Garth D Ehrlich

List of Publications by Year
in descending order

133	6,673	71102	66911
papers	citations	41 h-index	78 g-index

139	139	139	8374
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Direct Detection of Bacterial Biofilms on the Middle-Ear Mucosa of Children With Chronic Otitis Media. JAMA - Journal of the American Medical Association, 2006, 296, 202.	7.4	754
2	Structure and dynamics of the pan-genome of Streptococcus pneumoniae and closely related species. Genome Biology, 2010, 11, R107.	8.8	321
3	Comparative Genomic Analyses of Seventeen Streptococcus pneumoniae Strains: Insights into the Pneumococcal Supragenome. Journal of Bacteriology, 2007, 189, 8186-8195.	2.2	249
4	Interaction between the microbiome and TP53 in human lung cancer. Genome Biology, 2018, 19, 123.	8.8	247
5	Biofilms and Chronic Infections. JAMA - Journal of the American Medical Association, 2008, 299, 2682.	7.4	232
6	Characterization and modeling of the Haemophilus influenzae core and supragenomes based on the complete genomic sequences of Rd and 12 clinical nontypeable strains. Genome Biology, 2007, 8, R103.	9.6	228
7	The microbiome of chronic rhinosinusitis: culture, molecular diagnostics and biofilm detection. BMC Infectious Diseases, 2013, 13, 210.	2.9	223
8	Characterization of biofilm matrix, degradation by DNase treatment and evidence of capsule downregulation in Streptococcus pneumoniae clinical isolates. BMC Microbiology, 2008, 8, 173.	3.3	211
9	Diagnosis of Periprosthetic Joint Infection. Journal of Arthroplasty, 2014, 29, 77-83.	3.1	193
10	Meta-omic Characterization of the Marine Invertebrate Microbial Consortium That Produces the Chemotherapeutic Natural Product ET-743. ACS Chemical Biology, 2011, 6, 1244-1256.	3.4	171
11	Comparative Genomic Analyses of 17 Clinical Isolates of Gardnerella vaginalis Provide Evidence of Multiple Genetically Isolated Clades Consistent with Subspeciation into Genovars. Journal of Bacteriology, 2012, 194, 3922-3937.	2.2	147
12	Generation of Genic Diversity among Streptococcus pneumoniae Strains via Horizontal Gene Transfer during a Chronic Polyclonal Pediatric Infection. PLoS Pathogens, 2010, 6, e1001108.	4.7	141
13	Biofilm-Based Implant Infections in Orthopaedics. Advances in Experimental Medicine and Biology, 2015, 830, 29-46.	1.6	134
14	Orthopaedic biofilm infections. Current Orthopaedic Practice, 2011, 22, 558-563.	0.2	133
15	Successful Identification of Pathogens by Polymerase Chain Reaction (PCR)-Based Electron Spray Ionization Time-of-Flight Mass Spectrometry (ESI-TOF-MS) in Culture-Negative Periprosthetic Joint Infection. Journal of Bone and Joint Surgery - Series A, 2012, 94, 2247-2254.	3.0	129
16	Bacterial Plurality as a General Mechanism Driving Persistence in Chronic Infections. Clinical Orthopaedics and Related Research, 2005, &NA;, 20-24.	1.5	120
17	Species-level bacterial community profiling of the healthy sinonasal microbiome using Pacific Biosciences sequencing of full-length 16S rRNA genes. Microbiome, 2018, 6, 190.	11.1	117
18	PCR-based detection of bacterial DNA after antimicrobial treatment is indicative of persistent, viable bacteria in the chinchilla model of otitis media. American Journal of Otolaryngology - Head and Neck Medicine and Surgery, 1996, 17, 106-111.	1.3	111

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19	Engineering Approaches for the Detection and Control of Orthopaedic Biofilm Infections. Clinical Orthopaedics and Related Research, 2005, &NA;, 59-66.	1.5	105
20	Chronic Surgical Site Infection Due to Suture-Associated Polymicrobial Biofilm. Surgical Infections, 2009, 10, 457-461.	1.4	101
21	Characterization of a mixed MRSA/MRSE biofilm in an explanted total ankle arthroplasty. FEMS Immunology and Medical Microbiology, 2011, 62, 66-74.	2.7	96
22	Salicylic acid-releasing polyurethane acrylate polymers as anti-biofilm urological catheter coatings. Acta Biomaterialia, 2012, 8, 1869-1880.	8.3	93
23	Assessment of the Lower Urinary Tract Microbiota during Symptom Flare in Women with Urologic Chronic Pelvic Pain Syndrome: A MAPP Network Study. Journal of Urology, 2016, 195, 356-362.	0.4	92
24	Deletion and acquisition of genomic content during early stage adaptation of Pseudomonas aeruginosa to a human host environment. Environmental Microbiology, 2012, 14, 2200-2211.	3.8	88
25	Can We Trust Intraoperative Culture Results in Nonunions?. Journal of Orthopaedic Trauma, 2014, 28, 384-390.	1.4	81
26	The distributed genome hypothesis as a rubric for understanding evolutionin situduring chronic bacterial biofilm infectious processes. FEMS Immunology and Medical Microbiology, 2010, 59, 269-279.	2.7	80
27	Extensive Genomic Plasticity in Pseudomonas aeruginosa Revealed by Identification and Distribution Studies of Novel Genes among Clinical Isolates. Infection and Immunity, 2006, 74, 5272-5283.	2.2	78
28	Search for Microorganisms in Men with Urologic Chronic Pelvic Pain Syndrome: A Culture-Independent Analysis in the MAPP Research Network. Journal of Urology, 2015, 194, 127-135.	0.4	75
29	Genome of alkaliphilic Bacillus pseudofirmus OF4 reveals adaptations that support the ability to grow in an external pH range from 7.5 to 11.4. Environmental Microbiology, 2011, 13, 3289-3309.	3.8	73
30	Characterization of Bacterial Communities in Venous Insufficiency Wounds by Use of Conventional Culture and Molecular Diagnostic Methods. Journal of Clinical Microbiology, 2011, 49, 3812-3819.	3.9	65
31	Comparative Evaluation of Culture and PCR for the Detection and Determination of Persistence of Bacterial Strains and DNAS in the Chinchilla Laniger Model of Otitis Media. Annals of Otology, Rhinology and Laryngology, 1998, 107, 508-513.	1.1	64
32	Diagnosis of Periprosthetic Joint Infection. Journal of Orthopaedic Research, 2014, 32, S98-107.	2.3	64
33	Phenotypic diversity and genotypic flexibility of Burkholderia cenocepacia during long-term chronic infection of cystic fibrosis lungs. Genome Research, 2017, 27, 650-662.	5.5	64
34	Age of Child, More than HPV Type, Is Associated with Clinical Course in Recurrent Respiratory Papillomatosis. PLoS ONE, 2008, 3, e2263.	2.5	63
35	Prevalence of Propionibacterium acnes in Intervertebral Discs of Patients Undergoing Lumbar Microdiscectomy: A Prospective Cross-Sectional Study. PLoS ONE, 2016, 11, e0161676.	2.5	63
36	What makes pathogens pathogenic. Genome Biology, 2008, 9, 225.	9.6	60

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37	Identification, Distribution, and Expression of Novel Genes in 10 Clinical Isolates of Nontypeable Haemophilus influenzae. Infection and Immunity, 2005, 73, 3479-3491.	2.2	59
38	Pan-genome analysis provides much higher strain typing resolution than multi-locus sequence typing. Microbiology (United Kingdom), 2010, 156, 1060-1068.	1.8	50
39	Comparative supragenomic analyses among the pathogens Staphylococcus aureus, Streptococcus pneumoniae, and Haemophilus influenzae Using a modification of the finite supragenome model. BMC Genomics, 2011, 12, 187.	2.8	50
40	Comparative analysis and supragenome modeling of twelve Moraxella catarrhalis clinical isolates. BMC Genomics, 2011, 12, 70.	2.8	50
41	Characterization, Distribution, and Expression of Novel Genes among Eight Clinical Isolates of Streptococcuspneumoniae. Infection and Immunity, 2006, 74, 321-330.	2.2	45
42	Molecular characterization of Î²-lactamase genes in clinical isolates of carbapenem-resistant Acinetobacter baumannii. Annals of Clinical Microbiology and Antimicrobials, 2017, 16, 75.	3.8	44
43	Age at diagnosis, but not HPV type, is strongly associated with clinical course in recurrent respiratory papillomatosis. PLoS ONE, 2019, 14, e0216697.	2.5	43
44	Comparing culture and molecular methods for the identification of microorganisms involved in necrotizing soft tissue infections. BMC Infectious Diseases, 2016, 16, 652.	2.9	41
45	Antagonistic Pleiotropy in the Bifunctional Surface Protein FadL (OmpP1) during Adaptation of Haemophilus influenzae to Chronic Lung Infection Associated with Chronic Obstructive Pulmonary Disease. MBio, 2018, 9, .	4.1	39
46	A Culture-Independent Analysis of the Microbiota of Female Interstitial Cystitis/Bladder Pain Syndrome Participants in the MAPP Research Network. Journal of Clinical Medicine, 2019, 8, 415.	2.4	37
47	Strain-Specific Virulence Phenotypes of Streptococcus pneumoniae Assessed Using the Chinchilla laniger Model of Otitis Media. PLoS ONE, 2008, 3, e1969.	2.5	35
48	Direct Demonstration of <i>Staphylococcus</i> Biofilm in an External Ventricular Drain in a Patient with a History of Recurrent Ventriculoperitoneal Shunt Failure. Pediatric Neurosurgery, 2010, 46, 127-132.	0.7	35
49	Genetic Stabilization of the Drug-Resistant PMEN1 Pneumococcus Lineage by Its Distinctive DpnIII Restriction-Modification System. MBio, 2015, 6, e00173.	4.1	31
50	The Tsk2/+ Mouse Fibrotic Phenotype Is Due to a Gain-of-Function Mutation in the PIIINP Segment of the Col3a1 Gene. Journal of Investigative Dermatology, 2015, 135, 718-727.	0.7	30
51	HIV-1 Genetic Variation Resulting in the Development of New Quasispecies Continues to Be Encountered in the Peripheral Blood of Well-Suppressed Patients. PLoS ONE, 2016, 11, e0155382.	2.5	29
52	Bacterial Biofilm Growth on 3D-Printed Materials. Frontiers in Microbiology, 2021, 12, 646303.	3.5	29
53	Broad-Spectrum and Personalized Guide RNAs for CRISPR/Cas9 HIV-1 Therapeutics. AIDS Research and Human Retroviruses, 2018, 34, 950-960.	1.1	26
54	Nontypeable Haemophilus influenzae Genetic Islands Associated with Chronic Pulmonary Infection. PLoS ONE, 2012, 7, e44730.	2.5	25

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55	Virulence phenotypes of low-passage clinical isolates of Nontypeable Haemophilus influenzae assessed using the chinchilla laniger model of otitis media. BMC Microbiology, 2007, 7, 56.	3.3	24
56	Population-level virulence factors amongst pathogenic bacteria: relation to infection outcome. Future Microbiology, 2008, 3, 31-42.	2.0	24
57	Fibroblasts from phenotypically normal palmar fascia exhibit molecular profiles highly similar to fibroblasts from active disease in Dupuytren's Contracture. BMC Medical Genomics, 2012, 5, 15.	1.5	24
58	Differences in Genotype and Virulence among Four Multidrug-Resistant Streptococcus pneumoniae Isolates Belonging to the PMEN1 Clone. PLoS ONE, 2011, 6, e28850.	2.5	23
59	Novel gRNA design pipeline to develop broad-spectrum CRISPR/Cas9 gRNAs for safe targeting of the HIV-1 quasispecies in patients. Scientific Reports, 2019, 9, 17088.	3.3	23
60	Century of Jackson-Weiss syndrome: Further definition of clinical and radiographic findings in ?lost? descendants of the original kindred. American Journal of Medical Genetics Part A, 2001, 100, 315-324.	2.4	22
61	What role do periodontal pathogens play in osteoarthritis and periprosthetic joint infections of the knee?. Journal of Applied Biomaterials and Functional Materials, 2014, 12, 13-20.	1.6	22
62	In Vivo Capsular Switch in Streptococcus pneumoniae â€“ Analysis by Whole Genome Sequencing. PLoS ONE, 2012, 7, e47983.	2.5	22
63	Multicenter Initiative Seeking Critical Genes in Respiratory Papillomatosis. Laryngoscope, 2004, 114, 349-357.	2.0	21
64	From Koch's Postulates to Biofilm Theory. The Lesson of Bill Costerton. International Journal of Artificial Organs, 2012, 35, 695-699.	1.4	21
65	The Chinchilla Research Resource Database: resource for an otolaryngology disease model. Database: the Journal of Biological Databases and Curation, 2016, 2016, baw073.	3.0	20
66	Comparative Genomic Analyses of theMoraxella catarrhalisSerosensitive and Seroresistant Lineages Demonstrate Their Independent Evolution. Genome Biology and Evolution, 2016, 8, 955-974.	2.5	20
67	Identification of a patient withStreptococcus pneumoniaebacteremia and meningitis by the polymerase chain reaction (PCR). Molecular and Cellular Probes, 1995, 9, 157-160.	2.1	19
68	Detection of methicillin-resistant and methicillin-susceptible Staphylococcus aureus colonization of healthy military personnel by traditional culture, PCR, and mass spectrometry. Scandinavian Journal of Infectious Diseases, 2013, 45, 752-759.	1.5	19
69	Demonstration of Bacillus cereus in Orthopaedic-Implant-Related Infection with Use of a Multi-Primer Polymerase Chain Reaction-Mass Spectrometric Assay. Journal of Bone and Joint Surgery - Series A, 2011, 93, e85.	3.0	18
70	Specific amino acids in HIV-1 Vpr are significantly associated with differences in patient neurocognitive status. Journal of NeuroVirology, 2017, 23, 113-124.	2.1	18
71	Genome rearrangements induce biofilm formation in Escherichia coli C â€“ an old model organism with a new application in biofilm research. BMC Genomics, 2019, 20, 767.	2.8	18
72	Mechanical effects, antimicrobial efficacy and cytotoxicity of usnic acid as a biofilm prophylaxis in PMMA. Journal of Materials Science: Materials in Medicine, 2011, 22, 2773-2780.	3.6	17

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73	Comparison of PCR/Electron spray Ionization-Time-of-Flight-Mass Spectrometry versus Traditional Clinical Microbiology for active surveillance of organisms contaminating high-use surfaces in a burn intensive care unit, an orthopedic ward and healthcare workers. BMC Infectious Diseases, 2012, 12, 252.	2.9	16
74	The use of PCR/Electrospray Ionization-Time-of-Flight-Mass Spectrometry (PCR/ESI-TOF-MS) to detect bacterial and fungal colonization in healthy military service members. BMC Infectious Diseases, 2016, 16, 338.	2.9	16
75	Polymerase Chain Reactionâ€"Electrosprayâ€"Time-of-Flight Mass Spectrometry Versus Culture for Bacterial Detection in Septic Arthritis and Osteoarthritis. Genetic Testing and Molecular Biomarkers, 2016, 20, 721-731.	0.7	16
76	Circular RNAs as Diagnostic Biomarkers for Osteoarthritis. Genetic Testing and Molecular Biomarkers, 2019, 23, 701-702.	0.7	16
77	The Time Is Now for Gene- and Genome-Based Bacterial Diagnostics. JAMA Internal Medicine, 2013, 173, 1405.	5.1	15
78	Identification and Characterization of msf, a Novel Virulence Factor in Haemophilus influenzae. PLoS ONE, 2016, 11, e0149891.	2.5	15
79	Design and validation of a supragenome array for determination of the genomic content of Haemophilus influenzae isolates. BMC Genomics, 2013, 14, 484.	2.8	14
80	Complete Genome Sequence of Haemophilus influenzae Strain 375 from the Middle Ear of a Pediatric Patient with Otitis Media. Genome Announcements, 2014, 2, .	0.8	14
81	Urinary fungi associated with urinary symptom severity among women with interstitial cystitis/bladder pain syndrome (IC/BPS). World Journal of Urology, 2020, 38, 433-446.	2.2	14
82	PCR Is Changing Clinical Diagnostics. Microbe Magazine, 2013, 8, 15-20.	0.4	14
83	Virulence Potential and Genome-Wide Characterization of Drug Resistant Streptococcus pneumoniae Clones Selected In Vivo by the 7-Valent Pneumococcal Conjugate Vaccine. PLoS ONE, 2013, 8, e74867.	2.5	13
84	Refined localization of a gene for pediatric gastroesophageal reflux makes HTR2A an unlikely candidate gene. Human Genetics, 2000, 107, 519-525.	3.8	12
85	There is a specific response to pH by isolates of Haemophilus influenzae and this has a direct influence on biofilm formation. BMC Microbiology, 2014, 14, 47.	3.3	11
86	Deletion of genes involved in the ketogluconate metabolism, Entner-Doudoroff pathway, and glucose dehydrogenase increase local and invasive virulence phenotypes in Streptococcus pneumoniae. PLoS ONE, 2019, 14, e0209688.	2.5	10
87	Codon usage comparison of novel genes in clinical isolates of Haemophilus influenzae. Nucleic Acids Research, 2005, 33, 3644-3658.	14.5	9
88	The Development of a Pipeline for the Identification and Validation of Small-Molecule RelA Inhibitors for Use as Anti-Biofilm Drugs. Microorganisms, 2020, 8, 1310.	3.6	9
89	Whole genome sequencing of Streptomyces actuosus ISP-5337, Streptomyces sioyaensis B-5408, and Actinospica acidiphila B-2296 reveals secondary metabolomes with antibiotic potential. Biotechnology Reports (Amsterdam, Netherlands), 2021, 29, e00596.	4.4	9
90	Microarray Analysis of Transcriptomic Response of <i>Escherichia coli</i> to Nonthermal Plasma-Treated PBS Solution. Advances in Bioscience and Biotechnology (Print), 2015, 06, 49-62.	0.7	9

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91	Death and Transfiguration in Static Staphylococcus epidermidis Cultures. PLoS ONE, 2014, 9, e100002.	2.5	8
92	Phase Variation in HMW1A Controls a Phenotypic Switch in Haemophilus influenzae Associated with Pathoadaptation during Persistent Infection. MBio, 2021, 12, e0078921.	4.1	8
93	Construction and characterization of a highly redundant Pseudomonas aeruginosa genomic library prepared from 12 clinical isolates: Application to studies of gene distribution among populations. International Journal of Pediatric Otorhinolaryngology, 2006, 70, 1891-1900.	1.0	6
94	A micropatterned substrate for on-surface enzymatic labelling of linearized long DNA molecules. Scientific Reports, 2019, 9, 15059.	3.3	6
95	The bacterial microbiota of Hunner lesion interstitial cystitis/bladder pain syndrome. BJU International, 2022, 129, 104-112.	2.5	6
96	Panel 3: Genomics, precision medicine and targeted therapies. International Journal of Pediatric Otorhinolaryngology, 2020, 130, 109835.	1.0	5
97	Species-Level Profiling of Ixodes pacificus Bacterial Microbiomes Reveals High Variability Across Short Spatial Scales at Different Taxonomic Resolutions. Genetic Testing and Molecular Biomarkers, 2021, 25, 551-562.	0.7	5
98	Beyond the pan-genome: current perspectives on the functional and practical outcomes of the distributed genome hypothesis. Biochemical Society Transactions, 2020, 48, 2437-2455.	3.4	5
99	Preliminary study: Treatment with intramuscular interferon beta-1a results in increased levels of IL-12RÎ²2+ and decreased levels of IL23R+ CD4+ T - Lymphocytes in multiple sclerosis. BMC Neurology, 2011, 11, 155.	1.8	4
100	Genome-wide analysis of DNA uptake across the outer membrane of naturally competent Haemophilus influenzae. IScience, 2021, 24, 102007.	4.1	4
101	Streptococcus pneumoniae Supragenome Hybridization Arrays for Profiling of Genetic Content and Gene Expression. Current Protocols in Microbiology, 2015, 36, 9D.4.1-9D.4.20.	6.5	4
102	Development and Validation of an Haemophilus influenzae Supragenome Hybridization (SGH) Array for Transcriptomic Analyses. PLoS ONE, 2014, 9, e105493.	2.5	4
103	The healthy urinary microbiome in asymptomatic participants in the MAPP Network Study: Relation to gender, age, and menopausal status. Canadian Urological Association Journal, 2022, 16, .	0.6	4
104	Bill Costerton: leader as servant. FEMS Immunology and Medical Microbiology, 2012, 66, 269-272.	2.7	3
105	Complete Genome Sequence of Aggregatibacter actinomycetemcomitans Strain IDH781. Genome Announcements, 2016, 4, .	0.8	3
106	Transition of Serotype 35B Pneumococci From Commensal to Prevalent Virulent Strain in Children. Frontiers in Cellular and Infection Microbiology, 2021, 11, 744742.	3.9	3
107	Bacterial Identification and Visualization of Bacterial Biofilms Adjacent to Fracture Sites After Internal Fixation. Genetic Testing and Molecular Biomarkers, 2022, 26, 70-80.	0.7	3
108	1147 APPLICATION OF STATE-OF-THE-ART METHODS TO SEARCH FOR MICROBIAL CONTRIBUTIONS TO THE ETIOLOGY OF UROLOGICAL CHRONIC PELVIC PAIN SYNDROME (UCPPS). Journal of Urology, 2013, 189, .	0.4	2

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109	Editorial: Otitis Media Genomics and the Middle Ear Microbiome. Frontiers in Genetics, 2021, 12, 763688.	2.3	2
110	Expanding the Scope of our Journal to Include Molecular Diagnostics for Infectious Diseases. Genetic Testing and Molecular Biomarkers, 2015, 19, 225-225.	0.7	1
111	Next-Generation Molecular Diagnostics Provide Evidence Suggestive of a Role for Nontraditional Bacterial Pathogens in Osteoarthritis of the Knee. Genetic Testing and Molecular Biomarkers, 2016, 20, 719-720.	0.7	1
112	Novel Genetic Markers for Common Degenerative Orthopedic Diseases. Genetic Testing and Molecular Biomarkers, 2017, 21, 577-577.	0.7	1
113	Development of a Trio of Potential Biomarkers for Cancer Prognosis. Genetic Testing and Molecular Biomarkers, 2018, 22, 1-2.	0.7	1
114	High-Fidelity Point-of-Care Diagnostic Test for Periprosthetic Joint Infection. Journal of Bone and Joint Surgery - Series A, 2018, 100, e7.	3.0	1
115	Correction: Structure and dynamics of the pan-genome of Streptococcus pneumoniae and closely related species. Genome Biology, 2011, 12, 140.	9.6	0
116	MP16-08 SEARCH FOR MICROORGANISMS IN MEN WITH UROLOGIC CHRONIC PELVIC PAIN SYNDROME: A CULTURE-INDEPENDENT ANALYSIS OF CASES AND CONTROLS ENROLLED IN THE TRANS-MAPP EPIDEMIOLOGY/PHENOTYPING (EP) STUDY. Journal of Urology, 2014, 191, .	0.4	0
117	MicroRNAs Come of Age in Diagnostics. Genetic Testing and Molecular Biomarkers, 2015, 19, 647-647.	0.7	0
118	Diagnostics Are the Vanguard of Medicine. Genetic Testing and Molecular Biomarkers, 2015, 19, 173-173.	0.7	0
119	Developing the Scientific Infrastructure to Produce Ethnogenetically-Specific Personalized Medicine. Genetic Testing and Molecular Biomarkers, 2015, 19, 465-466.	0.7	0
120	A Potential Role for Aromatase Levels in Coronary Heart Disease. Genetic Testing and Molecular Biomarkers, 2016, 20, 1-1.	0.7	0
121	Debunking the Myth of the Genetic Superman. Genetic Testing and Molecular Biomarkers, 2016, 20, 273-273.	0.7	0
122	The Paradox of Dickkopf-1: Tumor Suppressor and Tumor Enhancer. Genetic Testing and Molecular Biomarkers, 2016, 20, 163-164.	0.7	0
123	MP82-18 SHARED ALTERATIONS IN URINARY BACTERIAL COMMUNITIES IN PATIENTS WITH INTERSTITIAL CYSTITIS AND OVERACTIVE BLADDER. Journal of Urology, 2017, 197, .	0.4	0
124	Biobanking: Where Science and Society Meet. Genetic Testing and Molecular Biomarkers, 2017, 21, 127-127.	0.7	0
125	Changing of the Guard at GTMB. Genetic Testing and Molecular Biomarkers, 2017, 21, 455-455.	0.7	0
126	The Future Is Today. Genetic Testing and Molecular Biomarkers, 2018, 22, 279-280.	0.7	0

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127	The Rise of Noninvasive Diagnostic Technologies. Genetic Testing and Molecular Biomarkers, 2019, 23, 229-229.	0.7	0
128	Raising the Bar at GTMB. Genetic Testing and Molecular Biomarkers, 2019, 23, 151-152.	0.7	0
129	Genetic Testing and Molecular Biomarkers's Policy on Database-Derived Articles. Genetic Testing and Molecular Biomarkers, 2020, 24, 457-457.	0.7	0
130	Metabolic Markers of Chronic Disease States. Genetic Testing and Molecular Biomarkers, 2020, 24, 533-534.	0.7	0
131	LncRNAs H19 and MEG3 as Universal Indicators of Metabolic Derangements?. Genetic Testing and Molecular Biomarkers, 2020, 24, 319-320.	0.7	0
132	Abstract 4925: Microbiome-TP53 gene interaction in human lung cancer. , 2017, , .		0
133	The Human Genetics of Infection. Genetic Testing and Molecular Biomarkers, 2022, 26, 251-252.	0.7	0