

Shicheng Guo

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Awards on Publons

September 2019: Top 1% of reviewers in Molecular Biology and Genetics

September 2017: Top reviewers for University of California, San Diego (in field General)

September 2017: Top reviewers for University of California, San Diego (in field Biochemistry, Genetics and Molecular Biology)

September 2017: Top reviewers for journal 'Scientific Reports'

September 2017: Top reviewers in General September 2017: Top reviewers in Medicine

September 2016: Sentinels of Science: Biochemistry, Genetics and Molecular Biology

Peer Review Summary

Performed 135 reviews for journals including *Clinical Epigenetics* and *Frontiers in Genetics* between January 2009 and September 2019.

1 of these reviews was rated "Excellent" in quality by an editor.

27 Clinical Epigenetics

21 Frontiers in Genetics

15 Scientific Reports

12 Medicine 쉾 (1 excellent review)

Journal of Thoracic Oncology

5 Functional & Integrative Genomics

4 Tumor Biology

3 Oncotarget

3 BMC Genetics

Journal of Genetics and Genomics

Frontiers in Oncology

2 BMC Research Notes

wiley 2 Cancer

2 BioData Mining

Biomarkers

2 Pathology Research and Practice

2 Journal of Clinical Laboratory Analysis

Journal of Pediatric Endocrinology and Metabolism

Open Journal of Rheumatology and Autoimmune Diseases

Statistics and Its Interface

Plos One

Brain, Behavior, and Immunity

Cancer Letters

@ 1 Genes and Immunity

Epigenomics

Cancer Investigation

Molecular Biology Reports

Journal of International Medical Research

Journal of Thoracic Disease

- Frontiers in Cell and Developmental Biology

 Molecular Genetics & Genomic Medicine

 Journal of Cancer
- Disease Markers
- International Journal of Dermatology and Venereology

Publication List

2019

2017

2017

A list of all publications authored between January 2009 and September 2019.

2019 Frontiers in Genetics

2019 Blood

Scientific Reports

A gene-based recessive diplotype exome scan discovers FGF6, a novel hepcidin-regulating iron-metabolism gene

(5R)-5-Hydroxytriptolide (LLDT-8) induces substantial epigenetic mediated immune response network changes in fibroblast-like synoviocytes from rheumatoid arthritis patients

Genome-Wide DNA Methylation Profiles Reveal Common Epigenetic Patterns of Interferon-Related Genes in Multiple Autoimmune Diseases

The Journal of Rheumatology

May 2018 Journal of Investigative Dermatology

Associations of Multiple NOTCH4 Exonic Variants with Systemic Sclerosis

Genome-Wide DNA Methylation Analysis in Systemic Sclerosis Reveals Hypomethylation of IFN-Associated Genes in CD4+ and CD8+ T Cells

2018 Frontiers in Genetics

Identification of Hyper-Methylated Tumor Suppressor Genes-Based Diagnostic Panel for Esophageal Squamous Cell Carcinoma (ESCC) in a Chinese Han Population

2018 Clinical Rheumatology

Increased DOT1L in synovial biopsies of patients with OA and RA

2018 FASEB Journal

Hypoxic-stabilized EPAS1 proteins transactivate DNMT1 and cause promoter hypermethylation and transcription inhibition of EPAS1 in non-small cell lung cancer

2018 Cell Death and Disease

Epigenetic silencing of ZNF132 mediated by methylation-sensitive Sp1 binding promotes cancer progression in esophageal squamous cell carcinoma

Genetic variant of miR-4293 rs12220909 is associated with susceptibility to non-small cell lung cancer in a Chinese Han population

2017 Modern Rheumatology
 Genome-wide DNA methylation patterns in CD4+T cells from Chinese Han patients with rheumatoid arthritis

2017 International Journal of Clinical and Experimental Pathology

Alterations and diagnosis potential of serum lipid profiles in rheumatoid arthritis patients

2017 Oncotarget

Quantitative assessment of the diagnostic role of FHIT promoter methylation in non-small cell lung cancer

Identification of methylation haplotype blocks aids in deconvolution of heterogeneous tissue samples and tumor tissue-of-origin mapping from plasma DNA

2017 Journal of Solid State Electrochemistry

A new lead single flow battery in a composite perchloric acid system with high specific surface capacity for large-scale energy storage

2017 Clinical Epigenetics

Toggeted highlifte aggregates identified a panel of DNA methylation based highertylers for econhaggel aggregate call aggregate (ESCC)

Targeted bisulfite sequencing identified a panel of DNA methylation-based biomarkers for esophageal squamous cell carcinoma (ESCC)

Hdac7 promotes lung tumorigenesis by inhibiting Stat3 activation

Molecular Cancer

Arthritis & Rheumatology

Genome-Wide DNA Methylation Analysis in Systemic Sclerosis Reveals Hypomethylation of Interferon-Associated Genes in CD4+and CD8+T Cells

Jun 2016 Oncology Letters

Hypermethylation reduces the expression of PNPLA7 in hepatocellular carcinoma

Apr 2016 The Journal of Rheumatology

Copy Number Variation of HLA-DQA1 and APOBEC3A/3B Contribute to the Susceptibility of Systemic Sclerosis in the Chinese Han Population

2016 Journal of Cancer

Aberrant methylation of CDH13 can be a diagnostic biomarker for lung adenocarcinoma

2016 Clinical Epigenetics

The detective, prognostic, and predictive value of DNA methylation in human esophageal squamous cell carcinoma

nature 2016 Nature

In vivo genome editing via CRISPR/Cas9 mediated homology-independent targeted integration

2016 Clinical Epigenetics

Obesity-related DNA methylation at imprinted genes in human sperm: Results from the TIEGER study

2016 Tumor Biology

Genetic variants in miR-196a2 and miR-499 are associated with susceptibility to esophageal squamous cell carcinoma in Chinese Han population

📳 Sep 2015 Oncology Letters

Abnormal methylation status of FBXW10 and SMPD3, and associations with clinical characteristics in clear cell renal cell carcinoma

Aug 2015 Quantitative Biology

Multiple functional linear model for association analysis of RNA-seq with imaging

May 2015 Journal of Thoracic Oncology

Erratum

2015 Plos One

Functional Principal Component Analysis and Randomized Sparse Clustering Algorithm for Medical Image Analysis

witev 2015 Arthritis & Rheumatology

Genome-Wide DNA Methylation Patterns in CD4+T Reveal Significant Contribution of DNA Methylation to Rheumatoid Arthritis

Z015 Tumor Biology

miR-449b rs10061133 and miR-4293 rs12220909 polymorphisms are associated with decreased esophageal squamous cell carcinoma in a Chinese population

pp 2015 European Journal of Human Genetics

Positional cloning and next-generation sequencing identified a TGM6 mutation in a large Chinese pedigree with acute myeloid leukaemia

A Stress-Related Chromosome Region, Contributes to Reducing Lung Squamous Cell Carcinoma Risk (vol 7, pg 1041, 2014)

2015 Clinical Epigenetics

Identification and validation of the methylation biomarkers of non-small cell lung cancer (NSCLC)

2015 International Journal of Rheumatic Diseases

Association between ABCG2 Q141K polymorphism and gout risk affected by ethnicity and gender: a systematic review and meta-analysis

Apr 2014 Investigative Ophthalmology & Visual Science

Different Hereditary Contribution of the CFH Gene Between Polypoidal Choroidal Vasculopathy and Age-Related Macular Degeneration in Chinese Han People

Mar 2014 Cancer Medicine

Significant SNPs have limited prediction ability for thyroid cancer

2014 Plos One

Association of the HLA-DRB1 with Scleroderma in Chinese Population

2014 Journal of Genetics

Association between HLA-DQA1 gene copy number polymorphisms and susceptibility to rheumatoid arthritis in Chinese Han population

2014 Clinical Epigenetics

Genome-wide methylation profiling of the different stages of hepatitis B virus-related hepatocellular carcinoma development in plasma cell-free DNA reveals potential biomarkers for early detection and high-risk monitoring of hepatocellular carcinoma

Quantitative assessment of the diagnostic role of APC promoter methylation in non-small cell lung cancer

2014 Clinical Epigenetics

 $High-frequency\ aberrantly\ methylated\ targets\ in\ pancreatic\ adenocarcinoma\ identified\ via\ global\ DNA\ methylation\ analysis\ using\ methyl Cap-seq$

Acta Biochimica et Biophysica Sinica

Immunological effects of interleukin 12 administered by bolus intravenous injection to patients with cancer

Molecular Medicine Reports

Association study of miR-149 rs2292832 and miR-608 rs4919510 and the risk of hepatocellular carcinoma in a large-scale population

9q33.3, A Stress-Related Chromosome Region, Contributes to Reducing Lung Squamous Cell Carcinoma Risk

M Sep 2013 Genes & Immunity

Association between copy number variations of HLA-DQA1 and ankylosing spondylitis in the Chinese Han population

BMJ Jul 2013 Journal of Medical Genetics

Confirmation of papillary thyroid cancer susceptibility loci identified by genome-wide association studies of chromosomes 14q13, 9q22, 2q35 and 8p12 in a Chinese population

묡 Jan 2013 Oncology Letters

Abnormal methylation of seven genes and their associations with clinical characteristics in early stage non-small cell lung cancer

2013 FASEB Journal

Hypermethylation reduces expression of tumor-suppressor PLZF and regulates proliferation and apoptosis in non-small-cell lund cancers

2013 Plos One

Prognostic Role of MicroRNA-181a/b in Hematological Malignancies: A Meta-Analysis

Jun 2012 Clinical Cancer Research

Predictive Value of XRCC1 Gene Polymorphisms on Platinum-Based Chemotherapy in Advanced Non-Small Cell Lung Cancer Patients: A Systematic Review and Meta-analysis

2012 Acta Diabetologica

Quantitative assessment of the variation in IGF2BP2 gene and type 2 diabetes risk

2012 Plos One

Methylcap-Seq Reveals Novel DNA Methylation Markers for the Diagnosis and Recurrence Prediction of Bladder Cancer in a Chinese Population

2011 Neonlasia

Hypomethylation of the hsa-miR-191 Locus Causes High Expression of hsa-miR-191 and Promotes the Epithelial-to-Mesenchymal Transition in Hepatocellular Carcinoma

Jul 2010 Nature Biotechnology

Erratum: Single base-resolution methylome of the silkworm reveals a sparse epigenomic map

May 2010 Nature Biotechnology

Single base-resolution methylome of the silkworm reveals a sparse epigenomic map

Zhonghua Zhong Liu za Zhi [Chinese Journal of Oncology]

[Correlation of the methylation status of CpG islands in the promoter region of 10 genes with the 5-Fu chemosensitivity in 3 breast cancer cell lines]

2010 PLoS Biology

The DNA Methylome of Human Peripheral Blood Mononuclear Cells