## Jeremy A Garson

List of Publications by Year in descending order

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48 papers 15,901 citations

28 h-index 206112 48 g-index

53 all docs 53 docs citations

53 times ranked 26356 citing authors

#	Article	IF	Citations
1	The MIQE Guidelines: Minimum Information for Publication of Quantitative Real-Time PCR Experiments. Clinical Chemistry, 2009, 55, 611-622.	3.2	12,487
2	The Digital MIQE Guidelines: Minimum Information for Publication of Quantitative Digital PCR Experiments. Clinical Chemistry, 2013, 59, 892-902.	3.2	723
3	Diagnosis of Neuroinvasive Astrovirus Infection in an Immunocompromised Adult With Encephalitis by Unbiased Next-Generation Sequencing. Clinical Infectious Diseases, 2015, 60, 919-923.	5.8	262
4	The need for transparency and good practices in the qPCR literature. Nature Methods, 2013, 10, 1063-1067.	19.0	251
5	The Digital MIQE Guidelines Update: Minimum Information for Publication of Quantitative Digital PCR Experiments for 2020. Clinical Chemistry, 2020, 66, 1012-1029.	3.2	247
6	Differential susceptibility of PCR reactions to inhibitors: an important and unrecognised phenomenon. BMC Research Notes, 2008, 1, 70.	1.4	191
7	Disease-associated XMRV sequences are consistent with laboratory contamination. Retrovirology, 2010, 7, 111.	2.0	141
8	Real-time PCR quantitation of hepatitis B virus DNA using automated sample preparation and murine cytomegalovirus internal control. Journal of Virological Methods, 2005, 126, 207-213.	2.1	134
9	Comparative Study of Sensitivity, Linearity, and Resistance to Inhibition of Digital and Nondigital Polymerase Chain Reaction and Loop Mediated Isothermal Amplification Assays for Quantification of Human Cytomegalovirus. Analytical Chemistry, 2014, 86, 4387-4394.	6.5	126
10	Detection of SARS Coronavirus in Plasma by Real-Time RT-PCR. New England Journal of Medicine, 2003, 349, 2468-2469.	27.0	124
11	Novel non-isotopic in situ hybridization technique detects small (1 Kb) unique sequences in routinely G-banded human chromosomes: fine mapping of N-myc and $\hat{I}^2$ -NGF genes. Nucleic Acids Research, 1987, 15, 4761-4770.	14.5	122
12	Minor groove binder modification of widely used TaqMan probe for hepatitis E virus reduces risk of false negative real-time PCR results. Journal of Virological Methods, 2012, 186, 157-160.	2.1	101
13	Evaluation of diagnostic methods for the detection of cytomegalovirus in recipients of allogeneic stem cell transplants. Journal of Clinical Virology, 2001, 20, 59-70.	3.1	68
14	Primer Sequence Disclosure: A Clarification of the MIQE Guidelines. Clinical Chemistry, 2011, 57, 919-921.	3.2	63
15	Transmission of hepatitis C virus by organ transplantation in the United Kingdom. Journal of Hepatology, 1994, 20, 768-772.	3.7	55
16	Detection of Rare Drug Resistance Mutations by Digital PCR in a Human Influenza A Virus Model System and Clinical Samples. Journal of Clinical Microbiology, 2016, 54, 392-400.	3.9	52
17	Application of a novel in vitro selection technique to isolate and characterise high affinity DNA aptamers binding mammalian prion proteins. Journal of Virological Methods, 2008, 151, 107-115.	2.1	47
18	Cautionary Note on Contamination of Reagents Used for Molecular Detection of SARS-CoV-2. Clinical Chemistry, 2020, 66, 1369-1372.	3.2	46

#	Article	IF	CITATIONS
19	Quantitative analysis of human endogenous retrovirus-K transcripts in postmortem premotor cortex fails to confirm elevated expression of HERV-K RNA in amyotrophic lateral sclerosis. Acta Neuropathologica Communications, 2019, 7, 45.	5.2	44
20	Analysis of XMRV integration sites from human prostate cancer tissues suggests PCR contamination rather than genuine human infection. Retrovirology, 2011, 8, 13.	2.0	43
21	Digital PCR dynamic range is approaching that of real-time quantitative PCR. Biomolecular Detection and Quantification, 2016, 10, 31-33.	7.0	43
22	Development and evaluation of a real-time RT-PCR assay for quantification of cell-free human immunodeficiency virus type 2 using a Brome Mosaic Virus internal control. Journal of Virological Methods, 2006, 135, 102-108.	2.1	41
23	Quality Control Assessment of Human Immunodeficiency Virus Type 2 (HIV-2) Viral Load Quantification Assays: Results from an International Collaboration on HIV-2 Infection in 2006. Journal of Clinical Microbiology, 2008, 46, 2088-2091.	3.9	40
24	Instability of 8E5 calibration standard revealed by digital PCR risks inaccurate quantification of HIV DNA in clinical samples by qPCR. Scientific Reports, 2017, 7, 1209.	3.3	35
25	Development of an ethidium monoazide–enhanced internally controlled universal 16S rDNA realâ€time polymerase chain reaction assay for detection of bacterial contamination in platelet concentrates. Transfusion, 2012, 52, 1423-1432.	1.6	33
26	Development of an improved RT-qPCR Assay for detection of Japanese encephalitis virus (JEV) RNA including a systematic review and comprehensive comparison with published methods. PLoS ONE, 2018, 13, e0194412.	2.5	32
27	A Second Outbreak of Hepatitis C Virus Infection from Anti-D Immunoglobulin in Ireland. Vox Sanguinis, 1999, 76, 175-180.	1.5	31
28	Unreliable Real-Time PCR Analysis of Human Endogenous Retrovirus-W (HERV-W) RNA Expression and DNA Copy Number in Multiple Sclerosis. AIDS Research and Human Retroviruses, 2009, 25, 377-378.	1.1	29
29	An International Collaboration To Standardize HIV-2 Viral Load Assays: Results from the 2009 ACHI E V 2E Quality Control Study. Journal of Clinical Microbiology, 2011, 49, 3491-3497.	3.9	29
30	No Evidence of XMRV or Related Retroviruses in a London HIV-1-Positive Patient Cohort. PLoS ONE, 2011, 6, e18096.	2.5	25
31	Development and clinical application of a fully controlled quantitative PCR assay for cell-free cytomegalovirus in human plasma. Journal of Clinical Virology, 2003, 26, 49-59.	3.1	24
32	Quantitation of hepatitis C virus using an in-house real-time reverse transcriptase polymerase chain reaction in plasma samples. Diagnostic Microbiology and Infectious Disease, 2008, 61, 415-420.	1.8	24
33	Lack of susceptibility of the cottontop tamarin to hepatitis C infection. Journal of Medical Virology, 1997, 52, 286-288.	5.0	23
34	Proof-of-Principle for Immune Control of Global HIV-1 Reactivation In Vivo. Clinical Infectious Diseases, 2015, 61, 120-128.	5.8	17
35	Optimized protocol for a quantitative SARS-CoV-2 duplex RT-qPCR assay with internal human sample sufficiency control. Journal of Virological Methods, 2021, 294, 114174.	2.1	16
36	A Second Outbreak of Hepatitis C Virus Infection from Anti–D Immunoglobulin in Ireland. Vox Sanguinis, 1999, 76, 175-180.	1.5	16

#	Article	IF	CITATIONS
37	Highly sensitive and specific detection of the SARS-CoV-2 Delta variant by double-mismatch allele-specific real time reverse transcription PCR. Journal of Clinical Virology, 2022, 146, 105049.	3.1	15
38	A HML6 endogenous retrovirus on chromosome 3 is upregulated in amyotrophic lateral sclerosis motor cortex. Scientific Reports, 2021, 11, 14283.	3.3	13
39	Complement fixation by IgM and IgG autoantibodies on cultured human glial cells. Journal of Neurosurgery, 1981, 55, 19-26.	1.6	11
40	Development and evaluation of an internally controlled semiautomated PCR assay for quantification of cell-free cytomegalovirus. Journal of Medical Virology, 2002, 66, 518-523.	5.0	11
41	Multiple sclerosis-associated retrovirus and related human endogenous retrovirus-W in patients with multiple sclerosis. Journal of Neuroimmunology, 2014, 266, 87-88.	2.3	11
42	Evaluation of an ethidium monoazide–enhanced 16 S r DNA realâ€time polymerase chain reaction assay for bacterial screening of platelet concentrates and comparison with automated culture. Transfusion, 2014, 54, 870-878.	1.6	9
43	Response to the Letter from Garcia-Montojo and colleagues concerning our paper entitled, Quantitative analysis of human endogenous retrovirus-K transcripts in postmortem premotor cortex fails to confirm elevated expression of HERV-K RNA in amyotrophic lateral sclerosis. Acta Neuropathologica Communications, 2019, 7, 102.	5.2	9
44	Absence of human retrovirus 5 in French patients with rheumatoid arthritis: Comment on the article by Griffiths et al. Arthritis and Rheumatism, 1999, 42, 2492-2493.	6.7	8
45	Viral RNA Degradation Makes Urine a Challenging Specimen for Detection of Japanese Encephalitis Virus in Patients With Suspected CNS Infection. Open Forum Infectious Diseases, 2019, 6, ofz048.	0.9	7
46	Whole Blood as an Alternative to Plasma for Detection of Hepatitis C Virus RNA. Journal of Clinical Microbiology, 2008, 46, 3791-3794.	3.9	5
47	Minor groove binder modification of widely used TaqMan hydrolysis probe for detection of dengue virus reduces risk of false-negative real-time PCR results for serotype 4. Journal of Virological Methods, 2019, 268, 17-23.	2.1	2
48	Lack of susceptibility of the cottontop tamarin to hepatitis C infection., 1997, 52, 286.		2