Fusheng Si

List of Publications by Year in descending order

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FUSHENC SI

#	Article	lF	CITATIONS
1	TLR8-Mediated Metabolic Control of Human Treg Function: A Mechanistic Target for Cancer Immunotherapy. Cell Metabolism, 2019, 29, 103-123.e5.	16.2	149
2	Reprogramming lipid metabolism prevents effector T cell senescence and enhances tumor immunotherapy. Science Translational Medicine, 2021, 13, .	12.4	101
3	Molecular detection of hepatitis E virus in sheep from southern Xinjiang, China. Virus Genes, 2015, 50, 410-417.	1.6	46
4	Porcine Epidemic Diarrhea Virus (PEDV) ORF3 Enhances Viral Proliferation by Inhibiting Apoptosis of Infected Cells. Viruses, 2020, 12, 214.	3.3	41
5	SALL1 functions as a tumor suppressor in breast cancer by regulating cancer cell senescence and metastasis through the NuRD complex. Molecular Cancer, 2018, 17, 78.	19.2	40
6	Baicalein and Baicalin Promote Melanoma Apoptosis and Senescence via Metabolic Inhibition. Frontiers in Cell and Developmental Biology, 2020, 8, 836.	3.7	29
7	NK and NKT cells have distinct properties and functions in cancer. Oncogene, 2021, 40, 4521-4537.	5.9	29
8	Tumor-derived ILT4 induces T cell senescence and suppresses tumor immunity. , 2021, 9, e001536.		23
9	Identification of host cell proteins that interact with the M protein of porcine epidemic diarrhea virus. Veterinary Microbiology, 2020, 246, 108729.	1.9	16
10	Citrate Promotes Excessive Lipid Biosynthesis and Senescence in Tumor Cells for Tumor Therapy. Advanced Science, 2022, 9, e2101553.	11.2	16
11	Study on the Characteristic Codon Usage Pattern in Porcine Epidemic Diarrhea Virus Genomes and Its Host Adaptation Phenotype. Frontiers in Microbiology, 2021, 12, 738082.	3.5	14
12	Reduced prevalence of genotype 3 HEV in Shanghai pig farms and hypothetical homeostasis of porcine HEV reservoir. Veterinary Microbiology, 2009, 137, 184-189.	1.9	12
13	Molecular epidemiology of hepatitis E virus infections in Shanghai, China. Virology Journal, 2011, 8, 541.	3.4	11
14	Porcine Epidemic Diarrhea Virus ORF3 Protein Is Transported through the Exocytic Pathway. Journal of Virology, 2020, 94, .	3.4	11
15	Infectivity of a genotype 4 hepatitis E virus cDNA clone by intrahepatic inoculation of laboratory rats. Veterinary Microbiology, 2013, 166, 405-411.	1.9	9
16	Identification of cellular proteins interacting with PEDV M protein through APEX2 labeling. Journal of Proteomics, 2021, 240, 104191.	2.4	8
17	Adaptation of Genotype 3 Hepatitis E Virus in Eastern China and Inverse Correlation with Genotype 4 Hepatitis E Virus. Intervirology, 2012, 55, 356-364.	2.8	6
18	Determination of the full-genome sequence of hepatitis E virus (HEV) SAAS-FX17 and use as a reference to identify putative HEV genotype 4 virulence determinants. Virology Journal, 2012, 9, 264.	3.4	4

#	Article	IF	CITATIONS
19	Construction of an Infectious cDNA Clone of a Swine Genotype 3 HEV Strain Isolated in Shanghai, China. Intervirology, 2014, 57, 74-82.	2.8	3