Journal of Human Genetics</i> , 2019 , 27, 1296-1303	5.3	6
239	Diabetic cats have decreased gut microbial diversity and a lack of butyrate producing bacteria. <i>Scientific Reports</i> , 2019 , 9, 4822	4.9	26
238	Metformin-induced changes of the gut microbiota in healthy young men: results of a non-blinded, one-armed intervention study. <i>Diabetologia</i> , 2019 , 62, 1024-1035	10.3	79
237	Association of genetic variants previously implicated in coronary artery disease with age at onset of coronary artery disease requiring revascularizations. <i>PLoS ONE</i> , 2019 , 14, e0211690	3.7	2
236	Reference values for fasting serum concentrations of thyroid-stimulating hormone and thyroid hormones in healthy Danish/North-European white children and adolescents. <i>Scandinavian Journal of Clinical and Laboratory Investigation</i> , 2019 , 79, 129-135	2	11
235	Increased frequency of rare missense PPP1R3B variants among Danish patients with type 2 diabetes. <i>PLoS ONE</i> , 2019 , 14, e0210114	3.7	6
234	Genome-wide meta-analysis of macronutrient intake of 91,114 European ancestry participants from the cohorts for heart and aging research in genomic epidemiology consortium. <i>Molecular Psychiatry</i> , 2019 , 24, 1920-1932	15.1	30

233	Sequencing reveals protective and pathogenic effects on development of diabetes of rare GLIS3 variants. <i>PLoS ONE</i> , 2019 , 14, e0220805	3.7	2
232	A trans-ancestral meta-analysis of genome-wide association studies reveals loci associated with childhood obesity. <i>Human Molecular Genetics</i> , 2019 , 28, 3327-3338	5.6	30
231	Genetic predisposition to higher body fat yet lower cardiometabolic risk in children and adolescents. <i>International Journal of Obesity</i> , 2019 , 43, 2007-2016	5.5	5
230	Urinary nucleic acid oxidation product levels show differential associations with pharmacological treatment in patients with type 2 diabetes. <i>Free Radical Research</i> , 2019 , 53, 694-703	4	4
229	Human pancreatic islet three-dimensional chromatin architecture provides insights into the genetics of type 2 diabetes. <i>Nature Genetics</i> , 2019 , 51, 1137-1148	36.3	111
228	Effects of Calcium, Magnesium, and Potassium Concentrations on Ventricular Repolarization in Unselected Individuals. <i>Journal of the American College of Cardiology</i> , 2019 , 73, 3118-3131	15.1	12
227	The expression of four pyridoxal kinase (PDXK) human variants in Drosophila impacts on genome integrity. <i>Scientific Reports</i> , 2019 , 9, 14188	4.9	5
226	Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9. <i>BMC Cardiovascular Disorders</i> , 2019 , 19, 240	2.3	8
225	Genetic Aspects of Latent Autoimmune Diabetes in Adults: A Mini-Review. <i>Current Diabetes Reviews</i> , 2019 , 15, 194-198	2.7	7
224	Protein-coding variants implicate novel genes related to lipid homeostasis contributing to body-fat distribution. <i>Nature Genetics</i> , 2019 , 51, 452-469	36.3	44
223	Impaired glucose metabolism and altered gut microbiome despite calorie restriction of ob/ob mice. <i>Animal Microbiome</i> , 2019 , 1, 11	4.1	5
222	Associations between thyroid-stimulating hormone, blood pressure and adiponectin are attenuated in children and adolescents with overweight or obesity. <i>Journal of Pediatric Endocrinology and Metabolism</i> , 2019 , 32, 1351-1358	1.6	0
221	Association of alcohol consumption with allergic disease and asthma: a multi-centre Mendelian randomization analysis. <i>Addiction</i> , 2019 , 114, 216-225	4.6	7
220	A novel rare CUBN variant and three additional genes identified in Europeans with and without diabetes: results from an exome-wide association study of albuminuria. <i>Diabetologia</i> , 2019 , 62, 292-305	10.3	17
219	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. <i>American Journal of Human Genetics</i> , 2019 , 104, 112-138	11	54
218	The antidepressant-like effect of probiotics and their faecal abundance may be modulated by the cohabiting gut microbiota in rats. <i>European Neuropsychopharmacology</i> , 2019 , 29, 98-110	1.2	14
217	Effect modification of FADS2 polymorphisms on the association between breastfeeding and intelligence: results from a collaborative meta-analysis. <i>International Journal of Epidemiology</i> , 2019 , 48, 45-57	7.8	2
216	ADAMTS9 Regulates Skeletal Muscle Insulin Sensitivity Through Extracellular Matrix Alterations. Diabetes, 2019, 68, 502-514	0.9	11

(2018-2018)

Genetic architecture of obesity and related metabolic traits-recent insights from isolated populations. <i>Current Opinion in Genetics and Development</i> , 2018 , 50, 74-78	4.9	3
Genetic determinants of glycated hemoglobin levels in the Greenlandic Inuit population. <i>European Journal of Human Genetics</i> , 2018 , 26, 868-875	5.3	3
Refining the accuracy of validated target identification through coding variant fine-mapping in type 2 diabetes. <i>Nature Genetics</i> , 2018 , 50, 559-571	36.3	221
Impact of a vegan diet on the human salivary microbiota. <i>Scientific Reports</i> , 2018 , 8, 5847	4.9	50
Prospective Studies Exploring the Possible Impact of an ID3 Polymorphism on Changes in Obesity Measures. <i>Obesity</i> , 2018 , 26, 747-754	8	1
Aberrant intestinal microbiota in individuals with prediabetes. <i>Diabetologia</i> , 2018 , 61, 810-820	10.3	163
Re-analysis of public genetic data reveals a rare X-chromosomal variant associated with type 2 diabetes. <i>Nature Communications</i> , 2018 , 9, 321	17.4	50
Bone turnover, calcium homeostasis, and vitamin D status in Danish vegans. <i>European Journal of Clinical Nutrition</i> , 2018 , 72, 1046-1054	5.2	17
Associations of genetic determinants of serum vitamin B12 and folate concentrations with hay fever and asthma: a Mendelian randomization meta-analysis. <i>European Journal of Clinical Nutrition</i> , 2018 , 72, 264-271	5.2	6
Evidence of a liver-alpha cell axis in humans: hepatic insulin resistance attenuates relationship between fasting plasma glucagon and glucagonotropic amino acids. <i>Diabetologia</i> , 2018 , 61, 671-680	10.3	41
Loss-of-function variants in ADCY3 increase risk of obesity and type 2 diabetes. <i>Nature Genetics</i> , 2018 , 50, 172-174	36.3	97
Patient profiling for success after weight loss surgery (GO Bypass study): An interdisciplinary study protocol. <i>Contemporary Clinical Trials Communications</i> , 2018 , 10, 121-130	1.8	14
A Genome-Wide Association Study of Diabetic Kidney Disease in Subjects With Type 2 Diabetes. <i>Diabetes</i> , 2018 , 67, 1414-1427	0.9	71
Genome-Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. <i>Molecular Nutrition and Food Research</i> , 2018 , 62, 1700347	5.9	5
Identification of novel high-impact recessively inherited type 2 diabetes risk variants in the Greenlandic population. <i>Diabetologia</i> , 2018 , 61, 2005-2015	10.3	11
Brugada Syndrome-Associated Genetic Loci Are Associated With J-Point Elevation and an Increased Risk of Cardiac Arrest. <i>Frontiers in Physiology</i> , 2018 , 9, 894	4.6	1
Genome-wide association and HLA fine-mapping studies identify risk loci and genetic pathways underlying allergic rhinitis. <i>Nature Genetics</i> , 2018 , 50, 1072-1080	36.3	52
Exome-chip meta-analysis identifies novel loci associated with cardiac conduction, including ADAMTS6. <i>Genome Biology</i> , 2018 , 19, 87	18.3	25
	Genetic determinants of glycated hemoglobin levels in the Greenlandic Inuit population. <i>European Journal of Human Genetics</i> , 2018, 26, 868-875 Refining the accuracy of validated target identification through coding variant fine-mapping in type 2 diabetes. <i>Nature Genetics</i> , 2018, 50, 559-571 Impact of a vegan diet on the human salivary microbiota. <i>Scientific Reports</i> , 2018, 8, 5847 Prospective Studies Exploring the Possible Impact of an ID3 Polymorphism on Changes in Obesity Measures. <i>Obesity</i> , 2018, 26, 747-754 Aberrant intestinal microbiota in individuals with prediabetes. <i>Diabetologia</i> , 2018, 61, 810-820 Re-analysis of public genetic data reveals a rare X-chromosomal variant associated with type 2 diabetes. <i>Nature Communications</i> , 2018, 9, 321 Bone turnover, calcium homeostasis, and vitamin D status in Danish vegans. <i>European Journal of Clinical Nutrition</i> , 2018, 72, 1046-1054 Associations of genetic determinants of serum vitamin B12 and folate concentrations with hay fever and asthma: a Mendellan randomization meta-analysis. <i>European Journal of Clinical Nutrition</i> , 2018, 72, 642-271 Evidence of a liver-alpha cell axis in humans: hepatic insulin resistance attenuates relationship between fasting plasma glucagon and glucagonotropic amino acids. <i>Diabetologia</i> , 2018, 61, 671-680 Loss-of-function variants in ADCY3 increase risk of obesity and type 2 diabetes. <i>Nature Genetics</i> , 2018, 50, 172-174 Patient profiling for success after weight loss surgery (GO Bypass study): An interdisciplinary study protocol. <i>Contemporary Clinical Trials Communications</i> , 2018, 10, 121-130 A Genome-Wide Association Study of Diabetic Kidney Disease in Subjects With Type 2 Diabetes. <i>Diabetes</i> , 2018, 67, 1414-1427 Genome-Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. <i>Molecular Nutrition and Food Research</i> , 2018, 62, 1700347 Identification of novel high-impact recessively inherited type 2 diabetes risk variants in the Greenlandic population. <i>Diabetologia</i> , 2018, 61, 2005-20	populations. Current Opinion in Genetics and Development, 2018, 50, 74-78 Genetic determinants of glycated hemoglobin levels in the Greenlandic Inuit population. European Journal of Human Genetics, 2018, 26, 868-875 Refining the accuracy of validated target identification through coding variant fine-mapping in type 2 diabetes. Nature Genetics, 2018, 50, 559-571 Impact of a vegan diet on the human salivary microbiota. Scientific Reports, 2018, 8, 5847 Prospective Studies Exploring the Possible Impact of an ID3 Polymorphism on Changes in Obesity Measures. Obesity, 2018, 26, 747-754 Aberrant intestinal microbiota in individuals with prediabetes. Diabetologia, 2018, 61, 810-820 Re-analysis of public genetic data reveals a rare X-chromosomal variant associated with type 2 diabetes. Nature Communications, 2018, 9, 321 Bone turnover, calcium homeostasis, and vitamin D status in Danish vegans. European Journal of Clinical Nutrition, 2018, 72, 1045-1054 Associations of genetic determinants of serum vitamin B12 and folate concentrations with hay fever and asthma: a Mendelian randomization meta-analysis. European Journal of Clinical Nutrition, 2018, 72, 264-271 Evidence of a liver-alpha cell axis in humans: hepatic insulin resistance attenuates relationship between fasting plasma glucagon and glucagonotropic amino acids. Diabetologia, 2018, 61, 671-680 Loss-of-function variants in ADCY3 increase risk of obesity and type 2 diabetes. Nature Genetics, 2018, 50, 172-174 Patient profiling for success after weight loss surgery (CO Bypass study): An interdisciplinary study protocol. Contemporary Clinical Trials Communications, 2018, 10, 121-130 A Genome-Wide Association Study of Diabetic Kidney Disease in Subjects With Type 2 Diabetes. Diabetes, 2018, 67, 1414-1427 Genome-Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. Molecular Nutrition and Food Research, 2018, 61, 2005-2015 Genome-Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. Molecul

197	Gestational diabetes is associated with change in the gut microbiota composition in third trimester of pregnancy and postpartum. <i>Microbiome</i> , 2018 , 6, 89	16.6	155
196	Common variants in the hERG (KCNH2) voltage-gated potassium channel are associated with altered fasting and glucose-stimulated plasma incretin and glucagon responses. <i>BMC Genetics</i> , 2018 , 19, 15	2.6	8
195	Genetics of metabolic traits in Greenlanders: lessons from an isolated population. <i>Journal of Internal Medicine</i> , 2018 , 284, 464-477	10.8	4
194	Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. <i>Nature Communications</i> , 2018 , 9, 2976	17.4	45
193	Meta-analysis of exome array data identifies six novel genetic loci for lung function. <i>Wellcome Open Research</i> , 2018 , 3, 4	4.8	16
192	Childhood obesity treatment; Effects on BMI SDS, body composition, and fasting plasma lipid concentrations. <i>PLoS ONE</i> , 2018 , 13, e0190576	3.7	7
191	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. <i>Nature Genetics</i> , 2018 , 50, 26-41	36.3	186
190	Impaired fasting glucose and the metabolic profile in Danish children and adolescents with normal weight, overweight, or obesity. <i>Pediatric Diabetes</i> , 2018 , 19, 356-365	3.6	8
189	Obesity is associated with vitamin D deficiency in Danish children and adolescents. <i>Journal of Pediatric Endocrinology and Metabolism</i> , 2018 , 31, 53-61	1.6	26
188	Genetic risk scores for body fat distribution attenuate weight loss in women during dietary intervention. <i>International Journal of Obesity</i> , 2018 , 42, 370-375	5.5	12
187	The effect of impaired glucose metabolism on weight loss in multidisciplinary childhood obesity treatment. <i>Pediatric Diabetes</i> , 2018 , 19, 366-374	3.6	6
186	Genetic insights into fetal growth and measures of glycaemic regulation and adiposity in adulthood: a family-based study. <i>BMC Medical Genetics</i> , 2018 , 19, 207	2.1	1
185	The effect of drinking water pH on the human gut microbiota and glucose regulation: results of a randomized controlled cross-over intervention. <i>Scientific Reports</i> , 2018 , 8, 16626	4.9	16
184	Identification of novel LEPR mutations in Pakistani families with morbid childhood obesity. <i>BMC Medical Genetics</i> , 2018 , 19, 199	2.1	4
183	Genetic Susceptibility for Childhood BMI has no Impact on Weight Loss Following Lifestyle Intervention in Danish Children. <i>Obesity</i> , 2018 , 26, 1915-1922	8	7
182	A low-gluten diet induces changes in the intestinal microbiome of healthy Danish adults. <i>Nature Communications</i> , 2018 , 9, 4630	17.4	69
181	Hypertension genetic risk score is associated with burden of coronary heart disease among patients referred for coronary angiography. <i>PLoS ONE</i> , 2018 , 13, e0208645	3.7	6
180	Glimepiride monotherapy versus combination of glimepiride and linagliptin therapy in patients with HNF1A-diabetes: a protocol for a randomised, double-blinded, placebo-controlled trial. <i>BMJ Open</i> , 2018 , 8, e022517	3	5

179	Fine-mapping type 2 diabetes loci to single-variant resolution using high-density imputation and islet-specific epigenome maps. <i>Nature Genetics</i> , 2018 , 50, 1505-1513	36.3	675
178	Population-based studies of relationships between dietary acidity load, insulin resistance and incident diabetes in Danes. <i>Nutrition Journal</i> , 2018 , 17, 91	4.3	9
177	First Genome-Wide Association Study of Latent Autoimmune Diabetes in Adults Reveals Novel Insights Linking Immune and Metabolic Diabetes. <i>Diabetes Care</i> , 2018 , 41, 2396-2403	14.6	57
176	Genome-wide analyses identify a role for SLC17A4 and AADAT in thyroid hormone regulation. <i>Nature Communications</i> , 2018 , 9, 4455	17.4	75
175	A computational framework to integrate high-throughput QomicsQdatasets for the identification of potential mechanistic links. <i>Nature Protocols</i> , 2018 , 13, 2781-2800	18.8	44
174	Recovery of gut microbiota of healthy adults following antibiotic exposure. <i>Nature Microbiology</i> , 2018 , 3, 1255-1265	26.6	246
173	Spatial QRS-T angle variants for prediction of all-cause mortality. <i>Journal of Electrocardiology</i> , 2018 , 51, 768-775	1.4	7
172	Cardiolipin Synthesis in Brown and Beige Fat Mitochondria Is Essential for Systemic Energy Homeostasis. <i>Cell Metabolism</i> , 2018 , 28, 159-174.e11	24.6	67
171	Patients with Obesity Caused by Melanocortin-4 Receptor Mutations Can Be Treated with a Glucagon-like Peptide-1 Receptor Agonist. <i>Cell Metabolism</i> , 2018 , 28, 23-32.e3	24.6	49
170	Birth weight variants are associated with variable fetal intrauterine growth from 20 weeks of gestation. <i>Scientific Reports</i> , 2018 , 8, 8376	4.9	3
169	An adult-based insulin resistance genetic risk score associates with insulin resistance, metabolic traits and altered fat distribution in Danish children and adolescents who are overweight or obese. <i>Diabetologia</i> , 2018 , 61, 1769-1779	10.3	8
168	Genetic Variations in the Human G Protein-coupled Receptor Class C, Group 6, Member A (GPRC6A) Control Cell Surface Expression and Function. <i>Journal of Biological Chemistry</i> , 2017 , 292, 1524-1534	5.4	16
167	Reference intervals for C-peptide and insulin derived from a general adult Danish population. <i>Clinical Biochemistry</i> , 2017 , 50, 408-413	3.5	9
166	Rare and low-frequency coding variants alter human adult height. <i>Nature</i> , 2017 , 542, 186-190	50.4	412
165	A study of associations between early DHA status and fatty acid desaturase (FADS) SNP and developmental outcomes in children of obese mothers. <i>British Journal of Nutrition</i> , 2017 , 117, 278-286	3.6	9
164	Patients With Long-QT Syndrome Caused by Impaired -Encoded K11.1 Potassium Channel Have Exaggerated Endocrine Pancreatic and Incretin Function Associated With Reactive Hypoglycemia. <i>Circulation</i> , 2017 , 135, 1705-1719	16.7	20
163	Genetic evidence of a causal effect of insulin resistance on branched-chain amino acid levels. <i>Diabetologia</i> , 2017 , 60, 873-878	10.3	79
162	Discovery of novel heart rate-associated loci using the Exome Chip. <i>Human Molecular Genetics</i> , 2017 , 26, 2346-2363	5.6	17

161	Selection in Europeans on Fatty Acid Desaturases Associated with Dietary Changes. <i>Molecular Biology and Evolution</i> , 2017 , 34, 1307-1318	8.3	50
160	Carriers of a enhancer polymorphism selectively binding CHOP/DDIT3 are predisposed to increased circulating levels of thyroid-stimulating hormone. <i>Journal of Medical Genetics</i> , 2017 , 54, 166-175	5.8	8
159	A Genome-Wide Association Study of IVGTT-Based Measures of First-Phase Insulin Secretion Refines the Underlying Physiology of Type 2 Diabetes Variants. <i>Diabetes</i> , 2017 , 66, 2296-2309	0.9	69
158	FGF21 Is a Sugar-Induced Hormone Associated with Sweet Intake and Preference in Humans. <i>Cell Metabolism</i> , 2017 , 25, 1045-1053.e6	24.6	123
157	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. <i>Nature Communications</i> , 2017 , 8, 14977	17.4	105
156	The Effect of an Extreme and Prolonged Population Bottleneck on Patterns of Deleterious Variation: Insights from the Greenlandic Inuit. <i>Genetics</i> , 2017 , 205, 787-801	4	31
155	PCSK9 genetic variants and risk of type 2 diabetes: a mendelian randomisation study. <i>Lancet Diabetes and Endocrinology,the</i> , 2017 , 5, 97-105	18.1	225
154	Ponderal index at birth associates with later risk of gestational diabetes mellitus. <i>Archives of Gynecology and Obstetrics</i> , 2017 , 296, 249-256	2.5	8
153	Dyslipidemia and reference values for fasting plasma lipid concentrations in Danish/North-European White children and adolescents. <i>BMC Pediatrics</i> , 2017 , 17, 116	2.6	39
152	Self-Reported Versus Accelerometer-Assessed Daily Physical Activity in Childhood Obesity Treatment. <i>Perceptual and Motor Skills</i> , 2017 , 124, 795-811	2.2	1
151	An Expanded Genome-Wide Association Study of Type 2 Diabetes in Europeans. <i>Diabetes</i> , 2017 , 66, 288	38 - 290	2 414
150	A Low-Frequency Inactivating Variant Enriched in the Finnish Population Is Associated With Fasting Insulin Levels and Type 2 Diabetes Risk. <i>Diabetes</i> , 2017 , 66, 2019-2032	0.9	29
149	The effect of smoking on the urinary excretion of 8-oxodG and 8-oxoGuo in patients with type 2 diabetes. <i>Scandinavian Journal of Clinical and Laboratory Investigation</i> , 2017 , 77, 253-258	2	7
148	and Loci Identified through Large-Scale Exome Chip Analysis Regulate Kidney Development and Function. <i>Journal of the American Society of Nephrology: JASN</i> , 2017 , 28, 981-994	12.7	30
147	Dietary Assessment in the MetaCardis Study: Development and Relative Validity of an Online Food Frequency Questionnaire. <i>Journal of the Academy of Nutrition and Dietetics</i> , 2017 , 117, 878-888	3.9	18
146	Genetic determinants of serum vitamin B12 and their relation to body mass index. <i>European Journal of Epidemiology</i> , 2017 , 32, 125-134	12.1	22
145	Cardiovascular and All-Cause Mortality Risk Associated With Urinary Excretion of 8-oxoGuo, a Biomarker for RNA Oxidation, in Patients With Type 2 Diabetes: A Prospective Cohort Study. <i>Diabetes Care</i> , 2017 , 40, 1771-1778	14.6	36
144	Exome-wide association study of plasma lipids in >300,000 individuals. <i>Nature Genetics</i> , 2017 , 49, 1758-	1 36 .6	310

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143	Higher intake of fish and fat is associated with lower plasma s-adenosylhomocysteine: a cross-sectional study. <i>Nutrition Research</i> , 2017 , 46, 78-87	4	1
142	New Blood Pressure-Associated Loci Identified in Meta-Analyses of 475 000 Individuals. <i>Circulation: Cardiovascular Genetics</i> , 2017 , 10,		33
141	Urinary metabolomics reveals glycemic and coffee associated signatures of thyroid function in two population-based cohorts. <i>PLoS ONE</i> , 2017 , 12, e0173078	3.7	15
140	Causal relationship between obesity and serum testosterone status in men: A bi-directional mendelian randomization analysis. <i>PLoS ONE</i> , 2017 , 12, e0176277	3.7	47
139	Alcohol consumption and its interaction with adiposity-associated genetic variants in relation to subsequent changes in waist circumference and body weight. <i>Nutrition Journal</i> , 2017 , 16, 51	4.3	5
138	Cohort description: The Danish study of Functional Disorders. <i>Clinical Epidemiology</i> , 2017 , 9, 127-139	5.9	50
137	Effect of the interaction between diet composition and the genetic variant on insulin resistance and Itell function markers during weight loss: results from the Nutrient Gene Interactions in Human Obesity: implications for dietary guidelines (NUGENOB) randomized trial. American Journal	7	19
136	of Clinical Nutrition, 2017 , 106, 902-908 Large-scale GWAS identifies multiple loci for hand grip strength providing biological insights into muscular fitness. <i>Nature Communications</i> , 2017 , 8, 16015	17.4	80
135	Investigating the causal effect of smoking on hay fever and asthma: a Mendelian randomization meta-analysis in the CARTA consortium. <i>Scientific Reports</i> , 2017 , 7, 2224	4.9	24
134	Environmental spread of microbes impacts the development of metabolic phenotypes in mice transplanted with microbial communities from humans. <i>ISME Journal</i> , 2017 , 11, 676-690	11.9	41
133	Heterogeneity in glucose response curves during an oral glucose tolerance test and associated cardiometabolic risk. <i>Endocrine</i> , 2017 , 55, 427-434	4	18
132	Sequence data and association statistics from 12,940 type 2 diabetes cases and controls. <i>Scientific Data</i> , 2017 , 4, 170179	8.2	22
131	High Prevalence of Diabetes-Predisposing Variants in MODY Genes Among Danish Women With Gestational Diabetes Mellitus. <i>Journal of the Endocrine Society</i> , 2017 , 1, 681-690	0.4	23
130	Genome-wide physical activity interactions in adiposity - A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017 , 13, e1006528	6	103
129	Comparison of HapMap and 1000 Genomes Reference Panels in a Large-Scale Genome-Wide Association Study. <i>PLoS ONE</i> , 2017 , 12, e0167742	3.7	21
128	A genome-wide association study of thyroid stimulating hormone and free thyroxine in Danish children and adolescents. <i>PLoS ONE</i> , 2017 , 12, e0174204	3.7	14
127	Human biallelic MFN2 mutations induce mitochondrial dysfunction, upper body adipose hyperplasia, and suppression of leptin expression. <i>ELife</i> , 2017 , 6,	8.9	42
126	Sequencing and de novo assembly of 150 genomes from Denmark as a population reference. <i>Nature</i> , 2017 , 548, 87-91	50.4	87

125	Archaic Adaptive Introgression in TBX15/WARS2. Molecular Biology and Evolution, 2017, 34, 509-524	8.3	63
124	Genetic investigations of sudden unexpected deaths in infancy using next-generation sequencing of 100 genes associated with cardiac diseases. <i>European Journal of Human Genetics</i> , 2016 , 24, 817-22	5.3	44
123	Interactions between genetic variants associated with adiposity traits and soft drinks in relation to longitudinal changes in body weight and waist circumference. <i>American Journal of Clinical Nutrition</i> , 2016 , 104, 816-26	7	31
122	A principal component meta-analysis on multiple anthropometric traits identifies novel loci for body shape. <i>Nature Communications</i> , 2016 , 7, 13357	17.4	46
121	Plasma Alkylresorcinols Reflect Gluten Intake and Distinguish between Gluten-Rich and Gluten-Poor Diets in a Population at Risk of Metabolic Syndrome. <i>Journal of Nutrition</i> , 2016 , 146, 1991-	1 9 98	13
120	Trans-ancestry meta-analyses identify rare and common variants associated with blood pressure and hypertension. <i>Nature Genetics</i> , 2016 , 48, 1151-1161	36.3	181
119	Genome-Wide Association Study of the Modified Stumvoll Insulin Sensitivity Index Identifies BCL2 and FAM19A2 as Novel Insulin Sensitivity Loci. <i>Diabetes</i> , 2016 , 65, 3200-11	0.9	47
118	Human gut microbes impact host serum metabolome and insulin sensitivity. <i>Nature</i> , 2016 , 535, 376-81	50.4	977
117	Blood-based biomarkers of age-associated epigenetic changes in human islets associate with insulin secretion and diabetes. <i>Nature Communications</i> , 2016 , 7, 11089	17.4	145
116	A genomic approach to therapeutic target validation identifies a glucose-lowering GLP1R variant protective for coronary heart disease. <i>Science Translational Medicine</i> , 2016 , 8, 341ra76	17.5	77
115	Weight loss and weight maintenance obtained with or without GLP-1 analogue treatment decrease branched chain amino acid levels. <i>Metabolomics</i> , 2016 , 12, 1	4.7	
114	Transcriptional interactions suggest niche segregation among microorganisms in the human gut. <i>Nature Microbiology</i> , 2016 , 1, 16152	26.6	38
113	Colonic transit time is related to bacterial metabolism and mucosal turnover in the gut. <i>Nature Microbiology</i> , 2016 , 1, 16093	26.6	204
112	An Improved Method for High Quality Metagenomics DNA Extraction from Human and Environmental Samples. <i>Scientific Reports</i> , 2016 , 6, 26775	4.9	101
111	Insights into metabolic disease from studying genetics in isolated populations: stories from Greece to Greenland. <i>Diabetologia</i> , 2016 , 59, 938-41	10.3	8
110	Genome-wide association studies in the Japanese population identify seven novel loci for type 2 diabetes. <i>Nature Communications</i> , 2016 , 7, 10531	17.4	99
109	Obesity and Bariatric Surgery Drive Epigenetic Variation of Spermatozoa in Humans. <i>Cell Metabolism</i> , 2016 , 23, 369-78	24.6	320
108	Levels of the inflammation marker YKL-40 in young adults exposed to intrauterine hyperglycemia. <i>Diabetes Research and Clinical Practice</i> , 2016 , 114, 50-4	7.4	1

(2016-2016)

107	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. <i>Nature Communications</i> , 2016 , 7, 10495	17.4	180
106	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. <i>Nature Communications</i> , 2016 , 7, 10494	17.4	107
105	Genome-wide association analysis identifies three new susceptibility loci for childhood body mass index. <i>Human Molecular Genetics</i> , 2016 , 25, 389-403	5.6	202
104	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. <i>Human Molecular Genetics</i> , 2016 , 25, 358-70	5.6	54
103	Identification of Novel Genetic Determinants of Erythrocyte Membrane Fatty Acid Composition among Greenlanders. <i>PLoS Genetics</i> , 2016 , 12, e1006119	6	16
102	Effects of a Family-Based Childhood Obesity Treatment Program on Parental Weight Status. <i>PLoS ONE</i> , 2016 , 11, e0161921	3.7	11
101	Genetic Correlation between Body Fat Percentage and Cardiorespiratory Fitness Suggests Common Genetic Etiology. <i>PLoS ONE</i> , 2016 , 11, e0166738	3.7	8
100	Vejle Diabetes Biobank - a resource for studies of the etiologies of diabetes and its comorbidities. <i>Clinical Epidemiology</i> , 2016 , 8, 393-413	5.9	17
99	Serum 25-Hydroxyvitamin D Status and Longitudinal Changes in Weight and Waist Circumference: Influence of Genetic Predisposition to Adiposity. <i>PLoS ONE</i> , 2016 , 11, e0153611	3.7	5
98	The genetic architecture of type 2 diabetes. <i>Nature</i> , 2016 , 536, 41-47	50.4	704
98 97	The genetic architecture of type 2 diabetes. <i>Nature</i> , 2016 , 536, 41-47 Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. <i>American Journal of Human Genetics</i> , 2016 , 99, 8-21	50.4	7°4 47
	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. <i>American</i>		, ,
97	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. <i>American Journal of Human Genetics</i> , 2016 , 99, 8-21 Genetic risk scores link body fat distribution with specific cardiometabolic profiles. <i>Obesity</i> , 2016 ,	11	47
97 96	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. <i>American Journal of Human Genetics</i> , 2016 , 99, 8-21 Genetic risk scores link body fat distribution with specific cardiometabolic profiles. <i>Obesity</i> , 2016 , 24, 1778-85 DNA methylation and gene expression of HIF3A: cross-tissue validation and associations with BMI	11 8 7.7	47
97 96 95	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. <i>American Journal of Human Genetics</i> , 2016 , 99, 8-21 Genetic risk scores link body fat distribution with specific cardiometabolic profiles. <i>Obesity</i> , 2016 , 24, 1778-85 DNA methylation and gene expression of HIF3A: cross-tissue validation and associations with BMI and insulin resistance. <i>Clinical Epigenetics</i> , 2016 , 8, 89 Roux-en-Y gastric bypass surgery of morbidly obese patients induces swift and persistent changes	111 8 7-7	47 2 29
97 96 95 94	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. <i>American Journal of Human Genetics</i> , 2016 , 99, 8-21 Genetic risk scores link body fat distribution with specific cardiometabolic profiles. <i>Obesity</i> , 2016 , 24, 1778-85 DNA methylation and gene expression of HIF3A: cross-tissue validation and associations with BMI and insulin resistance. <i>Clinical Epigenetics</i> , 2016 , 8, 89 Roux-en-Y gastric bypass surgery of morbidly obese patients induces swift and persistent changes of the individual gut microbiota. <i>Genome Medicine</i> , 2016 , 8, 67	111 8 7-7	47 2 29 187
97 96 95 94 93	Exome Genotyping Identifies Pleiotropic Variants Associated with Red Blood Cell Traits. <i>American Journal of Human Genetics</i> , 2016 , 99, 8-21 Genetic risk scores link body fat distribution with specific cardiometabolic profiles. <i>Obesity</i> , 2016 , 24, 1778-85 DNA methylation and gene expression of HIF3A: cross-tissue validation and associations with BMI and insulin resistance. <i>Clinical Epigenetics</i> , 2016 , 8, 89 Roux-en-Y gastric bypass surgery of morbidly obese patients induces swift and persistent changes of the individual gut microbiota. <i>Genome Medicine</i> , 2016 , 8, 67 Genetics of Type 2 Diabetes: the Power of Isolated Populations. <i>Current Diabetes Reports</i> , 2016 , 16, 65 Genetic variants associated with subjective well-being, depressive symptoms, and neuroticism	11 8 7.7 14.4 5.6	47 2 29 187

89	Genome-wide associations for birth weight and correlations with adult disease. <i>Nature</i> , 2016 , 538, 248-2	2 5 2.4	266
88	FTO genotype and weight loss: systematic review and meta-analysis of 9563 individual participant data from eight randomised controlled trials. <i>BMJ, The</i> , 2016 , 354, i4707	5.9	70
87	GLP-1 Response to Oral Glucose Is Reduced in Prediabetes, Screen-Detected Type 2 Diabetes, and Obesity and Influenced by Sex: The ADDITION-PRO Study. <i>Diabetes</i> , 2015 , 64, 2513-25	0.9	171
86	Rare genetic variants previously associated with congenital forms of long QT syndrome have little or no effect on the QT interval. <i>European Heart Journal</i> , 2015 , 36, 2523-9	9.5	45
85	GLP-1 Receptor Agonist Treatment Increases Bone Formation and Prevents Bone Loss in Weight-Reduced Obese Women. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015 , 100, 2909-17	5.6	84
84	Inherited coding variants at the CDKN2A locus influence susceptibility to acute lymphoblastic leukaemia in children. <i>Nature Communications</i> , 2015 , 6, 7553	17.4	51
83	Identification and functional characterization of G6PC2 coding variants influencing glycemic traits define an effector transcript at the G6PC2-ABCB11 locus. <i>PLoS Genetics</i> , 2015 , 11, e1004876	6	76
82	Genetic and phenotypic correlations between surrogate measures of insulin release obtained from OGTT data. <i>Diabetologia</i> , 2015 , 58, 1006-12	10.3	6
81	Postprandial incretin and islet hormone responses and dipeptidyl-peptidase 4 enzymatic activity in patients with maturity onset diabetes of the young. <i>European Journal of Endocrinology</i> , 2015 , 173, 205-	1 5 .5	10
80	Impact of age, BMI and HbA1c levels on the genome-wide DNA methylation and mRNA expression patterns in human adipose tissue and identification of epigenetic biomarkers in blood. <i>Human Molecular Genetics</i> , 2015 , 24, 3792-813	5.6	168
79	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. <i>Nature Genetics</i> , 2015 , 47, 1282-1293	36.3	223
78	Genome-wide association studies of human adiposity: Zooming in on synapses. <i>Molecular and Cellular Endocrinology</i> , 2015 , 418 Pt 2, 90-100	4.4	7
77	Reference values for serum total adiponectin in healthy non-obese children and adolescents. <i>Clinica Chimica Acta</i> , 2015 , 450, 11-4	6.2	26
76	Reduced CD300LG mRNA tissue expression, increased intramyocellular lipid content and impaired glucose metabolism in healthy male carriers of Arg82Cys in CD300LG: a novel genometabolic cross-link between CD300LG and common metabolic phenotypes. <i>BMJ Open Diabetes Research and</i>	4.5	8
75	Greenlandic Inuit show genetic signatures of diet and climate adaptation. <i>Science</i> , 2015 , 349, 1343-7	33.3	298
74	Disentangling type 2 diabetes and metformin treatment signatures in the human gut microbiota. <i>Nature</i> , 2015 , 528, 262-266	50.4	1107
73	Genetic fine mapping and genomic annotation defines causal mechanisms at type 2 diabetes susceptibility loci. <i>Nature Genetics</i> , 2015 , 47, 1415-25	36.3	292
72	A glycogene mutation map for discovery of diseases of glycosylation. <i>Glycobiology</i> , 2015 , 25, 211-24	5.8	38

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Common variants in LEPR, IL6, AMD1, and NAMPT do not associate with risk of juvenile and childhood obesity in Danes: a case-control study. <i>BMC Medical Genetics</i> , 2015 , 16, 105	2.1	7
Multidisciplinary care of obese children and adolescents for one year reduces ectopic fat content in liver and skeletal muscle. <i>BMC Pediatrics</i> , 2015 , 15, 196	2.6	23
Intake of macro- and micronutrients in Danish vegans. <i>Nutrition Journal</i> , 2015 , 14, 115	4.3	64
The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015 , 11, e1005378	6	220
Effect of Antibiotics on Gut Microbiota, Gut Hormones and Glucose Metabolism. <i>PLoS ONE</i> , 2015 , 10, e0142352	3.7	61
Impact of PTBP1 rs11085226 on glucose-stimulated insulin release in adult Danes. <i>BMC Medical Genetics</i> , 2015 , 16, 17	2.1	4
Interactions of Lipid Genetic Risk Scores With Estimates of Metabolic Health in a Danish Population. <i>Circulation: Cardiovascular Genetics</i> , 2015 , 8, 465-72		25
Uncovering the genetic history of the present-day Greenlandic population. <i>American Journal of Human Genetics</i> , 2015 , 96, 54-69	11	61
Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. <i>Nature Communications</i> , 2015 , 6, 5897	17.4	147
1H-MRS Measured Ectopic Fat in Liver and Muscle in Danish Lean and Obese Children and Adolescents. <i>PLoS ONE</i> , 2015 , 10, e0135018	3.7	15
Comparative Analyses of QTLs Influencing Obesity and Metabolic Phenotypes in Pigs and Humans. <i>PLoS ONE</i> , 2015 , 10, e0137356	3.7	15
Diabetes in Population Isolates: Lessons from Greenland. <i>Review of Diabetic Studies</i> , 2015 , 12, 320-9	3.6	4
Identification of low-frequency and rare sequence variants associated with elevated or reduced risk of type 2 diabetes. <i>Nature Genetics</i> , 2014 , 46, 294-8	36.3	241
Incretin effect and glucagon responses to oral and intravenous glucose in patients with maturity-onset diabetes of the youngtype 2 and type 3. <i>Diabetes</i> , 2014 , 63, 2838-44	0.9	34
Cohort Profile: the Health2006 cohort, research centre for prevention and health. <i>International Journal of Epidemiology</i> , 2014 , 43, 568-75	7.8	76
Discovery of biomarkers for glycaemic deterioration before and after the onset of type 2 diabetes: rationale and design of the epidemiological studies within the IMI DIRECT Consortium. <i>Diabetologia</i> , 2014 , 57, 1132-42	10.3	39
Identification and assembly of genomes and genetic elements in complex metagenomic samples without using reference genomes. <i>Nature Biotechnology</i> , 2014 , 32, 822-8	44.5	624
An integrated catalog of reference genes in the human gut microbiome. <i>Nature Biotechnology</i> , 2014 , 32, 834-41	44.5	1088
	childhood obesity in Danes: a case-control study. <i>BMC Medical Genetics</i> , 2015 , 16, 105 Multidisciplinary care of obese children and adolescents for one year reduces ectopic fat content in liver and skeletal muscle. <i>BMC Pediatrics</i> , 2015 , 15, 196 Intake of macro- and micronutrients in Danish vegans. <i>Nutrition Journal</i> , 2015 , 14, 115 The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015 , 11, e1005378 Effect of Antibiotics on Gut Microbiota, Gut Hormones and Glucose Metabolism. <i>PLoS ONE</i> , 2015 , 10, e0142352 Impact of PTBP1 rs11085226 on glucose-stimulated insulin release in adult Danes. <i>BMC Medical Genetics</i> , 2015 , 16, 17 Interactions of Lipid Genetic Risk Scores With Estimates of Metabolic Health in a Danish Population. <i>Circulation: Cardiovascular Genetics</i> , 2015 , 8, 465-72 Uncovering the genetic history of the present-day Greenlandic population. <i>American Journal of Human Genetics</i> , 2015 , 96, 54-69 Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. <i>Nature Communications</i> , 2015 , 6, 5897 1H-MRS Measured Ectopic Fat in Liver and Muscle in Danish Lean and Obese Children and Adolescents. <i>PLoS ONE</i> , 2015 , 10, e0135018 Comparative Analyses of QTLs Influencing Obesity and Metabolic Phenotypes in Pigs and Humans. <i>PLoS ONE</i> , 2015 , 10, e0137356 Diabetes in Population Isolates: Lessons from Greenland. <i>Review of Diabetic Studies</i> , 2015 , 12, 320-9 Identification of low-frequency and rare sequence variants associated with elevated or reduced risk of type 2 diabetes. <i>Nature Genetics</i> , 2014 , 46, 294-8 Incretin effect and glucagon responses to oral and intravenous glucose in patients with maturity-onset diabetes of the young-type 2 and type 3. <i>Diabetes</i> , 2014 , 63, 2838-44 Cohort Profile: the Health2006 cohort, research centre for prevention and health. <i>International Journal of Epidemiology</i> , 2014 , 43, 568-75 Discovery	childhood obesity in Danes: a case-control study. <i>BMC Medical Genetics</i> , 2015, 16, 105 Multidisciplinary care of obese children and adolescents for one year reduces ectopic fat content in liver and skeletal muscle. <i>BMC Pediatrics</i> , 2015, 15, 196 Intake of macro- and micronutrients in Danish vegans. <i>Nutrition Journal</i> , 2015, 14, 115 43 The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , 2015, 11, e1005378 Effect of Antibiotics on Gut Microbiota, Gut Hormones and Glucose Metabolism. <i>PLoS ONE</i> , 2015, 10, e0142352 Impact of PTBP1 rs11085226 on glucose-stimulated insulin release in adult Danes. <i>BMC Medical Genetics</i> , 2015, 16, 17 Interactions of Lipid Genetic Risk Scores With Estimates of Metabolic Health in a Danish Population. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 465-72 Uncovering the genetic history of the present-day Greenlandic population. <i>American Journal of Human Genetics</i> , 2015, 96, 54-69 Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. <i>Nature Communications</i> , 2015, 6, 5897 17-4 1H-MRS Measured Ectopic Fat in Liver and Muscle in Danish Lean and Obese Children and Adolescents. <i>PLoS ONE</i> , 2015, 10, e0135018 Comparative Analyses of QTLs Influencing Obesity and Metabolic Phenotypes in Pigs and Humans. <i>PLoS ONE</i> , 2015, 10, e0137356 Jiabetes in Population Isolates: Lessons from Greenland. <i>Review of Diabetic Studies</i> , 2015, 12, 320-9 36 Identification of low-frequency and rare sequence variants associated with elevated or reduced risk of type 2 diabetes. <i>Nature Genetics</i> , 2014, 46, 294-8 Incretin effect and glucagon responses to oral and intravenous glucose in patients with maturity-onset diabetes of the young-type 2 and type 3. <i>Diabetes</i> , 2014, 63, 2838-44 Cohort Profile: the Health2006 cohort, research centre for prevention and health. <i>International Journal of Epidemiology</i> , 2014, 43, 568-75 Discovery of biomarkers fo

53	Evaluation of a target region capture sequencing platform using monogenic diabetes as a study-model. <i>BMC Genetics</i> , 2014 , 15, 13	2.6	42
52	Glucose-lowering effects and low risk of hypoglycemia in patients with maturity-onset diabetes of the young when treated with a GLP-1 receptor agonist: a double-blind, randomized, crossover trial. <i>Diabetes Care</i> , 2014 , 37, 1797-805	14.6	56
51	A common Greenlandic TBC1D4 variant confers muscle insulin resistance and type 2 diabetes. <i>Nature</i> , 2014 , 512, 190-3	50.4	258
50	Genetic susceptibility to type 2 diabetes and obesity: from genome-wide association studies to rare variants and beyond. <i>Diabetologia</i> , 2014 , 57, 1528-41	10.3	135
49	Blood pressure levels in male carriers of Arg82Cys in CD300LG. <i>PLoS ONE</i> , 2014 , 9, e109646	3.7	2
48	Impact of type 2 diabetes susceptibility variants on quantitative glycemic traits reveals mechanistic heterogeneity. <i>Diabetes</i> , 2014 , 63, 2158-71	0.9	235
47	Gene-lifestyle interaction and type 2 diabetes: the EPIC interact case-cohort study. <i>PLoS Medicine</i> , 2014 , 11, e1001647	11.6	149
46	FTO genetic variants, dietary intake and body mass index: insights from 177,330 individuals. <i>Human Molecular Genetics</i> , 2014 , 23, 6961-72	5.6	120
45	Variation and association to diabetes in 2000 full mtDNA sequences mined from an exome study in a Danish population. <i>European Journal of Human Genetics</i> , 2014 , 22, 1040-5	5.3	23
44	Pleiotropic genes for metabolic syndrome and inflammation. <i>Molecular Genetics and Metabolism</i> , 2014 , 112, 317-38	3.7	81
43	Interaction between genetic predisposition to adiposity and dietary protein in relation to subsequent change in body weight and waist circumference. <i>PLoS ONE</i> , 2014 , 9, e110890	3.7	13
42	Exome sequencing-driven discovery of coding polymorphisms associated with common metabolic phenotypes. <i>Diabetologia</i> , 2013 , 56, 298-310	10.3	102
41	Richness of human gut microbiome correlates with metabolic markers. <i>Nature</i> , 2013 , 500, 541-6	50.4	2584
40	Whole-exome sequencing of 2,000 Danish individuals and the role of rare coding variants in type 2 diabetes. <i>American Journal of Human Genetics</i> , 2013 , 93, 1072-86	11	109
39	New loci associated with birth weight identify genetic links between intrauterine growth and adult height and metabolism. <i>Nature Genetics</i> , 2013 , 45, 76-82	36.3	232
38	Does DNA methylation of PPARGC1A influence insulin action in first degree relatives of patients with type 2 diabetes?. <i>PLoS ONE</i> , 2013 , 8, e58384	3.7	22
37	A metagenome-wide association study of gut microbiota in type 2 diabetes. <i>Nature</i> , 2012 , 490, 55-60	50.4	3779
36	Low-grade inflammation in young adults exposed to intrauterine hyperglycemia. <i>Diabetes Research and Clinical Practice</i> , 2012 , 97, 322-30	7.4	8

(2009-2012)

35	The effect of FOXA2 rs1209523 on glucose-related phenotypes and risk of type 2 diabetes in Danish individuals. <i>BMC Medical Genetics</i> , 2012 , 13, 10	2.1	10
34	Novel loci for adiponectin levels and their influence on type 2 diabetes and metabolic traits: a multi-ethnic meta-analysis of 45,891 individuals. <i>PLoS Genetics</i> , 2012 , 8, e1002607	6	326
33	No interactions between previously associated 2-hour glucose gene variants and physical activity or BMI on 2-hour glucose levels. <i>Diabetes</i> , 2012 , 61, 1291-6	0.9	21
32	TFAP2B influences the effect of dietary fat on weight loss under energy restriction. <i>PLoS ONE</i> , 2012 , 7, e43212	3.7	28
31	Genome-wide population-based association study of extremely overweight young adultsthe GOYA study. <i>PLoS ONE</i> , 2011 , 6, e24303	3.7	90
30	Enterotypes of the human gut microbiome. <i>Nature</i> , 2011 , 473, 174-80	50.4	4240
29	Physical activity attenuates the influence of FTO variants on obesity risk: a meta-analysis of 218,166 adults and 19,268 children. <i>PLoS Medicine</i> , 2011 , 8, e1001116	11.6	379
28	Genetic variation in GIPR influences the glucose and insulin responses to an oral glucose challenge. <i>Nature Genetics</i> , 2010 , 42, 142-8	36.3	527
27	Twelve type 2 diabetes susceptibility loci identified through large-scale association analysis. <i>Nature Genetics</i> , 2010 , 42, 579-89	36.3	1449
26	A genome-wide association study in the Japanese population identifies susceptibility loci for type 2 diabetes at UBE2E2 and C2CD4A-C2CD4B. <i>Nature Genetics</i> , 2010 , 42, 864-8	36.3	214
25	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. <i>Nature Genetics</i> , 2010 , 42, 949-60	36.3	724
24	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010 , 42, 937-48	36.3	2267
23	New genetic loci implicated in fasting glucose homeostasis and their impact on type 2 diabetes risk. <i>Nature Genetics</i> , 2010 , 42, 105-16	36.3	1673
22	G-allele of intronic rs10830963 in MTNR1B confers increased risk of impaired fasting glycemia and type 2 diabetes through an impaired glucose-stimulated insulin release: studies involving 19,605 Europeans. <i>Diabetes</i> , 2009 , 58, 1450-6	0.9	111
21	Identification of novel variants in the hepatocyte nuclear factor-1alpha gene in South Indian patients with maturity onset diabetes of young. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2009 , 94, 1959-65	5.6	31
20	Genome-wide association yields new sequence variants at seven loci that associate with measures of obesity. <i>Nature Genetics</i> , 2009 , 41, 18-24	36.3	1085
19	A variant near MTNR1B is associated with increased fasting plasma glucose levels and type 2 diabetes risk. <i>Nature Genetics</i> , 2009 , 41, 89-94	36.3	466
18	Genetic variant near IRS1 is associated with type 2 diabetes, insulin resistance and hyperinsulinemia. <i>Nature Genetics</i> , 2009 , 41, 1110-5	36.3	356

17	Joint analysis of individual participants Qdata from 17 studies on the association of the IL6 variant -174G>C with circulating glucose levels, interleukin-6 levels, and body mass index. <i>Annals of Medicine</i> , 2009 , 41, 128-38	1.5	47
16	Overweight and the metabolic syndrome in adult offspring of women with diet-treated gestational diabetes mellitus or type 1 diabetes. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2009 , 94, 2464-7	o ^{5.6}	304
15	Variant near ADAMTS9 known to associate with type 2 diabetes is related to insulin resistance in offspring of type 2 diabetes patientsEUGENE2 study. <i>PLoS ONE</i> , 2009 , 4, e7236	3.7	43
14	Common nonsynonymous variants in PCSK1 confer risk of obesity. <i>Nature Genetics</i> , 2008 , 40, 943-5	36.3	242
13	Meta-analysis of genome-wide association data and large-scale replication identifies additional susceptibility loci for type 2 diabetes. <i>Nature Genetics</i> , 2008 , 40, 638-45	36.3	1496
12	High prevalence of type 2 diabetes and pre-diabetes in adult offspring of women with gestational diabetes mellitus or type 1 diabetes: the role of intrauterine hyperglycemia. <i>Diabetes Care</i> , 2008 , 31, 340-6	14.6	481
11	Low physical activity accentuates the effect of the FTO rs9939609 polymorphism on body fat accumulation. <i>Diabetes</i> , 2008 , 57, 95-101	0.9	386
10	A variant in CDKAL1 influences insulin response and risk of type 2 diabetes. <i>Nature Genetics</i> , 2007 , 39, 770-5	36.3	851
9	Melanocortin 4 receptor mutations in obese Czech children: studies of prevalence, phenotype development, weight reduction response, and functional analysis. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2007 , 92, 3689-96	5.6	59
8	Mutation analysis of the preproghrelin gene: no association with obesity and type 2 diabetes. <i>Clinical Biochemistry</i> , 2005 , 38, 420-4	3.5	43
7	The pathophysiology of diabetes involves a defective amplification of the late-phase insulin response to glucose by glucose-dependent insulinotropic polypeptide-regardless of etiology and phenotype. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2003 , 88, 4897-903	5.6	175
6	The genetic abnormality in the beta cell determines the response to an oral glucose load. <i>Diabetologia</i> , 2002 , 45, 427-35	10.3	197
5	Novel MODY3 mutations in the hepatocyte nuclear factor-1alpha gene: evidence for a hyperexcitability of pancreatic beta-cells to intravenous secretagogues in a glucose-tolerant carrier of a P447L mutation. <i>Diabetes</i> , 1997 , 46, 726-30	0.9	88
4	Meta-analysis of exome array data identifies six novel genetic loci for lung function. <i>Wellcome Open Research</i> , 3, 4	4.8	1
3	The Polygenic and Monogenic Basis of Blood Traits and Diseases		3
2	Fine-mapping of an expanded set of type 2 diabetes loci to single-variant resolution using high-density imputation and islet-specific epigenome maps		18
1	Human pancreatic islet 3D chromatin architecture provides insights into the genetics of type 2 diabete	s	7