Geoffrey S Ginsburg

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.

11,646 246 56 100 h-index g-index citations papers 8.2 6.36 289 13,953 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
246	Systematic comparison of published host gene expression signatures for bacterial/viral discrimination <i>Genome Medicine</i> , 2022 , 14, 18	14.4	3
245	Prospective Validation of a Rapid Host Gene Expression Test to Discriminate Bacterial From Viral Respiratory Infection <i>JAMA Network Open</i> , 2022 , 5, e227299	10.4	1
244	Harnessing the Power of Artificial Intelligence in Otolaryngology and the Communication Sciences JARO - Journal of the Association for Research in Otolaryngology, 2022, 23, 319	3.3	O
243	Comparing the Diagnostic Accuracy of Clinician Judgment to a Novel Host Response Diagnostic for Acute Respiratory Illness. <i>Open Forum Infectious Diseases</i> , 2021 , 8, ofab564	1	0
242	Genomic medicine year in review: 2021. American Journal of Human Genetics, 2021, 108, 2210-2214	11	1
241	The National Academies Roundtable on Genomics and Precision Health: Where we have been and where we are heading. <i>American Journal of Human Genetics</i> , 2021 , 108, 1817-1822	11	2
240	The Host Response to Viral Infections Reveals Common and Virus-Specific Signatures in the Peripheral Blood. <i>Frontiers in Immunology</i> , 2021 , 12, 741837	8.4	3
239	Establishing the value of genomics in medicine: the IGNITE Pragmatic Trials Network. <i>Genetics in Medicine</i> , 2021 , 23, 1185-1191	8.1	3
238	Clinical implementation of an oncology-specific family health history risk assessment tool. <i>Hereditary Cancer in Clinical Practice</i> , 2021 , 19, 20	2.3	0
237	Discriminating Bacterial and Viral Infection Using a Rapid Host Gene Expression Test. <i>Critical Care Medicine</i> , 2021 , 49, 1651-1663	1.4	11
236	An atlas connecting shared genetic architecture of human diseases and molecular phenotypes provides insight into COVID-19 susceptibility. <i>Genome Medicine</i> , 2021 , 13, 83	14.4	8
235	Association of Hepatic Steatosis With Major Adverse Cardiovascular Events, Independent of Coronary Artery Disease. <i>Clinical Gastroenterology and Hepatology</i> , 2021 , 19, 1480-1488.e14	6.9	20
234	A blood-based host gene expression assay for early detection of respiratory viral infection: an index-cluster prospective cohort study. <i>Lancet Infectious Diseases, The</i> , 2021 , 21, 396-404	25.5	13
233	Platelet reactivity in response to aspirin and ticagrelor in African-Americans and European-Americans. <i>Journal of Thrombosis and Thrombolysis</i> , 2021 , 51, 249-259	5.1	2
232	Family history assessment significantly enhances delivery of precision medicine in the genomics era. <i>Genome Medicine</i> , 2021 , 13, 3	14.4	6
231	Validation of a Host Gene Expression Test for Bacterial/Viral Discrimination in Immunocompromised Hosts. <i>Clinical Infectious Diseases</i> , 2021 , 73, 605-613	11.6	5
230	Association of Metabolic Phenotypes With Coronary Artery Disease and Cardiovascular Events in Patients With Stable Chest Pain. <i>Diabetes Care</i> , 2021 , 44, 1038-1045	14.6	1

(2020-2021)

229	Dysregulated transcriptional responses to SARS-CoV-2 in the periphery. <i>Nature Communications</i> , 2021 , 12, 1079	17.4	25	
228	The host transcriptional response to Candidemia is dominated by neutrophil activation and heme biosynthesis and supports novel diagnostic approaches. <i>Genome Medicine</i> , 2021 , 13, 108	14.4	3	
227	Strategies to Integrate Genomic Medicine into Clinical Care: Evidence from the IGNITE Network. Journal of Personalized Medicine, 2021 , 11,	3.6	1	
226	Utility of High-Sensitivity Troponin Among Stable Patients With Chest Pain Undergoing Stress Imaging (from PROMISE). <i>American Journal of Cardiology</i> , 2021 , 158, 148-149	3		
225	Adaptive Multi-Channel Event Segmentation and Feature Extraction for Monitoring Health Outcomes. <i>IEEE Transactions on Biomedical Engineering</i> , 2021 , 68, 2377-2388	5	1	
224	Heparin-based blood purification attenuates organ injury in baboons with pneumonia. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2021 , 321, L321-L335	5.8	1	
223	Assessment of the Feasibility of Using Noninvasive Wearable Biometric Monitoring Sensors to Detect Influenza and the Common Cold Before Symptom Onset. <i>JAMA Network Open</i> , 2021 , 4, e212853	34 ^{0.4}	4	
222	Welcome to the 18th volume of. <i>Personalized Medicine</i> , 2021 , 18, 1-3	2.2		
221	Digital Health-The Need to Assess Benefits, Risks, and Value. <i>JAMA - Journal of the American Medical Association</i> , 2021 , 325, 127-128	27.4	13	
220	A comparison of host response strategies to distinguish bacterial and viral infection <i>PLoS ONE</i> , 2021 , 16, e0261385	3.7	1	
219	The International Hundred Thousand Plus Cohort Consortium: integrating large-scale cohorts to address global scientific challenges. <i>The Lancet Digital Health</i> , 2020 , 2, e567-e568	14.4	9	
218	The Project Baseline Health Study: a step towards a broader mission to map human health. <i>Npj Digital Medicine</i> , 2020 , 3, 84	15.7	10	
217	Evaluation of family health history collection methods impact on data and risk assessment outcomes. <i>Preventive Medicine Reports</i> , 2020 , 18, 101072	2.6	4	
216	Previously Derived Host Gene Expression Classifiers Identify Bacterial and Viral Etiologies of Acute Febrile Respiratory Illness in a South Asian Population. <i>Open Forum Infectious Diseases</i> , 2020 , 7, ofaa194	1 ¹	1	
215	Enabling Technologies for Personalized and Precision Medicine. <i>Trends in Biotechnology</i> , 2020 , 38, 497-	5 1§ .1	71	
214	Promoting Wellness Through Mobile Health Technology in a College Student Population: Protocol Development and Pilot Study. <i>JMIR Research Protocols</i> , 2020 , 9, e16474	2	O	
213	Dysregulated transcriptional responses to SARS-CoV-2 in the periphery support novel diagnostic approaches 2020 ,		2	
212	1218. Comparing the Diagnostic Accuracy of Clinician Judgement to a Novel Host Response Diagnostic for Acute Respiratory Illness. <i>Open Forum Infectious Diseases</i> , 2020 , 7, S630-S630	1	_	

211	1226. Performance of a Host Response Test for Bacterial/Viral Discrimination in Immunocompromised Patients. <i>Open Forum Infectious Diseases</i> , 2020 , 7, S633-S634	1	
210	Chromatin remodeling in peripheral blood cells reflects COVID-19 symptom severity 2020 ,		4
209	An atlas connecting shared genetic architecture of human diseases and molecular phenotypes provides insight into COVID-19 susceptibility 2020 ,		19
208	Influence of Sex on Platelet Reactivity in Response to Aspirin. <i>Journal of the American Heart Association</i> , 2020 , 9, e014726	6	6
207	Genomic Medicine Year in Review: 2020. American Journal of Human Genetics, 2020, 107, 1007-1010	11	2
206	At the intersection of precision medicine and population health: an implementation-effectiveness study of family health history based systematic risk assessment in primary care. <i>BMC Health Services Research</i> , 2020 , 20, 1015	2.9	6
205	Single-Molecule hsTnI and Short-Term Risk in Stable Patients With Chest Pain. <i>Journal of the American College of Cardiology</i> , 2019 , 73, 251-260	15.1	19
204	Pilot study of myocardial ischemia-induced metabolomic changes in emergency department patients undergoing stress testing. <i>PLoS ONE</i> , 2019 , 14, e0211762	3.7	4
203	Primary Care PhysiciansCKnowledge, Attitudes, and Experience with Personal Genetic Testing. Journal of Personalized Medicine, 2019 , 9,	3.6	24
202	A Global Collaborative to Advance Genomic Medicine. <i>American Journal of Human Genetics</i> , 2019 , 104, 407-409	11	8
201	A host gene expression approach for identifying triggers of asthma exacerbations. <i>PLoS ONE</i> , 2019 , 14, e0214871	3.7	6
200	Implementation, adoption, and utility of family health history risk assessment in diverse care settings: evaluating implementation processes and impact with an implementation framework. <i>Genetics in Medicine</i> , 2019 , 21, 331-338	8.1	17
199	Opportunities to implement a sustainable genomic medicine program: lessons learned from the IGNITE Network. <i>Genetics in Medicine</i> , 2019 , 21, 743-747	8.1	24
198	Opportunities, resources, and techniques for implementing genomics in clinical care. <i>Lancet, The</i> , 2019 , 394, 511-520	4º	30
197	Family health history: underused for actionable risk assessment. <i>Lancet, The</i> , 2019 , 394, 596-603	40	35
196	Specific Immunologic Countermeasure Protocol for Deep-Space Exploration Missions. <i>Frontiers in Immunology</i> , 2019 , 10, 2407	8.4	16
195	Facilitating translational team science: The project leader model. <i>Journal of Clinical and Translational Science</i> , 2019 , 3, 140-146	0.4	4
194	Validation of a host response test to distinguish bacterial and viral respiratory infection. <i>EBioMedicine</i> , 2019 , 48, 453-461	8.8	22

(2018-2019)

193	Rapid, Sample-to-Answer Host Gene Expression Test to Diagnose Viral Infection. <i>Open Forum Infectious Diseases</i> , 2019 , 6, ofz466	1	5
192	1330. Evaluation of Multiple Host Response-Based Strategies to Classify Acute Respiratory Illness. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S481-S481	1	78
191	2595. Murine Models for the Host Response to Typical and Atypical Pneumonia. <i>Open Forum Infectious Diseases</i> , 2019 , 6, S902-S902	1	78
190	Genomic Medicine Year in Review: 2019. American Journal of Human Genetics, 2019, 105, 1072-1075	11	7
189	Effect of Sociodemographic Factors on Uptake of a Patient-Facing Information Technology Family Health History Risk Assessment Platform. <i>Applied Clinical Informatics</i> , 2019 , 10, 180-188	3.1	5
188	Qualitative study of system-level factors related to genomic implementation. <i>Genetics in Medicine</i> , 2019 , 21, 1534-1540	8.1	12
187	High-Sensitivity Troponin I and Coronary Computed Tomography in Symptomatic Outpatients With Suspected CAD: Insights From the PROMISE Trial. <i>JACC: Cardiovascular Imaging</i> , 2019 , 12, 1047-10)5 ⁸ 5 ⁴	22
186	Research Directions in the Clinical Implementation of Pharmacogenomics: An Overview of US Programs and Projects. <i>Clinical Pharmacology and Therapeutics</i> , 2018 , 103, 778-786	6.1	63
185	A community approach to mortality prediction in sepsis via gene expression analysis. <i>Nature Communications</i> , 2018 , 9, 694	17.4	106
184	Unsupervised Analysis of Transcriptomics in Bacterial Sepsis Across Multiple Datasets Reveals Three Robust Clusters. <i>Critical Care Medicine</i> , 2018 , 46, 915-925	1.4	115
183	Immune System Dysregulation During Spaceflight: Potential Countermeasures for Deep Space Exploration Missions. <i>Frontiers in Immunology</i> , 2018 , 9, 1437	8.4	132
182	Precision Medicine: From Science To Value. <i>Health Affairs</i> , 2018 , 37, 694-701	7	250
181	A proposed approach to accelerate evidence generation for genomic-based technologies in the context of a learning health system. <i>Genetics in Medicine</i> , 2018 , 20, 390-396	8.1	12
180	2015. Host Gene Expression Identifies Infectious Triggers of Asthma Exacerbation. <i>Open Forum Infectious Diseases</i> , 2018 , 5, S587-S587	1	78
179	2014. TLDA Validation of a Host Response Signature to Discriminate Bacterial, Viral, and Non-infectious Causes of Illness. <i>Open Forum Infectious Diseases</i> , 2018 , 5, S587-S587	1	1
178	2018. Host Gene Expression Classifiers Distinguish Bacterial and Viral Infections in Sri Lankan Patients with Acute Febrile Respiratory Illness. <i>Open Forum Infectious Diseases</i> , 2018 , 5, S588-S588	1	78
177	A miRNA Host Response Signature Accurately Discriminates Acute Respiratory Infection Etiologies. <i>Frontiers in Microbiology</i> , 2018 , 9, 2957	5.7	7
176	Insurance Coverage Policies for Pharmacogenomic and Multi-Gene Testing for Cancer. <i>Journal of Personalized Medicine</i> , 2018 , 8,	3.6	22

175	RAB11FIP5 Expression and Altered Natural Killer Cell Function Are Associated with Induction of HIV Broadly Neutralizing Antibody Responses. <i>Cell</i> , 2018 , 175, 387-399.e17	56.2	42
174	A crowdsourced analysis to identify ab initio molecular signatures predictive of susceptibility to viral infection. <i>Nature Communications</i> , 2018 , 9, 4418	17.4	9
173	An age- and sex-specific gene expression score is associated with revascularization and coronary artery disease: Insights from the Prospective Multicenter Imaging Study for Evaluation of Chest Pain (PROMISE) trial. <i>American Heart Journal</i> , 2017 , 184, 133-140	4.9	10
172	Human genetic and metabolite variation reveals that methylthioadenosine is a prognostic biomarker and an inflammatory regulator in sepsis. <i>Science Advances</i> , 2017 , 3, e1602096	14.3	35
171	Nasopharyngeal Protein Biomarkers of Acute Respiratory Virus Infection. <i>EBioMedicine</i> , 2017 , 17, 172-1	881. 8	10
170	Primary care providers Quise of pharmacist support for delivery of pharmacogenetic testing. <i>Pharmacogenomics</i> , 2017 , 18, 359-367	2.6	17
169	Making genomic medicine evidence-based and patient-centered: a structured review and landscape analysis of comparative effectiveness research. <i>Genetics in Medicine</i> , 2017 , 19, 1081-1091	8.1	42
168	Local and Global Challenges in the Clinical Implementation of Precision Medicine 2017 , 105-117		
167	Integrative health care services utilization as a function of body mass index: A BraveNet practice-based research network study. <i>Advances in Integrative Medicine</i> , 2017 , 4, 14-21	1.6	4
166	Genomics-Inspired Biomarkers and Diagnostics-Where Are They?. Clinical Chemistry, 2017, 63, 255-257	5.5	1
165	Challenges and strategies for implementing genomic services in diverse settings: experiences from the Implementing GeNomics In pracTicE (IGNITE) network. <i>BMC Medical Genomics</i> , 2017 , 10, 35	3.7	61
164	Impact of Genetic Testing and Family Health History Based Risk Counseling on Behavior Change and Cognitive Precursors for Type 2 Diabetes. <i>Journal of Genetic Counseling</i> , 2017 , 26, 133-140	2.5	16
163	The effective rate of influenza reassortment is limited during human infection. <i>PLoS Pathogens</i> , 2017 , 13, e1006203	7.6	27
162	Patient experiences with pharmacogenetic testing in a primary care setting. <i>Pharmacogenomics</i> , 2016 , 17, 1629-1636	2.6	24
161	Developing Biomarker Arrays Predicting Sleep and Circadian-Coupled Risks to Health. <i>Sleep</i> , 2016 , 39, 727-36	1.1	59
160	An individualized predictor of health and disease using paired reference and target samples. <i>BMC Bioinformatics</i> , 2016 , 17, 47	3.6	29
159	Host gene expression classifiers diagnose acute respiratory illness etiology. <i>Science Translational Medicine</i> , 2016 , 8, 322ra11	17.5	150
158	Opportunities for the Cardiovascular Community in the Precision Medicine Initiative. <i>Circulation</i> , 2016 , 133, 226-31	16.7	42

(2015-2016)

157	Transcriptomic Analysis of the Host Response and Innate Resilience to Enterotoxigenic Escherichia coli Infection in Humans. <i>Journal of Infectious Diseases</i> , 2016 , 213, 1495-504	7	9
156	Genomics, clinical research, and learning health care systems: Strategies to improve patient care. <i>Nursing Outlook</i> , 2016 , 64, 225-8	2.7	3
155	Clinical utility of a Web-enabled risk-assessment and clinical decision support program. <i>Genetics in Medicine</i> , 2016 , 18, 1020-8	8.1	27
154	A Genomic Signature of Influenza Infection Shows Potential for Presymptomatic Detection, Guiding Early Therapy, and Monitoring Clinical Responses. <i>Open Forum Infectious Diseases</i> , 2016 , 3, ofw007	1	24
153	The IGNITE network: a model for genomic medicine implementation and research. <i>BMC Medical Genomics</i> , 2016 , 9, 1	3.7	139
152	Philanthropy for Science: Is It a Viable Option?. <i>Circulation Research</i> , 2016 , 119, 1057-1059	15.7	8
151	Systems Pharmacogenomics Finds RUNX1 Is an Aspirin-Responsive Transcription Factor Linked to Cardiovascular Disease and Colon Cancer. <i>EBioMedicine</i> , 2016 , 11, 157-164	8.8	17
150	Realizing the Full Potential of Precision Medicine in Health and Health Care. <i>JAMA - Journal of the American Medical Association</i> , 2016 , 316, 1659-1660	27.4	51
149	Deep Sequencing of Influenza A Virus from a Human Challenge Study Reveals a Selective Bottleneck and Only Limited Intrahost Genetic Diversification. <i>Journal of Virology</i> , 2016 , 90, 11247-112	58 ^{.6}	68
148	Use of a patient-entered family health history tool with decision support in primary care: impact of identification of increased risk patients on genetic counseling attendance. <i>Journal of Genetic Counseling</i> , 2015 , 24, 179-88	2.5	20
147	What was old is new again: using the host response to diagnose infectious disease. <i>Expert Review of Molecular Diagnostics</i> , 2015 , 15, 1143-58	3.8	20
146	Gene Expression Signatures and the Spectrum of Coronary Artery Disease. <i>Journal of Cardiovascular Translational Research</i> , 2015 , 8, 339-52	3.3	9
145	Patient beliefs and behaviors about genomic risk for type 2 diabetes: implications for prevention. Journal of Health Communication, 2015 , 20, 728-35	2.5	3
144	Does Type 2 Diabetes Genetic Testing and Counseling Reduce Modifiable Risk Factors? A Randomized Controlled Trial of Veterans. <i>Journal of General Internal Medicine</i> , 2015 , 30, 1591-8	4	25
143	Aligning incentives to fulfil the promise of personalised medicine. <i>Lancet, The</i> , 2015 , 385, 2118-9	40	55
142	A Guide for a Cardiovascular Genomics Biorepository: the CATHGEN Experience. <i>Journal of Cardiovascular Translational Research</i> , 2015 , 8, 449-57	3.3	45
141	Making Personalized Health Care Even More Personalized: Insights From Activities of the IOM Genomics Roundtable. <i>Annals of Family Medicine</i> , 2015 , 13, 373-80	2.9	30
140	SLCO1B1 genetic variants, long-term low-density lipoprotein cholesterol levels and clinical events in patients following cardiac catheterization. <i>Pharmacogenomics</i> , 2015 , 16, 449-58	2.6	14

139	Perspectives on genetic and genomic technologies in an academic medical center: the duke experience. <i>Journal of Personalized Medicine</i> , 2015 , 5, 67-82	3.6	13
138	Gene Expression Profiles Link Respiratory Viral Infection, Platelet Response to Aspirin, and Acute Myocardial Infarction. <i>PLoS ONE</i> , 2015 , 10, e0132259	3.7	19
137	Global implementation of genomic medicine: We are not alone. <i>Science Translational Medicine</i> , 2015 , 7, 290ps13	17.5	112
136	Perceptions of Personalized Medicine in an Academic Health System: Educational Findings. <i>Journal of Contemporary Medical Education</i> , 2015 , 3, 14-19	2.5	11
135	Quality of family history collection with use of a patient facing family history assessment tool. <i>BMC Family Practice</i> , 2014 , 15, 31	2.6	32
134	Multiplex detection of disease biomarkers using SERS molecular sentinel-on-chip. <i>Analytical and Bioanalytical Chemistry</i> , 2014 , 406, 3335-44	4.4	40
133	Providing patient education: impact on quantity and quality of family health history collection. <i>Familial Cancer</i> , 2014 , 13, 325-32	3	27
132	Implementing family health history risk stratification in primary care: impact of guideline criteria on populations and resource demand. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics,</i> 2014 , 166C, 24-33	3.1	33
131	The current epidemiology and clinical decisions surrounding acute respiratory infections. <i>Trends in Molecular Medicine</i> , 2014 , 20, 579-88	11.5	34
130	Genetically guided statin therapy on statin perceptions, adherence, and cholesterol lowering: a pilot implementation study in primary care patients. <i>Journal of Personalized Medicine</i> , 2014 , 4, 147-62	3.6	26
129	Tumor acquisition for biomarker research in lung cancer. Cancer Investigation, 2014, 32, 291-8	2.1	3
128	An integrated transcriptome and expressed variant analysis of sepsis survival and death. <i>Genome Medicine</i> , 2014 , 6, 111	14.4	48
127	Pilot study of pharmacist-assisted delivery of pharmacogenetic testing in a primary care setting. <i>Pharmacogenomics</i> , 2014 , 15, 1677-86	2.6	27
126	Development of a novel preclinical model of pneumococcal pneumonia in nonhuman primates. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2014 , 50, 995-1004	5.7	16
125	Bayesian modeling of temporal properties of infectious disease in a college student population. Journal of Applied Statistics, 2014 , 41, 1358-1382	1	2
124	Genomics-enabled drug repositioning and repurposing: insights from an IOM Roundtable activity. JAMA - Journal of the American Medical Association, 2014 , 311, 2063-4	27.4	23
123	Preoperative CYP2D6 metabolism-dependent Eblocker use and mortality after coronary artery bypass grafting surgery. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2014 , 147, 1368-1375.e3	1.5	12
122	Medical genomics: Gather and use genetic data in health care. <i>Nature</i> , 2014 , 508, 451-3	50.4	39

(2013-2013)

121	Longitudinal analysis of leukocyte differentials in peripheral blood of patients with acute respiratory viral infections. <i>Journal of Clinical Virology</i> , 2013 , 58, 689-95	14.5	44
120	Aspirin exposure reveals novel genes associated with platelet function and cardiovascular events. Journal of the American College of Cardiology, 2013 , 62, 1267-1276	15.1	43
119	Genomic medicine: a decade of successes, challenges, and opportunities. <i>Science Translational Medicine</i> , 2013 , 5, 189sr4	17.5	165
118	Patient and primary care provider experience using a family health history collection, risk stratification, and clinical decision support tool: a type 2 hybrid controlled implementation-effectiveness trial. <i>BMC Family Practice</i> , 2013 , 14, 111	2.6	46
117	Application of Human Genome Information to Clinical Practice 2013, 204-215		
116	Implementing genomic medicine in the clinic: the future is here. <i>Genetics in Medicine</i> , 2013 , 15, 258-67	8.1	385
115	Surface-enhanced Raman scattering molecular sentinel nanoprobes for viral infection diagnostics. <i>Analytica Chimica Acta</i> , 2013 , 786, 153-8	6.6	26
114	Clinical utility of genetic risk testing in primary care: the example of Type 2 diabetes. <i>Personalized Medicine</i> , 2013 , 10, 549-563	2.2	12
113	Comparative effectiveness research in cancer genomics and precision medicine: current landscape and future prospects. <i>Journal of the National Cancer Institute</i> , 2013 , 105, 929-36	9.7	28
112	A host-based RT-PCR gene expression signature to identify acute respiratory viral infection. <i>Science Translational Medicine</i> , 2013 , 5, 203ra126	17.5	102
111	Realizing the opportunities of genomics in health care. <i>JAMA - Journal of the American Medical Association</i> , 2013 , 309, 1463-4	27.4	19
110	An integrated clinico-metabolomic model improves prediction of death in sepsis. <i>Science Translational Medicine</i> , 2013 , 5, 195ra95	17.5	282
109	Transforming epidemiology for 21st century medicine and public health. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 508-16	4	83
108	The genomic medicine model: an integrated approach to implementation of family health history in primary care. <i>Personalized Medicine</i> , 2013 , 10, 295-306	2.2	17
107	Comparing influenza and RSV viral and disease dynamics in experimentally infected adults predicts clinical effectiveness of RSV antivirals. <i>Antiviral Therapy</i> , 2013 , 18, 785-91	1.6	47
106	Health coaching and genomics-potential avenues to elicit behavior change in those at risk for chronic disease: protocol for personalized medicine effectiveness study in air force primary care. <i>Global Advances in Health and Medicine</i> , 2013 , 2, 26-38	1.9	44
105	Gene expression-based classifiers identify Staphylococcus aureus infection in mice and humans. <i>PLoS ONE</i> , 2013 , 8, e48979	3.7	39
104	A host transcriptional signature for presymptomatic detection of infection in humans exposed to influenza H1N1 or H3N2. <i>PLoS ONE</i> , 2013 , 8, e52198	3.7	121

103	Development and validation of a primary care-based family health history and decision support program (MeTree). <i>North Carolina Medical Journal</i> , 2013 , 74, 287-96	0.6	49
102	Collection of family health history for assessment of chronic disease risk in primary care. <i>North Carolina Medical Journal</i> , 2013 , 74, 279-86	0.6	20
101	The host response to infection: advancing a novel diagnostic paradigm. <i>Critical Care</i> , 2012 , 16, 168	10.8	6
100	Examining the impact of genetic testing for type 2 diabetes on health behaviors: study protocol for a randomized controlled trial. <i>Trials</i> , 2012 , 13, 121	2.8	15
99	High-Dimensional Longitudinal Genomic Data: An analysis used for monitoring viral infections. <i>IEEE Signal Processing Magazine</i> , 2012 , 29, 108-123	9.4	9
98	Clinical application of cardiovascular pharmacogenetics. <i>Journal of the American College of Cardiology</i> , 2012 , 60, 9-20	15.1	53
97	Genetic testing: clinical and personal utility. AMA Journal of Ethics, 2012, 14, 604-9	1.4	3
96	Time-dependent changes in non-COX-1-dependent platelet function with daily aspirin therapy. Journal of Thrombosis and Thrombolysis, 2012 , 33, 246-57	5.1	16
95	Comparative effectiveness research, genomics-enabled personalized medicine, and rapid learning health care: a common bond. <i>Journal of Clinical Oncology</i> , 2012 , 30, 4233-42	2.2	39
94	Conference Scene: Is personalized medicine ready for prime time?. <i>Personalized Medicine</i> , 2012 , 9, 475-4	1782	1
93	The MURDOCK Study: a long-term initiative for disease reclassification through advanced biomarker discovery and integration with electronic health records. <i>American Journal of Translational Research (discontinued)</i> , 2012 , 4, 291-301	3	20
92	The Measurement to Understand Reclassification of Disease of Cabarrus/Kannapolis (MURDOCK) Study Community Registry and Biorepository. <i>American Journal of Translational Research (discontinued)</i> , 2012 , 4, 458-70	3	16
91	Application of Molecular Technologies to Clinical Medicine 2012 , 199-203		
90	Personalized medicine: progress and promise. <i>Annual Review of Genomics and Human Genetics</i> , 2011 , 12, 217-44	9.7	183
89	Whole blood gene expression analyses in patients with single versus recurrent venous thromboembolism. <i>Thrombosis Research</i> , 2011 , 128, 536-40	8.2	33
88	H3N2 influenza infection elicits more cross-reactive and less clonally expanded anti-hemagglutinin antibodies than influenza vaccination. <i>PLoS ONE</i> , 2011 , 6, e25797	3.7	139
87	Consideration of patient preferences and challenges in storage and access of pharmacogenetic test results. <i>Genetics in Medicine</i> , 2011 , 13, 887-90	8.1	16
86	Genomic risk profiling: attitudes and use in personal and clinical care of primary care physicians who offer risk profiling. <i>Journal of General Internal Medicine</i> , 2011 , 26, 834-40	4	56

(2010-2011)

85	Discovery of biomarker candidates for coronary artery disease from an APOE-knock out mouse model using iTRAQ-based multiplex quantitative proteomics. <i>Proteomics</i> , 2011 , 11, 2763-76	4.8	28
84	High predictive accuracy of an unbiased proteomic profile for sustained virologic response in chronic hepatitis C patients. <i>Hepatology</i> , 2011 , 53, 1809-18	11.2	36
83	Detection of viruses via statistical gene expression analysis. <i>IEEE Transactions on Biomedical Engineering</i> , 2011 , 58, 468-79	5	20
82	Predicting Viral Infection From High-Dimensional Biomarker Trajectories. <i>Journal of the American Statistical Association</i> , 2011 , 106, 1259-1279	2.8	21
81	A hub for bench-to-bedside pharmacogenomic-based research. <i>Pharmacogenomics</i> , 2011 , 12, 1095-8	2.6	6
80	Temporal dynamics of host molecular responses differentiate symptomatic and asymptomatic influenza a infection. <i>PLoS Genetics</i> , 2011 , 7, e1002234	6	141
79	. IEEE Transactions on Signal Processing, 2011 , 59, 4447-4458	4.8	2
78	Academic medical centers: ripe for rapid-learning personalized health care. <i>Science Translational Medicine</i> , 2011 , 3, 101cm27	17.5	22
77	Opening up to precompetitive collaboration. <i>Science Translational Medicine</i> , 2010 , 2, 52cm26	17.5	40
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75 74 73	Intratumor heterogeneity and precision of microarray-based predictors of breast cancer biology and clinical outcome. <i>Journal of Clinical Oncology</i> , 2010 , 28, 2198-206 Genome-wide association study of Lp-PLA(2) activity and mass in the Framingham Heart Study. <i>PLoS Genetics</i> , 2010 , 6, e1000928 Identifying patients at high risk of a cardiovascular event in the near future: current status and future directions: report of a national heart, lung, and blood institute working group. <i>Circulation</i> , 2010 , 121, 1447-54 Cancer pharmacogenomics and pharmacoepidemiology: setting a research agenda to accelerate	6	795763
75 74 73 72	Intratumor heterogeneity and precision of microarray-based predictors of breast cancer biology and clinical outcome. <i>Journal of Clinical Oncology</i> , 2010 , 28, 2198-206 Genome-wide association study of Lp-PLA(2) activity and mass in the Framingham Heart Study. <i>PLoS Genetics</i> , 2010 , 6, e1000928 Identifying patients at high risk of a cardiovascular event in the near future: current status and future directions: report of a national heart, lung, and blood institute working group. <i>Circulation</i> , 2010 , 121, 1447-54 Cancer pharmacogenomics and pharmacoepidemiology: setting a research agenda to accelerate translation. <i>Journal of the National Cancer Institute</i> , 2010 , 102, 1698-705	6	795763
75 74 73 72 71	Intratumor heterogeneity and precision of microarray-based predictors of breast cancer biology and clinical outcome. <i>Journal of Clinical Oncology</i> , 2010 , 28, 2198-206 Genome-wide association study of Lp-PLA(2) activity and mass in the Framingham Heart Study. <i>PLoS Genetics</i> , 2010 , 6, e1000928 Identifying patients at high risk of a cardiovascular event in the near future: current status and future directions: report of a national heart, lung, and blood institute working group. <i>Circulation</i> , 2010 , 121, 1447-54 Cancer pharmacogenomics and pharmacoepidemiology: setting a research agenda to accelerate translation. <i>Journal of the National Cancer Institute</i> , 2010 , 102, 1698-705 Translational Genomics 2010 , 163-174 Response: Improving Development of the Molecular Signature for Diagnosis of Acute Respiratory	6 16.7 9.7	79576336

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7	Transcriptional regulation of the cholesteryl ester transfer protein gene by the orphan nuclear hormone receptor apolipoprotein AI regulatory protein-1. <i>Journal of Biological Chemistry</i> , 1995 , 270, 29916-22	5.4	37
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