Stephen Sc Chim

List of Publications by Year in descending order

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44 papers 4,194 citations

218381 26 h-index 253896 43 g-index

44 all docs

44 docs citations

44 times ranked 5770 citing authors

#	Article	IF	CITATIONS
1	Detection and Characterization of Placental MicroRNAs in Maternal Plasma. Clinical Chemistry, 2008, 54, 482-490.	1.5	775
2	Molecular Evolution of the SARS Coronavirus During the Course of the SARS Epidemic in China. Science, 2004, 303, 1666-1669.	6.0	695
3	Hypermethylated RASSF1A in Maternal Plasma: A Universal Fetal DNA Marker that Improves the Reliability of Noninvasive Prenatal Diagnosis. Clinical Chemistry, 2006, 52, 2211-2218.	1.5	319
4	Detection of the placental epigenetic signature of the maspin gene in maternal plasma. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 14753-14758.	3.3	307
5	Detection of SARS Coronavirus RNA in the Cerebrospinal Fluid of a Patient with Severe Acute Respiratory Syndrome. Clinical Chemistry, 2003, 49, 2108-2109.	1.5	233
6	Tissue and cellular tropism of the coronavirus associated with severe acute respiratory syndrome: an in-situ hybridization study of fatal cases. Journal of Pathology, 2004, 202, 157-163.	2.1	168
7	Noninvasive Prenatal Detection of Fetal Trisomy 18 by Epigenetic Allelic Ratio Analysis in Maternal Plasma: Theoretical and Empirical Considerations. Clinical Chemistry, 2006, 52, 2194-2202.	1.5	156
8	Quantitative Analysis and Prognostic Implication of SARS Coronavirus RNA in the Plasma and Serum of Patients with Severe Acute Respiratory Syndrome. Clinical Chemistry, 2003, 49, 1976-1980.	1.5	148
9	Hypermethylation of RASSF1A in Human and Rhesus Placentas. American Journal of Pathology, 2007, 170, 941-950.	1.9	128
10	Systematic Search for Placental DNA-Methylation Markers on Chromosome 21: Toward a Maternal Plasma-Based Epigenetic Test for Fetal Trisomy 21. Clinical Chemistry, 2008, 54, 500-511.	1.5	123
11	Systematic micro-array based identification of placental mRNA in maternal plasma: towards non-invasive prenatal gene expression profiling. Journal of Medical Genetics, 2004, 41, 461-467.	1.5	122
12	Endometrial microbiota in infertile women with and without chronic endometritis as diagnosed using a quantitative and reference range-based method. Fertility and Sterility, 2019, 112, 707-717.e1.	0.5	86
13	Serum Proteomic Fingerprints of Adult Patients with Severe Acute Respiratory Syndrome. Clinical Chemistry, 2006, 52, 421-429.	1.5	83
14	ACE2 Gene Polymorphisms Do Not Affect Outcome of Severe Acute Respiratory Syndrome. Clinical Chemistry, 2004, 50, 1683-1686.	1.5	76
15	Coronavirus Genomic-Sequence Variations and the Epidemiology of the Severe Acute Respiratory Syndrome. New England Journal of Medicine, 2003, 349, 187-188.	13.9	68
16	Genomic characterisation of the severe acute respiratory syndrome coronavirus of Amoy Gardens outbreak in Hong Kong. Lancet, The, 2003, 362, 1807-1808.	6.3	66
17	Quantitative aberrations of hypermethylated <i>RASSF1A</i> gene sequences in maternal plasma in preâ€eclampsia. Prenatal Diagnosis, 2007, 27, 1212-1218.	1.1	66
18	Systematic Comparison of Bacterial Colonization of Endometrial Tissue and Fluid Samples in Recurrent Miscarriage Patients: Implications for Future Endometrial Microbiome Studies. Clinical Chemistry, 2018, 64, 1743-1752.	1.5	45

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19	Systematic Identification of Spontaneous Preterm Birth-Associated RNA Transcripts in Maternal Plasma. PLoS ONE, 2012, 7, e34328.	1.1	43
20	Absence of association between angiotensin converting enzyme polymorphism and development of adult respiratory distress syndrome in patients with severe acute respiratory syndrome: a case control study. BMC Infectious Diseases, 2005, 5, 26.	1.3	41
21	Tracing SARS-Coronavirus Variant with Large Genomic Deletion. Emerging Infectious Diseases, 2005, 11, 168-170.	2.0	40
22	CHD5 Downregulation Associated with Poor Prognosis in Epithelial Ovarian Cancer. Gynecologic and Obstetric Investigation, 2011, 72, 203-207.	0.7	37
23	Characterization of a brain-specific nuclear LIM domain protein (FHL1B) which is an alternatively spliced variant of FHL1. Gene, 1999, 237, 253-263.	1.0	36
24	Calcyclin binding protein promotes DNA synthesis and differentiation in rat neonatal cardiomyocytes. Journal of Cellular Biochemistry, 2006, 98, 555-566.	1.2	33
25	Modulation by simvastatin of iberiotoxin-sensitive, Ca2+ -activated K+ channels of porcine coronary artery smooth muscle cells. British Journal of Pharmacology, 2007, 151, 987-997.	2.7	30
26	Epigenetic regulation of neonatal cardiomyocytes differentiation. Biochemical and Biophysical Research Communications, 2010, 400, 278-283.	1.0	28
27	Presence of Donor-Derived DNA and Cells in the Urine of Sex-Mismatched Hematopoietic Stem Cell Transplant Recipients: Implication for the Transrenal Hypothesis. Clinical Chemistry, 2009, 55, 715-722.	1.5	27
28	Systematic Selection of Reference Genes for the Normalization of Circulating RNA Transcripts in Pregnant Women Based on RNA-Seq Data. International Journal of Molecular Sciences, 2017, 18, 1709.	1.8	26
29	Systematic Identification of Placental Epigenetic Signatures for the Noninvasive Prenatal Detection of Edwards Syndrome. PLoS ONE, 2010, 5, e15069.	1.1	25
30	Proteomic analysis reveals platelet factor 4 and beta-thromboglobulin as prognostic markers in severe acute respiratory syndrome. Electrophoresis, 2012, 33, 1894-1900.	1.3	23
31	Molecular Epidemiology of SARS â€" From Amoy Gardens to Taiwan. New England Journal of Medicine, 2003, 349, 1875-1876.	13.9	21
32	Chromosomal Mapping of a Skeletal Muscle Specific LIM-Only Protein FHL3 to the Distal End of the Short Arm of Human Chromosome 1. Somatic Cell and Molecular Genetics, 1998, 24, 197-202.	0.7	19
33	Whole genome miRNA profiling revealed miR-199a as potential placental pathogenesis of selective fetal growth restriction in monochorionic twin pregnancies. Placenta, 2020, 92, 44-53.	0.7	18
34	Serum Amyloid A Is Not Useful in the Diagnosis of Severe Acute Respiratory Syndrome. Clinical Chemistry, 2006, 52, 1202-1204.	1.5	14
35	Reply to "STOX1 is not imprinted and is not likely to be involved in preeclampsia― Nature Genetics, 2007, 39, 280-281.	9.4	13
36	Mapping of the Human Cysteine-Rich Intestinal Protein GeneCRIP1to the Human Chromosomal Segment 7q11.23. Genomics, 1998, 47, 419-422.	1.3	12

#	Article	lF	CITATIONS
37	Genomic Sequencing of a SARS Coronavirus Isolate That Predated the Metropole Hotel Case Cluster in Hong Kong. Clinical Chemistry, 2004, 50, 231-233.	1.5	12
38	Differential gene expression of rat neonatal heart analyzed by suppression subtractive hybridization and expressed sequence tag sequencing. Journal of Cellular Biochemistry, 2001, 80, 24-36.	1.2	10
39	Role of monoamine oxidases in the exaggerated 5-hydroxytryptamine-induced tension development of human isolated preeclamptic umbilical artery. European Journal of Pharmacology, 2009, 605, 129-137.	1.7	7
40	Molecular cloning and characterization of a RING-H2 finger protein, ANAPC11, the human homolog of yeast Apc11p. Journal of Cellular Biochemistry, 2001, 83, 249-258.	1.2	6
41	Expression of replication factor C 40-kDa subunit is down-regulated during neonatal development in rat ventricular myocardium. Journal of Cellular Biochemistry, 2000, 78, 533-540.	1.2	4
42	A simple and rapid approach for screening of SARS-coronavirus genotypes: an evaluation study. BMC Infectious Diseases, 2005, 5, 87.	1.3	4
43	Potential application of fetal epigenetic markers on the non-invasive prenatal detection of chromosomal abnormality. Clinical Chemistry and Laboratory Medicine, 2014, 52, 585-8.	1.4	1
44	606: Maternal HBeAg status and hepatitis B viral activity in HBsAg positive mothers. American Journal of Obstetrics and Gynecology, 2012, 206, S273.	0.7	0