

Josep Quer

List of Publications by Year in descending order

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Version: 2024-02-01

137
papers

4,377
citations

126907

33
h-index

144013

57
g-index

153
all docs

153
docs citations

153
times ranked

4514
citing authors

#	ARTICLE	IF	CITATIONS
1	A year living with SARS-CoV-2: an epidemiological overview of viral lineage circulation by whole-genome sequencing in Barcelona city (Catalonia, Spain). <i>Emerging Microbes and Infections</i> , 2022, 11, 172-181.	6.5	17
2	Effect of Hepatitis E Virus RNA Universal Blood Donor Screening, Catalonia, Spain, 2017-2020. <i>Emerging Infectious Diseases</i> , 2022, 28, 157-165.	4.3	9
3	Inspecting the Ribozyme Region of Hepatitis Delta Virus Genotype 1: Conservation and Variability. <i>Viruses</i> , 2022, 14, 215.	3.3	4
4	Dynamics of SARS-CoV-2 Alpha (B.1.1.7) variant spread: The wastewater surveillance approach. <i>Environmental Research</i> , 2022, 208, 112720.	7.5	16
5	On the Reliability of the Notion of Native Signer and Its Risks. <i>Frontiers in Psychology</i> , 2022, 13, 716554.	2.1	5
6	Next-Generation Sequencing for Confronting Virus Pandemics. <i>Viruses</i> , 2022, 14, 600.	3.3	24
7	Characteristics of hepatitis C virus resistance in an international cohort after a decade of direct-acting antivirals. <i>JHEP Reports</i> , 2022, 4, 100462.	4.9	10
8	Hepatitis B Virus Variants with Multiple Insertions and/or Deletions in the X Open Reading Frame 3' End: Common Members of Viral Quasispecies in Chronic Hepatitis B Patients. <i>Biomedicines</i> , 2022, 10, 1194.	3.2	1
9	Characteristics of 24 SARS-CoV-2-Sequenced Reinfection Cases in a Tertiary Hospital in Spain. <i>Frontiers in Microbiology</i> , 2022, 13, .	3.5	1
10	Host-dependent editing of SARS-CoV-2 in COVID-19 patients. <i>Emerging Microbes and Infections</i> , 2021, 10, 1777-1789.	6.5	13
11	Sophisticated viral quasispecies with a genotype-related pattern of mutations in the hepatitis B X gene of HBeAg-ve chronically infected patients. <i>Scientific Reports</i> , 2021, 11, 4215.	3.3	3
12	Utility of the Cobas® Plasma Separation Card as a Sample Collection Device for Serological and Virological Diagnosis of Hepatitis C Virus Infection. <i>Diagnostics</i> , 2021, 11, 473.	2.6	7
13	Resistance-associated substitutions after sofosbuvir/velpatasvir/voxilaprevir triple therapy failure. <i>Journal of Viral Hepatitis</i> , 2021, 28, 1319-1324.	2.0	7
14	Population Disequilibrium as Promoter of Adaptive Explorations in Hepatitis C Virus. <i>Viruses</i> , 2021, 13, 616.	3.3	7
15	Next-generation sequencing for the diagnosis of hepatitis B: current status and future prospects. <i>Expert Review of Molecular Diagnostics</i> , 2021, 21, 381-396.	3.1	6
16	Viral populations of SARS-CoV-2 in upper respiratory tract, placenta, amniotic fluid and umbilical cord blood support viral replication in placenta. <i>Clinical Microbiology and Infection</i> , 2021, 27, 1542-1544.	6.0	2
17	Partial restoration of immune response in Hepatitis C patients after viral clearance by direct-acting antiviral therapy. <i>PLoS ONE</i> , 2021, 16, e0254243.	2.5	6
18	Monitoring Emergence of the SARS-CoV-2 B.1.1.7 Variant through the Spanish National SARS-CoV-2 Wastewater Surveillance System (VATar COVID-19). <i>Environmental Science & Technology</i> , 2021, 55, 11756-11766.	10.0	39

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19	Exploring the diversity of coronavirus in sewage during COVID-19 pandemic: Don't miss the forest for the trees. <i>Science of the Total Environment</i> , 2021, 800, 149562.	8.0	14
20	Study of Quasispecies Complexity and Liver Damage Progression after Liver Transplantation in Hepatitis C Virus Infected Patients. <i>Genes</i> , 2021, 12, 1731.	2.4	0
21	Cross-sectional evaluation of circulating hepatitis B virus RNA and DNA: Different quasispecies?. <i>World Journal of Gastroenterology</i> , 2021, 27, 7144-7158.	3.3	1
22	Microorganisms as Shapers of Human Civilization, from Pandemics to Even Our Genomes: Villains or Friends? A Historical Approach. <i>Microorganisms</i> , 2021, 9, 2518.	3.6	6
23	Deep-sequencing reveals broad subtype-specific HCV resistance mutations associated with treatment failure. <i>Antiviral Research</i> , 2020, 174, 104694.	4.1	39
24	Amino Acid Substitutions Associated with Treatment Failure for Hepatitis C Virus Infection. <i>Journal of Clinical Microbiology</i> , 2020, 58, .	3.9	15
25	Naturally occurring SARS-CoV-2 gene deletions close to the spike S1/S2 cleavage site in the viral quasispecies of COVID19 patients. <i>Emerging Microbes and Infections</i> , 2020, 9, 1900-1911.	6.5	57
26	Dissimilar Conservation Pattern in Hepatitis C Virus Mutant Spectra, Consensus Sequences, and Data Banks. <i>Journal of Clinical Medicine</i> , 2020, 9, 3450.	2.4	12
27	Unexpected long-lasting anti-HEV IgM positivity: Is HEV antigen a better serological marker for hepatitis E infection diagnosis?. <i>Journal of Viral Hepatitis</i> , 2020, 27, 747-753.	2.0	26
28	Broad and Dynamic Diversification of Infectious Hepatitis C Virus in a Cell Culture Environment. <i>Journal of Virology</i> , 2020, 94, .	3.4	20
29	A new implication of quasispecies dynamics: Broad virus diversification in absence of external perturbations. <i>Infection, Genetics and Evolution</i> , 2020, 82, 104278.	2.3	20
30	Conservation and variability of hepatitis B core at different chronic hepatitis stages. <i>World Journal of Gastroenterology</i> , 2020, 26, 2584-2598.	3.3	8
31	The Critical Role of Codon Composition on the Translation Efficiency Robustness of the Hepatitis A Virus Capsid. <i>Genome Biology and Evolution</i> , 2019, 11, 2439-2456.	2.5	12
32	<p><p>Deep-sequencing study of HCV G4a resistance-associated substitutions in Egyptian patients failing DAA treatment</p></p>. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 2799-2807.	2.7	5
33	Delimiting reported discourse: Cross-modal criteria. <i>Linguistic Typology</i> , 2019, 23, 221-228.	1.2	2
34	<p><p>Whole-genome characterization and resistance-associated substitutions in a new HCV genotype 1 subtype</p></p>. <i>Infection and Drug Resistance</i> , 2019, Volume 12, 947-955.	2.7	5
35	Handling Sign Language Data: The Impact of Modality. <i>Frontiers in Psychology</i> , 2019, 10, 483.	2.1	15
36	Significant Improvement in Diagnosis of Hepatitis C Virus Infection by a One-Step Strategy in a Central Laboratory: an Optimal Tool for Hepatitis C Elimination?. <i>Journal of Clinical Microbiology</i> , 2019, 58, .	3.9	21

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37	Viral Load Measurements in Individuals with Hepatitis C Virus Infection: on the European Association for the Study of the Liver Recommendations on Treatment of Hepatitis C 2018. <i>Journal of Clinical Microbiology</i> , 2019, 58, .	3.9	6
38	The increasing impact of lethal mutagenesis of viruses. <i>Future Medicinal Chemistry</i> , 2019, 11, 1645-1657.	2.3	30
39	Evidence for positive selection of hepatitis A virus antigenic variants in vaccinated men-having-sex-with men patients: Implications for immunization policies. <i>EBioMedicine</i> , 2019, 39, 348-357.	6.1	22
40	Synergistic Lethal Mutagenesis of Hepatitis C Virus. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	3.2	10
41	Characterization of hepatitis B virus X gene quasispecies complexity in mono-infection and hepatitis delta virus superinfection. <i>World Journal of Gastroenterology</i> , 2019, 25, 1566-1579.	3.3	7
42	Hepatitis C virus intrinsic molecular determinants may contribute to the development of cholestatic hepatitis after liver transplantation. <i>Journal of General Virology</i> , 2019, 100, 63-68.	2.9	1
43	The noun-verb distinction in Catalan Sign Language. <i>Sign Language and Linguistics (Online)</i> , 2019, 22, 1-43.	0.5	0
44	Hepatitis C virus deep sequencing for sub-genotype identification in mixed infections: A real-life experience. <i>International Journal of Infectious Diseases</i> , 2018, 67, 114-117.	3.3	23
45	Detection of hyper-conserved regions in hepatitis B virus X gene potentially useful for gene therapy. <i>World Journal of Gastroenterology</i> , 2018, 24, 2095-2107.	3.3	17
46	On categorizing types of role shift in Sign languages. <i>Theoretical Linguistics</i> , 2018, 44, 277-282.	0.2	6
47	Pipeline for specific subtype amplification and drug resistance detection in hepatitis C virus. <i>BMC Infectious Diseases</i> , 2018, 18, 446.	2.9	29
48	Baseline hepatitis C virus resistance-associated substitutions present at frequencies lower than 15% may be clinically significant. <i>Infection and Drug Resistance</i> , 2018, Volume 11, 2207-2210.	2.7	26
49	Thrombotic thrombocytopenic purpura relapse induced by acute hepatitis E transmitted by cryosupernatant plasma and successfully controlled with ribavirin. <i>Transfusion</i> , 2018, 58, 2501-2505.	1.6	14
50	Rare haplotype load as marker for lethal mutagenesis. <i>PLoS ONE</i> , 2018, 13, e0204877.	2.5	8
51	Hepatitis C virus early kinetics and resistance-associated substitution dynamics during antiviral therapy with direct-acting antivirals. <i>Journal of Viral Hepatitis</i> , 2018, 25, 1515-1525.	2.0	10
52	Resistance of high fitness hepatitis C virus to lethal mutagenesis. <i>Virology</i> , 2018, 523, 100-109.	2.4	30
53	Characterization of intra- and inter-host norovirus P2 genetic variability in linked individuals by amplicon sequencing. <i>PLoS ONE</i> , 2018, 13, e0201850.	2.5	10
54	HIV-1 Protease Evolvability Is Affected by Synonymous Nucleotide Recoding. <i>Journal of Virology</i> , 2018, 92, .	3.4	9

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55	Agent-backgrounding in Catalan Sign Language (LSC). <i>Sign Language and Linguistics (Online)</i> , 2018, 21, 334-349.	0.5	1
56	New hepatitis C virus genotype 1 subtype naturally harbouring resistance-associated mutations to NS5A inhibitors. <i>Journal of General Virology</i> , 2018, 99, 97-102.	2.9	6
57	Analysis of hepatitis B virus preS1 variability and prevalence of the rs2296651 polymorphism in a Spanish population. <i>World Journal of Gastroenterology</i> , 2018, 24, 680-692.	3.3	10
58	Internal Disequilibria and Phenotypic Diversification during Replication of Hepatitis C Virus in a Noncoevolving Cellular Environment. <i>Journal of Virology</i> , 2017, 91, .	3.4	42
59	Phylogenetic analysis of an epidemic outbreak of acute hepatitis C in HIV-infected patients by ultra-deep pyrosequencing. <i>Journal of Clinical Virology</i> , 2017, 92, 42-47.	3.1	17
60	Deep sequencing in the management of hepatitis virus infections. <i>Virus Research</i> , 2017, 239, 115-125.	2.2	23
61	Assessment of a Novel Automatic Real-Time PCR Assay on the Cobas 4800 Analyzer as a Screening Platform for Hepatitis C Virus Genotyping in Clinical Practice: Comparison with Massive Sequencing. <i>Journal of Clinical Microbiology</i> , 2017, 55, 504-509.	3.9	15
62	Epidemiological trends of HIV-1 infection in blood donors from Catalonia, Spain (2005-2014). <i>Transfusion</i> , 2017, 57, 2164-2173.	1.6	11
63	Red blood cell transfusion-transmitted acute hepatitis E in an immunocompetent subject in Europe: a case report. <i>Transfusion</i> , 2017, 57, 244-247.	1.6	41
64	Evolution of acute hepatitis C virus infection in a large European city: Trends and new patterns. <i>PLoS ONE</i> , 2017, 12, e0187893.	2.5	8
65	Lethal Mutagenesis of Hepatitis C Virus Induced by Favipiravir. <i>PLoS ONE</i> , 2016, 11, e0164691.	2.5	63
66	Identification of hepatitis C virus genotype 3 by a commercial assay challenged by natural polymorphisms detected in Spain from patients with diverse origins. <i>Journal of Clinical Virology</i> , 2016, 78, 14-19.	3.1	2
67	Subjunctive mood in Griko: A micro-comparative approach. <i>Lingua</i> , 2016, 174, 65-85.	1.0	5
68	Barrier-Independent, Fitness-Associated Differences in Sofosbuvir Efficacy against Hepatitis C Virus. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 3786-3793.	3.2	42
69	Viral quasispecies complexity measures. <i>Virology</i> , 2016, 493, 227-237.	2.4	109
70	Improving virus production through quasispecies genomic selection and molecular breeding. <i>Scientific Reports</i> , 2016, 6, 35962.	3.3	12
71	One or two derivations in (bimodal) bilinguals. <i>Linguistic Approaches To Bilingualism</i> , 2016, 6, 817-821.	0.9	0
72	Mood. , 2016, , 954-966.		3

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73	Evidence of an Exponential Decay Pattern of the Hepatitis Delta Virus Evolution Rate and Fluctuations in Quasispecies Complexity in Long-Term Studies of Chronic Delta Infection. <i>PLoS ONE</i> , 2016, 11, e0158557.	2.5	19
74	Reporting with and without role shift: sign language strategies of complementation. , 2016, , 204-230.		3
75	New real-time-PCR method to identify single point mutations in hepatitis C virus. <i>World Journal of Gastroenterology</i> , 2016, 22, 9604.	3.3	5
76	Ambiguities in Sign Languages. <i>Linguistic Review</i> , 2015, 32, .	0.4	3
77	Resistance of Hepatitis C Virus to Inhibitors: Complexity and Clinical Implications. <i>Viruses</i> , 2015, 7, 5746-5766.	3.3	44
78	Complex Genotype Mixtures Analyzed by Deep Sequencing in Two Different Regions of Hepatitis B Virus. <i>PLoS ONE</i> , 2015, 10, e0144816.	2.5	9
79	Ambiguities in sign languages. <i>Linguistic Review</i> , 2015, 32, .	0.4	6
80	Phylogenetic Demonstration of Hepatitis E Infection Transmitted by Pork Meat Ingestion. <i>Journal of Clinical Gastroenterology</i> , 2015, 49, 165-168.	2.2	80
81	High-Resolution Hepatitis C Virus Subtyping Using NS5B Deep Sequencing and Phylogeny, an Alternative to Current Methods. <i>Journal of Clinical Microbiology</i> , 2015, 53, 219-226.	3.9	74
82	Quasispecies dynamics in hepatitis C liver transplant recipients receiving grafts from hepatitis C virus infected donors. <i>Journal of General Virology</i> , 2015, 96, 3493-3498.	2.9	9
83	Clinical Application of Estimating Hepatitis B Virus Quasispecies Complexity by Massive Sequencing: Correlation between Natural Evolution and On-Treatment Evolution. <i>PLoS ONE</i> , 2014, 9, e112306.	2.5	26
84	Identification of host and viral factors involved in a dissimilar resolution of a hepatitis C virus infection. <i>Liver International</i> , 2014, 34, 896-906.	3.9	10
85	Inference with viral quasispecies diversity indices: clonal and NGS approaches. <i>Bioinformatics</i> , 2014, 30, 1104-1111.	4.1	56
86	Increased Replicative Fitness Can Lead to Decreased Drug Sensitivity of Hepatitis C Virus. <i>Journal of Virology</i> , 2014, 88, 12098-12111.	3.4	74
87	Cirrhosis, Liver Transplantation and HIV Infection Are Risk Factors Associated with Hepatitis E Virus Infection. <i>PLoS ONE</i> , 2014, 9, e103028.	2.5	46
88	Determining argument structure in sign languages. <i>Language Faculty and Beyond</i> , 2014, , 45-60.	0.1	3
89	A comparative study of ultra-deep pyrosequencing and cloning to quantitatively analyze the viral quasispecies using hepatitis B virus infection as a model. <i>Antiviral Research</i> , 2013, 98, 273-283.	4.1	53
90	The β 20 Isoform of the Complement Regulator C4b-Binding Protein Induces a Semimature, Anti-Inflammatory State in Dendritic Cells. <i>Journal of Immunology</i> , 2013, 190, 2857-2872.	0.8	33

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91	Exhaustive and non-exhaustive variation with free choice and referential vagueness: Evidence from Greek, Catalan, and Spanish. <i>Lingua</i> , 2013, 126, 120-149.	1.0	26
92	Response of Hepatitis C Virus to Long-Term Passage in the Presence of Alpha Interferon: Multiple Mutations and a Common Phenotype. <i>Journal of Virology</i> , 2013, 87, 7593-7607.	3.4	88
93	Extinction of Hepatitis C Virus by Ribavirin in Hepatoma Cells Involves Lethal Mutagenesis. <i>PLoS ONE</i> , 2013, 8, e71039.	2.5	60
94	Ultra-Deep Pyrosequencing (UDPS) Data Treatment to Study Amplicon HCV Minor Variants. <i>PLoS ONE</i> , 2013, 8, e83361.	2.5	54
95	Impersonal reference in Catalan Sign Language (LSC). , 2013, , 237-258.		11
96	Molecular epidemiology and putative origin of hepatitis C virus in random volunteers from Argentina. <i>World Journal of Gastroenterology</i> , 2013, 19, 5813.	3.3	15
97	T cell responses and viral variability in blood donation candidates with occult hepatitis B infection. <i>Journal of Hepatology</i> , 2012, 56, 765-774.	3.7	58
98	Ultra-Deep Pyrosequencing Detects Conserved Genomic Sites and Quantifies Linkage of Drug-Resistant Amino Acid Changes in the Hepatitis B Virus Genome. <i>PLoS ONE</i> , 2012, 7, e37874.	2.5	49
99	Legal Pathways to the Recognition of Sign Languages: A Comparison of the Catalan and Spanish Sign Language Acts. <i>Sign Language Studies</i> , 2012, 12, 565-582.	0.3	49
100	Quasispecies dynamics in main core epitopes of hepatitis B virus by ultra-deep-pyrosequencing. <i>World Journal of Gastroenterology</i> , 2012, 18, 6096.	3.3	15
101	Quantificational Strategies across Language Modalities. <i>Lecture Notes in Computer Science</i> , 2012, , 82-91.	1.3	38
102	Abacavir coadministration does not interfere with the suppressive activity of ribavirin in an HCV replicon system. <i>Antiviral Therapy</i> , 2011, 16, 887-893.	1.0	2
103	Ultra-deep pyrosequencing analysis of the hepatitis B virus preCore region and main catalytic motif of the viral polymerase in the same viral genome. <i>Nucleic Acids Research</i> , 2011, 39, 8457-8471.	14.5	42
104	When agreeing to disagree is not enough: Further arguments for the linguistic status of sign language agreement. <i>Theoretical Linguistics</i> , 2011, 37, .	0.2	14
105	HBV core region variability: effect of antiviral treatments on main epitopic regions. <i>Antiviral Therapy</i> , 2011, 16, 37-49.	1.0	14
106	Quantitative longitudinal evaluations of hepatitis delta virus RNA and hepatitis B virus DNA shows a dynamic, complex replicative profile in chronic hepatitis B and D. <i>Journal of Hepatology</i> , 2010, 52, 658-664.	3.7	155
107	The dynamic DNA methylomes of double-stranded DNA viruses associated with human cancer. <i>Genome Research</i> , 2009, 19, 438-451.	5.5	218
108	Hepatitis C virus (HCV)â€specific Tâ€cell responses among recombinant immunoblot assayâ€indeterminate blood donors: a confirmatory evidence of HCV exposure. <i>Transfusion</i> , 2009, 49, 1296-1305.	1.6	23

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109	Mood management: An updated toolkit. <i>Lingua</i> , 2009, 119, 1909-1913.	1.0	6
110	Transmission of low-density hepatitis C viral particles during sexually transmitted acute resolving infection. <i>Journal of Medical Virology</i> , 2008, 80, 242-246.	5.0	12
111	Naturally occurring NS3-protease-inhibitor resistant mutant A156T in the liver of an untreated chronic hepatitis C patient. <i>Virology</i> , 2008, 370, 237-245.	2.4	67
112	Lexical access in Catalan Signed Language (LSC) production. <i>Cognition</i> , 2008, 108, 856-865.	2.2	91
113	The changing epidemiology of hepatitis C virus infection in Europe. <i>Journal of Hepatology</i> , 2008, 48, 148-162.	3.7	379
114	Prevalence of hepatitis C in Romania: Different from European rates? Reply. <i>Journal of Hepatology</i> , 2008, 49, 663.	3.7	0
115	Nosocomial transmission of hepatitis C virus during contrast-enhanced computed tomography scanning. <i>European Journal of Gastroenterology and Hepatology</i> , 2008, 20, 73-78.	1.6	17
116	Valency in classifier predicates: A syntactic analysis. <i>Lingua</i> , 2007, 117, 1202-1215.	1.0	17
117	Induction of potent and long-lasting CD4 and CD8 T-cell responses against hepatitis C virus by immunization with viral antigens plus poly(I:C) and anti-CD40. <i>Antiviral Research</i> , 2007, 74, 25-35.	4.1	12
118	Effect of Bottlenecking on Evolution of the Nonstructural Protein 3 Gene of Hepatitis C Virus during Sexually Transmitted Acute Resolving Infection. <i>Journal of Virology</i> , 2005, 79, 15131-15141.	3.4	31
119	Improved Attachment of Natural HCV Isolate to Daudi Cells upon Elimination of Immune Complexes and Close pH Control. <i>Intervirology</i> , 2005, 48, 285-291.	2.8	4
120	Sexual Transmission of Hepatitis C Virus From a Patient With Chronic Disease to His Sex Partner After Removal of an Intrauterine Device. <i>Sexually Transmitted Diseases</i> , 2003, 30, 470-471.	1.7	19
121	The Syntax and Semantics of Unselected Embedded Questions. <i>Language</i> , 2001, 77, 107-133.	0.6	50
122	Contingent Neutrality in Competing Viral Populations. <i>Journal of Virology</i> , 2001, 75, 7315-7320.	3.4	41
123	Interpreting mood. <i>Probus</i> , 2001, 13, .	0.2	51
124	Genetic alterations in the S gene of hepatitis B virus in patients with acute hepatitis B, chronic hepatitis B and hepatitis B liver cirrhosis before and after liver transplantation. <i>Liver International</i> , 1999, 19, 177-182.	3.9	42
125	Red Queen Dynamics, Competition and Critical Points in a Model of RNA Virus Quasispecies. <i>Journal of Theoretical Biology</i> , 1999, 198, 47-59.	1.7	56
126	Hepatitis delta genotypes in chronic delta infection in the northeast of Spain (Catalonia). <i>Journal of Hepatology</i> , 1998, 28, 971-977.	3.7	29

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127	In the Cause of Subjunctive. <i>Linguistics in the Netherlands</i> , 1997, 14, 171-182.	0.1	1
128	Reproducible Nonlinear Population Dynamics and Critical Points During Replicative Competitions of RNA Virus Quasispecies. <i>Journal of Molecular Biology</i> , 1996, 264, 465-471.	4.2	70
129	Transmission of Hepatitis C Virus by a Cardiac Surgeon. <i>New England Journal of Medicine</i> , 1996, 334, 555-561.	27.0	360
130	Hepatitis C virus infection in renal transplant recipients: epidemiology, clinical impact, serological confirmation and viral replication. <i>Journal of Hepatology</i> , 1995, 22, 272-277.	3.7	30
131	Transmission of sign languages in Mediterranean Europe. , 0, , 95-112.		5
132	Nonmanuals: their grammatical and prosodic roles. , 0, , 381-402.		192
133	Language policy and planning in Deaf communities. , 0, , 120-145.		5
134	Subjunctives. , 0, , 660-684.		10
135	Nominal referential values of semantic classifiers and role shift in signed narratives. <i>Linguistik Aktuell</i> , 0, , 251-274.	0.6	5
136	Searching for imperatives in European sign languages. <i>Studies in Language Companion Series</i> , 0, , .	0.4	22
137	On the syntax of negation and modals in Catalan Sign Language and German Sign Language. <i>Trends in Linguistics Studies and Monographs</i> , 0, , .	0.1	58