

Sophie Visvikis-Siest

List of Publications by Year in Descending Order

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Version: 2024-04-23

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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.

273
papers

24,949
citations

56
h-index

155
g-index

312
ext. papers

29,769
ext. citations

6.6
avg, IF

7.47
L-index

#	Paper	IF	Citations
273	The association of vascular endothelial growth factor related SNPs and circulating iron levels might depend on body mass index.. <i>Frontiers in Bioscience</i> , 2022 , 27, 27		1
272	A genetic determinant of VEGF-A levels is associated with telomere attrition. <i>Aging</i> , 2021 , 13, 23517-23526	5.26	1
271	Heterogeneous contributions of change in population distribution of body mass index to change in obesity and underweight. <i>ELife</i> , 2021 , 10,	8.9	10
270	Effect of Mastiha supplementation on NAFLD: The MAST4HEALTH Randomised, Controlled Trial. <i>Molecular Nutrition and Food Research</i> , 2021 , 65, e2001178	5.9	6
269	Nutrigenetic Interactions Might Modulate the Antioxidant and Anti-Inflammatory Status in Mastiha-Supplemented Patients With NAFLD. <i>Frontiers in Immunology</i> , 2021 , 12, 683028	8.4	4
268	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. <i>Nature Communications</i> , 2021 , 12, 24	17.4	30
267	Genome-wide association study of circulating interleukin 6 levels identifies novel loci. <i>Human Molecular Genetics</i> , 2021 , 30, 393-409	5.6	6
266	Epigenome-wide association study detects a novel loci associated with central obesity in healthy subjects. <i>BMC Medical Genomics</i> , 2021 , 14, 233	3.7	0
265	Worldwide trends in hypertension prevalence and progress in treatment and control from 1990 to 2019: a pooled analysis of 1201 population-representative studies with 104 million participants. <i>Lancet, The</i> , 2021 , 398, 957-980	40	154
264	Dietary Patterns, Blood Pressure and the Glycemic and Lipidemic Profile of Two Teenage, European Populations. <i>Nutrients</i> , 2021 , 13,	6.7	2
263	Height and body-mass index trajectories of school-aged children and adolescents from 1985 to 2019 in 200 countries and territories: a pooled analysis of 2181 population-based studies with 65 million participants. <i>Lancet, The</i> , 2020 , 396, 1511-1524	40	73
262	Repositioning of the global epicentre of non-optimal cholesterol. <i>Nature</i> , 2020 , 582, 73-77	50.4	48
261	Epigenome-wide association study in healthy individuals identifies significant associations with DNA methylation and PBMC extract VEGF-A concentration. <i>Clinical Epigenetics</i> , 2020 , 12, 79	7.7	3
260	Increased risk of hypercholesterolemia in a French and Lebanese population due to an interaction between rs2569190 in CD14 and gender. <i>Clinica Chimica Acta</i> , 2020 , 509, 172-176	6.2	1
259	Variants Associated with Short Leukocyte Telomeres: Implication of Higher Early Life Leukocyte Telomere Attrition as Assessed by the Blood-and-Muscle Model. <i>Cells</i> , 2020 , 9,	7.9	2
258	Obesity status modifies the association between rs7556897T>C in the intergenic region SLC19A3-CCL20 and blood pressure in French children. <i>Clinical Chemistry and Laboratory Medicine</i> , 2020 , 58, 1819-1827	5.9	0
257	Milestones in Personalized Medicine: From the Ancient Time to Nowadays-the Provocation of COVID-19. <i>Frontiers in Genetics</i> , 2020 , 11, 569175	4.5	6

256	National trends in total cholesterol obscure heterogeneous changes in HDL and non-HDL cholesterol and total-to-HDL cholesterol ratio: a pooled analysis of 458 population-based studies in Asian and Western countries. <i>International Journal of Epidemiology</i> , 2020 , 49, 173-192	7.8	25
255	Telomere length determinants in childhood. <i>Clinical Chemistry and Laboratory Medicine</i> , 2020 , 58, 162-177	9	16
254	Rising rural body-mass index is the main driver of the global obesity epidemic in adults. <i>Nature</i> , 2019 , 569, 260-264	50.4	278
253	Genetic determinants of circulating haptoglobin concentration. <i>Clinica Chimica Acta</i> , 2019 , 494, 138-142	6.2	7
252	Epigenome-Wide Association Study (EWAS) of Blood Lipids in Healthy Population from STANISLAS Family Study (SFS). <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	11
251	Peripheral blood mononuclear cells extracts VEGF protein levels and VEGF mRNA: Associations with inflammatory molecules in a healthy population. <i>PLoS ONE</i> , 2019 , 14, e0220902	3.7	2
250	Pleiotropy of ABO gene: correlation of rs644234 with E-selectin and lipid levels. <i>Clinical Chemistry and Laboratory Medicine</i> , 2018 , 56, 748-754	5.9	7
249	Contributions of mean and shape of blood pressure distribution to worldwide trends and variations in raised blood pressure: a pooled analysis of 1018 population-based measurement studies with 88.6 million participants. <i>International Journal of Epidemiology</i> , 2018 , 47, 872-883i	7.8	40
248	The Relationship Between Vascular Endothelial Growth Factor Cis- and Trans-Acting Genetic Variants and Metabolic Syndrome. <i>American Journal of the Medical Sciences</i> , 2018 , 355, 559-565	2.2	6
247	Effect of SLCO1B1 gene polymorphisms and vitamin D on statin-induced myopathy. <i>Drug Metabolism and Personalized Therapy</i> , 2018 , 33, 41-47	2	5
246	The polymorphism rs6918289 located in the downstream region of the TREM2 gene is associated with TNF- α levels and IMT-F. <i>Scientific Reports</i> , 2018 , 8, 7160	4.9	1
245	Epistatic interaction of apolipoprotein E and lipolysis-stimulated lipoprotein receptor genetic variants is associated with Alzheimer's disease. <i>Neurobiology of Aging</i> , 2018 , 69, 292.e1-292.e5	5.6	3
244	Effect of LSR polymorphism on blood lipid levels and age-specific epistatic interaction with the APOE common polymorphism. <i>Clinical Genetics</i> , 2018 , 93, 846-852	4	3
243	A transnational collaborative network dedicated to the study and applications of the vascular endothelial growth factor-A in medical practice: the VEGF Consortium. <i>Clinical Chemistry and Laboratory Medicine</i> , 2018 , 56, 83-86	5.9	0
242	VEGF-A is associated with early degenerative changes in cartilage and subchondral bone. <i>Growth Factors</i> , 2018 , 36, 263-273	1.6	8
241	Personalised Medicine: The Odyssey from Hope to Practice. <i>Journal of Personalized Medicine</i> , 2018 , 8,	3.6	7
240	Genome Analyses of >200,000 Individuals Identify 58 Loci for Chronic Inflammation and Highlight Pathways that Link Inflammation and Complex Disorders. <i>American Journal of Human Genetics</i> , 2018 , 103, 691-706	11	151
239	The future of telomere length in personalized medicine. <i>Frontiers in Bioscience - Landmark</i> , 2018 , 23, 1628-1654	2.8	18

238	VEGF-related polymorphisms identified by GWAS and risk for major depression. <i>Translational Psychiatry</i> , 2017 , 7, e1055	8.6	23
237	VEGF, the underlying factor for metabolic syndrome; fact or fiction?. <i>Diabetes and Metabolic Syndrome: Clinical Research and Reviews</i> , 2017 , 11 Suppl 1, S61-S64	8.9	25
236	Worldwide trends in body-mass index, underweight, overweight, and obesity from 1975 to 2016: a pooled analysis of 2416 population-based measurement studies in 128.9 million children, adolescents, and adults. <i>Lancet, The</i> , 2017 , 390, 2627-2642	40	2980
235	Pharmacogenomic Challenges in Cardiovascular Diseases: Examples of Drugs and Considerations for Future Integration in Clinical Practice. <i>Current Pharmaceutical Biotechnology</i> , 2017 , 18, 231-241	2.6	6
234	Next generation sequencing and immuno-histochemistry profiling identify numerous biomarkers for personalized therapy of endometrioid endometrial carcinoma. <i>Clinical Chemistry and Laboratory Medicine</i> , 2017 , 56, e19-e22	5.9	1
233	IL6R haplotype rs4845625*T/rs4537545*C is a risk factor for simultaneously high CRP, LDL and ApoB levels. <i>Genes and Immunity</i> , 2017 , 18, 163-169	4.4	3
232	Worldwide trends in blood pressure from 1975 to 2015: a pooled analysis of 1479 population-based measurement studies with 19.1 million participants. <i>Lancet, The</i> , 2017 , 389, 37-55	40	1100
231	TREM-1 SNP rs2234246 regulates TREM-1 protein and mRNA levels and is associated with plasma levels of L-selectin. <i>PLoS ONE</i> , 2017 , 12, e0182226	3.7	16
230	Plasma VEGF-related polymorphisms are implicated in autoimmune thyroid diseases. <i>Autoimmunity</i> , 2016 , 49, 229-35	3	4
229	Six Novel Loci Associated with Circulating VEGF Levels Identified by a Meta-analysis of Genome-Wide Association Studies. <i>PLoS Genetics</i> , 2016 , 12, e1005874	6	43
228	La Pharmacogénomique, meilleur exemple de médecine personnalisée. <i>HEGEL - HEpato-GastroEntérologie Libérale</i> , 2016 , N° 1, 10	0.1	
227	Cardiovascular diseases-related C825T polymorphism has a significant sex-specific effect on serum soluble E-selectin levels. <i>Journal of Inflammation</i> , 2016 , 13, 39	6.7	2
226	Worldwide trends in diabetes since 1980: a pooled analysis of 751 population-based studies with 4.4 million participants. <i>Lancet, The</i> , 2016 , 387, 1513-1530	40	2039
225	Trends in adult body-mass index in 200 countries from 1975 to 2014: a pooled analysis of 1698 population-based measurement studies with 19.2 million participants. <i>Lancet, The</i> , 2016 , 387, 1377-1396	40	2787
224	Pro- and anti-angiogenic VEGF mRNAs in autoimmune thyroid diseases. <i>Autoimmunity</i> , 2016 , 49, 366-373		5
223	Effects of diabetes definition on global surveillance of diabetes prevalence and diagnosis: a pooled analysis of 96 population-based studies with 331,288 participants. <i>Lancet Diabetes and Endocrinology, the</i> , 2015 , 3, 624-37	18.1	109
222	Genetic determinants of leucocyte telomere length in children: a neglected and challenging field. <i>Paediatric and Perinatal Epidemiology</i> , 2015 , 29, 146-50	2.7	9
221	Genetic determined low response to thienopyridines is associated with higher systemic inflammation in smokers. <i>Pharmacogenomics</i> , 2015 , 16, 459-69	2.6	

220	Angiogenesis related genes NOS3, CD14, MMP3 and IL4R are associated to VEGF gene expression and circulating levels in healthy adults. <i>BMC Medical Genetics</i> , 2015 , 16, 90	2.1	19
219	Beyond genome-wide association studies: identifying variants using -omics approaches. <i>Personalized Medicine</i> , 2015 , 12, 529-531	2.2	
218	Low copy number of the salivary amylase gene predisposes to obesity. <i>Nature Genetics</i> , 2014 , 46, 492-7	36.3	177
217	Human cytochrome P450 epoxygenases: variability in expression and role in inflammation-related disorders. <i>Pharmacology & Therapeutics</i> , 2014 , 144, 134-61	13.9	66
216	Public biobanks: calculation and recovery of costs. <i>Science Translational Medicine</i> , 2014 , 6, 261fs45	17.5	39
215	Conference scene: pharmacogenomics: from cell to clinic (part 2). <i>Pharmacogenomics</i> , 2014 , 15, 739-44	2.6	0
214	Influence of inflammation on cardiovascular protective effects of cytochrome P450 epoxygenase-derived epoxyeicosatrienoic acids. <i>Drug Metabolism Reviews</i> , 2014 , 46, 33-56	7	19
213	Epistatic study reveals two genetic interactions in blood pressure regulation. <i>BMC Medical Genetics</i> , 2013 , 14, 2	2.1	10
212	Associations of vascular endothelial growth factor (VEGF) with adhesion and inflammation molecules in a healthy population. <i>Cytokine</i> , 2013 , 61, 602-7	4	21
211	Dairy product consumption, calcium intakes, and metabolic syndrome-related factors over 5 years in the STANISLAS study. <i>Nutrition</i> , 2013 , 29, 519-24	4.8	47
210	Relationship between catalase haplotype and arterial aging. <i>Atherosclerosis</i> , 2013 , 227, 100-5	3.1	11
209	Newly identified synergy between clopidogrel and calcium-channel blockers for blood pressure regulation possibly involves CYP2C19 rs4244285. <i>International Journal of Cardiology</i> , 2013 , 168, 3057-8	3.2	2
208	A common variant highly associated with plasma VEGFA levels also contributes to the variation of both LDL-C and HDL-C. <i>Journal of Lipid Research</i> , 2013 , 54, 535-41	6.3	21
207	Causal relationship between obesity and vitamin D status: bi-directional Mendelian randomization analysis of multiple cohorts. <i>PLoS Medicine</i> , 2013 , 10, e1001383	11.6	592
206	Mining the human phenome using allelic scores that index biological intermediates. <i>PLoS Genetics</i> , 2013 , 9, e1003919	6	58
205	Influence of genetic variations on levels of inflammatory markers of healthy subjects at baseline and one week after clopidogrel therapy; results of a preliminary study. <i>International Journal of Molecular Sciences</i> , 2013 , 14, 16402-13	6.3	0
204	What is the contribution of two genetic variants regulating VEGF levels to type 2 diabetes risk and to microvascular complications?. <i>PLoS ONE</i> , 2013 , 8, e55921	3.7	32
203	Association of TERC and OBFC1 haplotypes with mean leukocyte telomere length and risk for coronary heart disease. <i>PLoS ONE</i> , 2013 , 8, e83122	3.7	34

202	Genetic biomarkers of hypertension and future challenges integrating epigenomics. <i>Clinica Chimica Acta</i> , 2012 , 414, 259-65	6.2	29
201	Human formyl peptide receptor 1 C32T SNP interacts with age and is associated with blood pressure levels. <i>Clinica Chimica Acta</i> , 2012 , 413, 34-8	6.2	14
200	Clinical necessity of partitioning of human plasma haptoglobin reference intervals by recently-discovered rs2000999. <i>Clinica Chimica Acta</i> , 2012 , 413, 1618-24	6.2	13
199	High prevalence of metabolic syndrome in Iran in comparison with France: what are the components that explain this?. <i>Metabolic Syndrome and Related Disorders</i> , 2012 , 10, 181-8	2.6	42
198	A genome-wide association search for type 2 diabetes genes in African Americans. <i>PLoS ONE</i> , 2012 , 7, e29202	3.7	138
197	Genome-wide association for abdominal subcutaneous and visceral adipose reveals a novel locus for visceral fat in women. <i>PLoS Genetics</i> , 2012 , 8, e1002695	6	199
196	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
195	Influence of pre-analytical variables on VEGF gene expression and circulating protein concentrations. <i>Biopreservation and Biobanking</i> , 2012 , 10, 454-61	2.1	14
194	Novel association approach for variable number tandem repeats (VNTRs) identifies DOCK5 as a susceptibility gene for severe obesity. <i>Human Molecular Genetics</i> , 2012 , 21, 3727-38	5.6	29
193	Alcohol Consumption, Beverage Preference, and Diet in Middle-Aged Men from the STANISLAS Study. <i>Journal of Nutrition and Metabolism</i> , 2012 , 2012, 987243	2.7	13
192	Need for pharmacogenomic information also for generic medications: recommendation of the European Society of Pharmacogenomics and Theranostics (ESPT). <i>Drug Metabolism and Drug Interactions</i> , 2012 , 27, 119		
191	Genome-wide meta-analysis points to CTC1 and ZNF676 as genes regulating telomere homeostasis in humans. <i>Human Molecular Genetics</i> , 2012 , 21, 5385-94	5.6	162
190	A genome-wide association study identifies rs2000999 as a strong genetic determinant of circulating haptoglobin levels. <i>PLoS ONE</i> , 2012 , 7, e32327	3.7	27
189	Functional epistatic interaction between rs6046G>A in F7 and rs5355C>T in SELE modifies systolic blood pressure levels. <i>PLoS ONE</i> , 2012 , 7, e40777	3.7	7
188	Functional genomics towards personalized healthcare and systems medicine. <i>Personalized Medicine</i> , 2011 , 8, 227-242	2.2	1
187	Childhood obesity is associated with shorter leukocyte telomere length. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2011 , 96, 1500-5	5.6	104
186	Mirror extreme BMI phenotypes associated with gene dosage at the chromosome 16p11.2 locus. <i>Nature</i> , 2011 , 478, 97-102	50.4	322
185	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. <i>Nature Genetics</i> , 2011 , 43, 1131-8	36.3	415

184	Klotho KL-VS genotype is involved in blood pressure regulation. <i>Clinica Chimica Acta</i> , 2011 , 412, 1773-7	6.2	17
183	Cardiovascular diseases and genome-wide association studies. <i>Clinica Chimica Acta</i> , 2011 , 412, 1697-701	6.2	19
182	The Lin28/let-7 axis regulates glucose metabolism. <i>Cell</i> , 2011 , 147, 81-94	56.2	649
181	Genome-wide association identifies nine common variants associated with fasting proinsulin levels and provides new insights into the pathophysiology of type 2 diabetes. <i>Diabetes</i> , 2011 , 60, 2624-34	0.9	285
180	Genome-wide association analysis identifies variants associated with nonalcoholic fatty liver disease that have distinct effects on metabolic traits. <i>PLoS Genetics</i> , 2011 , 7, e1001324	6	629
179	Substantial variation in qPCR measured mean blood telomere lengths in young men from eleven European countries. <i>American Journal of Human Biology</i> , 2011 , 23, 228-31	2.7	33
178	Availability of pharmacogenetic and pharmacogenomic information in anticancer drug monographs in France: personalized cancer therapy. <i>Pharmacogenomics</i> , 2011 , 12, 681-91	2.6	4
177	Identification of cis- and trans-acting genetic variants explaining up to half the variation in circulating vascular endothelial growth factor levels. <i>Circulation Research</i> , 2011 , 109, 554-63	15.7	57
176	Association of genetic Loci with glucose levels in childhood and adolescence: a meta-analysis of over 6,000 children. <i>Diabetes</i> , 2011 , 60, 1805-12	0.9	83
175	Biological and genetic factors associated with ABCB1 and pregnane-X-receptor expressions in peripheral blood mononuclear cells in the STANISLAS cohort. <i>Drug Metabolism and Drug Interactions</i> , 2011 , 26, 27-32		3
174	Expression of inflammatory molecules and associations with BMI in children. <i>European Journal of Clinical Investigation</i> , 2010 , 40, 388-92	4.6	20
173	Sex-dependent associations of leptin with metabolic syndrome-related variables: the Stanislas study. <i>Obesity</i> , 2010 , 18, 196-201	8	20
172	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
171	Prevalence of loss-of-function FTO mutations in lean and obese individuals. <i>Diabetes</i> , 2010 , 59, 311-8	0.9	83
170	Interactions of dietary whole-grain intake with fasting glucose- and insulin-related genetic loci in individuals of European descent: a meta-analysis of 14 cohort studies. <i>Diabetes Care</i> , 2010 , 33, 2684-91	14.6	112
169	Association study of gene polymorphisms involved in vascular alterations in elderly hypertensives with subjective memory complaints. <i>Dementia and Geriatric Cognitive Disorders</i> , 2010 , 30, 440-8	2.6	10
168	Metabolic syndrome-related composite factors over 5 years in the STANISLAS family study: genetic heritability and common environmental influences. <i>Clinica Chimica Acta</i> , 2010 , 411, 833-9	6.2	11
167	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

166	Association between angiotensin II type 1 receptor gene polymorphism and metabolic syndrome in a young female Iranian population. <i>Archives of Medical Research</i> , 2010 , 41, 343-9	6.6	13
165	A multi-stage multi-design strategy provides strong evidence that the BAI3 locus is associated with early-onset venous thromboembolism. <i>Journal of Thrombosis and Haemostasis</i> , 2010 , 8, 2671-9	15.4	35
164	Visfatin: the link between inflammation and childhood obesity. <i>Diabetes Care</i> , 2009 , 32, e71	14.6	14
163	Personalized therapy and pharmacogenomics: future perspective. <i>Pharmacogenomics</i> , 2009 , 10, 927-30	2.6	6
162	Capillary isotachopheresis study of lipoprotein network sensitive to apolipoprotein E phenotype. 2. ApoE and apoC-III relations in triglyceride clearance. <i>Molecular and Cellular Biochemistry</i> , 2009 , 325, 25-40	4.2	2
161	Capillary isotachopheresis study of lipoprotein network sensitive to apolipoprotein E phenotype. 1. ApoE distribution between lipoproteins. <i>Molecular and Cellular Biochemistry</i> , 2009 , 325, 41-51	4.2	3
160	Genetic profiling of human cell lines used as in vitro model to study cardiovascular pathophysiology and pharmacotoxicology. <i>Cell Biology and Toxicology</i> , 2009 , 25, 561-71	7.4	5
159	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
158	Genome-wide association study for early-onset and morbid adult obesity identifies three new risk loci in European populations. <i>Nature Genetics</i> , 2009 , 41, 157-9	36.3	521
157	Association between EGF and lipid concentrations: a benefit role in the atherosclerotic process?. <i>Clinica Chimica Acta</i> , 2009 , 402, 196-8	6.2	8
156	Association of ABCB1 gene polymorphisms with plasma lipid and apolipoprotein concentrations in the STANISLAS cohort. <i>Clinica Chimica Acta</i> , 2009 , 403, 198-202	6.2	20
155	Adipokine expression in adipose tissue and in peripheral blood mononuclear cells in children Correlation with BMI and fatty acid content. <i>Clinica Chimica Acta</i> , 2009 , 410, 85-9	6.2	6
154	Association of human cathelicidin (hCAP-18/LL-37) gene expression with cardiovascular disease risk factors. <i>Nutrition, Metabolism and Cardiovascular Diseases</i> , 2009 , 19, 720-8	4.5	24
153	Human formyl peptide receptor 1 (FPR1) c.32C>T SNP is associated with decreased soluble E-selectin levels. <i>Pharmacogenomics</i> , 2009 , 10, 951-9	2.6	8
152	Parental precocious influences on offspring cardiovascular risk markers: an exploratory study in the STANISLAS Cohort. <i>Personalized Medicine</i> , 2009 , 6, 343-352	2.2	
151	Candidate gene microarray analysis in peripheral blood cells for studying hypertension/obesity. <i>Personalized Medicine</i> , 2009 , 6, 269-291	2.2	4
150	Functional genomics towards personalized healthcare. <i>Personalized Medicine</i> , 2009 , 6, 19-32	2.2	3
149	Drug metabolizing enzymes and transporters mRNA in peripheral blood mononuclear cells of healthy subjects: biological variations and importance of pre-analytical steps. <i>Current Drug Metabolism</i> , 2009 , 10, 410-9	3.5	5

148	Genomics and the prospects of existing and emerging therapeutics for cardiovascular diseases. <i>Current Pharmaceutical Design</i> , 2009 , 15, 3193-206	3.3	12
147	Five-year alterations in BMI are associated with clustering of changes in cardiovascular risk factors in a gender-dependant way: the Stanislas study. <i>International Journal of Obesity</i> , 2008 , 32, 1279-88	5.5	24
146	Genetic Polymorphism of CYP2C19 gene in the Stanislas cohort. A link with inflammation. <i>Annals of Human Genetics</i> , 2008 , 72, 178-83	2.2	31
145	P-selectin polymorphisms influences on P-selectin serum concentrations and on their familial correlation: the STANISLAS family study. <i>Journal of Thrombosis and Haemostasis</i> , 2008 , 6, 920-7	15.4	8
144	Visfatin, low-grade inflammation and body mass index (BMI). <i>Clinical Endocrinology</i> , 2008 , 69, 568-74	3.4	33
143	Statins as effectors of key activities involved in apoE-dependent VLDL metabolism: review and hypothesis. <i>Vascular Pharmacology</i> , 2008 , 48, 70-5	5.9	7
142	Pharmacogenomics and Cardiovascular Drugs. <i>Methods in Pharmacology and Toxicology</i> , 2008 , 413-446	1.1	
141	Leptin expression in Peripheral Blood Mononuclear Cells (PBMCs) is related with blood pressure variability. <i>Clinica Chimica Acta</i> , 2008 , 395, 47-50	6.2	15
140	Transcription factor and drug-metabolizing enzyme gene expression in lymphocytes from healthy human subjects. <i>Drug Metabolism and Disposition</i> , 2008 , 36, 182-9	4	71
139	A new single nucleotide polymorphism genotyping method based on gap ligase chain reaction and a microsphere detection assay. <i>Clinical Chemistry and Laboratory Medicine</i> , 2008 , 46, 486-9	5.9	4
138	The STANISLAS Cohort: a 10-year follow-up of supposed healthy families. Gene-environment interactions, reference values and evaluation of biomarkers in prevention of cardiovascular diseases. <i>Clinical Chemistry and Laboratory Medicine</i> , 2008 , 46, 733-47	5.9	45
137	Genetic profiling in healthy subjects from the Stanislas cohort based on 24 polymorphisms: effects on biological variables. <i>Clinical Chemistry and Laboratory Medicine</i> , 2008 , 46, 64-72	5.9	1
136	The association of telomere length with paternal history of premature myocardial infarction in the European Atherosclerosis Research Study II. <i>Journal of Molecular Medicine</i> , 2008 , 86, 815-24	5.5	47
135	The lipoprotein lipase serine 447 stop polymorphism is associated with altered serum carotenoid concentrations in the Stanislas Family Study. <i>Journal of the American College of Nutrition</i> , 2007 , 26, 655-62	2.5	24
134	Peripheral blood mononuclear cells (PBMCs): a possible model for studying cardiovascular biology systems. <i>Clinical Chemistry and Laboratory Medicine</i> , 2007 , 45, 1154-68	5.9	45
133	From human genetic variations to prediction of risks and responses to drugs and the environment. <i>Personalized Medicine</i> , 2007 , 4, 95-104	2.2	2
132	Heritability of serum hs-CRP concentration and 5-year changes in the Stanislas family study: association with apolipoprotein E alleles. <i>Genes and Immunity</i> , 2007 , 8, 352-9	4.4	10
131	Lack of association between EGF 61A>G polymorphism and plasma EGF concentration in the STANISLAS family study. <i>Journal of Investigative Dermatology</i> , 2007 , 127, 969-70	4.3	1

130	Heritability for plasma VEGF concentration in the Stanislas family study. <i>Annals of Human Genetics</i> , 2007 , 71, 54-63	2.2	31
129	Determination of ABCB1 polymorphisms and haplotypes frequencies in a French population. <i>Fundamental and Clinical Pharmacology</i> , 2007 , 21, 411-8	3.1	36
128	Compared effect of immunosuppressive drugs cyclosporine A and rapamycin on cholesterol homeostasis key enzymes CYP27A1 and HMG-CoA reductase. <i>Basic and Clinical Pharmacology and Toxicology</i> , 2007 , 100, 392-7	3.1	32
127	Association of classical and related inflammatory markers with high-sensitivity C-reactive protein in healthy individuals: results from the Stanislas cohort. <i>Clinical Chemistry and Laboratory Medicine</i> , 2007 , 45, 1339-46	5.9	11
126	Effect of HMGCoA reductase inhibitors on cytochrome P450 expression in endothelial cell line. <i>Journal of Cardiovascular Pharmacology</i> , 2007 , 49, 306-15	3.1	20
125	Enzymes and pharmacogenetics of cardiovascular drugs. <i>Clinica Chimica Acta</i> , 2007 , 381, 26-31	6.2	31
124	Analysis of the effect of multiple genetic variants of cardiovascular disease risk on insulin concentration variability in healthy adults of the STANISLAS cohort. The role of FGB-455 G/A polymorphism. <i>Atherosclerosis</i> , 2007 , 191, 369-76	3.1	12
123	Association between TNF and IL-1 bloc polymorphisms and plasma MCP-1 concentration. <i>Atherosclerosis</i> , 2007 , 192, 348-53	3.1	10
122	Pharmacogenomics and antihypertensive drugs: a path toward personalized medicine. <i>Personalized Medicine</i> , 2007 , 4, 393-412	2.2	4
121	Natriuretic peptide Val7Met substitution and risk of coronary artery disease in Greek patients with familial hypercholesterolemia. <i>Journal of Clinical Laboratory Analysis</i> , 2006 , 20, 98-104	3	10
120	Biological determinants of and reference values for plasma interleukin-8, monocyte chemoattractant protein-1, epidermal growth factor, and vascular endothelial growth factor: Results from the STANISLAS cohort. <i>Clinical Chemistry</i> , 2006 , 52, 504-10	5.5	51
119	Polymorphisms associated with apolipoprotein B levels in Greek patients with familial hypercholesterolemia. <i>Clinical Chemistry and Laboratory Medicine</i> , 2006 , 44, 799-806	5.9	3
118	Different genes and polymorphisms affecting high-density lipoprotein cholesterol levels in Greek familial hypercholesterolemia patients. <i>Genetic Testing and Molecular Biomarkers</i> , 2006 , 10, 192-9		10
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