Klaus Lehnert

List of Publications by Year in descending order

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302126
39
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3198
citing authors

#	Article	IF	CITATIONS
1	Haploinsufficiency of the NF-κB1 Subunit p50 in Common Variable Immunodeficiency. American Journal of Human Genetics, 2015, 97, 389-403.	6.2	232
2	Cross-Comparison of Exome Analysis, Next-Generation Sequencing of Amplicons, and the iPLEX® ADME PGx Panel for Pharmacogenomic Profiling. Frontiers in Pharmacology, 2016, 7, 1.	3.5	231
3	Modulation of the maternal immune system by the pre-implantation embryo. BMC Genomics, 2010, 11, 474.	2.8	112
4	Sequence-based Association Analysis Reveals an MGST1 eQTL with Pleiotropic Effects on Bovine Milk Composition. Scientific Reports, 2016, 6, 25376.	3.3	103
5	De Novo Pathogenic Variants in CACNA1E Cause Developmental and Epileptic Encephalopathy with Contractures, Macrocephaly, and Dyskinesias. American Journal of Human Genetics, 2018, 103, 666-678.	6.2	87
6	Review: Diagnosing Common Variable Immunodeficiency Disorder in the Era of Genome Sequencing. Clinical Reviews in Allergy and Immunology, 2018, 54, 261-268.	6.5	69
7	Expression Variants of the Lipogenic AGPAT6 Gene Affect Diverse Milk Composition Phenotypes in Bos taurus. PLoS ONE, 2014, 9, e85757.	2.5	58
8	Epistatic interactions between mutations of TACI (<i>TNFRSF13B</i>) and <i>TCF3</i> result in a severe primary immunodeficiency disorder and systemic lupus erythematosus. Clinical and Translational Immunology, 2017, 6, e159.	3.8	54
9	MAdCAM-1 costimulates T cell proliferation exclusively through integrin î±4î²7, whereas VCAM-1 and CS-1 peptide use î±4β1: evidence for "remote―costimulation and induction of hyperresponsiveness to B7 molecules. European Journal of Immunology, 1998, 28, 3605-3615.	2.9	45
10	Clinical Implications of Digenic Inheritance and Epistasis in Primary Immunodeficiency Disorders. Frontiers in Immunology, 2017, 8, 1965.	4.8	44
11	Glycoprotein biosynthesis in Saccharomyces cerevisiae: ngd29, an N-glycosylation mutant allelic tooch 1 having a defect in the initiation of outer chain formation. FEBS Letters, 1995, 370, 41-45.	2.8	42
12	Brain dopamineâ€serotonin vesicular transport disease presenting as a severe infantile hypotonic parkinsonian disorder. Journal of Inherited Metabolic Disease, 2016, 39, 305-308.	3.6	41
13	Common Variable Immunodeficiency Disorders, T-Cell Responses to SARS-CoV-2 Vaccines, and the Risk of Chronic COVID-19. Journal of Allergy and Clinical Immunology: in Practice, 2021, 9, 3575-3583.	3.8	41
14	Metabolic proteomics of the liver and mammary gland during lactation. Journal of Proteomics, 2012, 75, 4429-4435.	2.4	39
15	Profiling the metabolic proteome of bovine mammary tissue. Proteomics, 2008, 8, 1502-1515.	2.2	32
16	Angiostatin enhances B7.1-mediated cancer immunotherapy independently of effects on vascular endothelial growth factor expression. Cancer Gene Therapy, 2001, 8, 719-727.	4.6	30
17	Cloning, Sequence Analysis, and Chromosomal Localization of the Novel Human Integrin $\hat{l}\pm 11$ Subunit (ITGA11). Genomics, 1999, 60, 179-187.	2.9	29
18	Proteomic Analysis of Microsomes from Lactating Bovine Mammary Gland. Journal of Proteome Research, 2008, 7, 1427-1432.	3.7	29

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19	Keeping it in the family: the case for considering late-onset combined immunodeficiency a subset of common variable immunodeficiency disorders. Expert Review of Clinical Immunology, 2018, 14, 549-556.	3.0	27
20	Estimation of genetic and crossbreeding parameters of fatty acid concentrations in milk fat predicted by mid-infrared spectroscopy in New Zealand dairy cattle. Journal of Dairy Research, 2014, 81, 340-349.	1.4	25
21	Genome-wide association studies of lactation yields of milk, fat, protein and somatic cell score in New Zealand dairy goats. Journal of Animal Science and Biotechnology, 2020, 11, 55.	5. 3	25
22	Perspective: diagnostic laboratories should urgently develop T cell assays for SARS-CoV-2 infection. Expert Review of Clinical Immunology, 2021, 17, 421-430.	3.0	24
23	LPAM-1 (integrin $\hat{l}\pm4\hat{l}^27$)-ligand binding: overlapping binding sites recognizing VCAM-1, MAdCAM-1 and CS-1 are blocked by fibrinogen, a fibronectin-like polymer and RGD-like cyclic peptides. European Journal of Immunology, 1998, 28, 995-1004.	2.9	23
24	Rapid, quantitative analysis of 3′- and 6′-sialyllactose in milk by flow-injection analysis–mass spectrometry: Screening of milks for naturally elevated sialyllactose concentration. Journal of Dairy Science, 2013, 96, 7684-7691.	3.4	21
25	Compound Heterozygous Inheritance of Mutations in Coenzyme Q8A Results in Autosomal Recessive Cerebellar Ataxia and Coenzyme Q10 Deficiency in a Female Sib-Pair. JIMD Reports, 2017, 42, 31-36.	1.5	21
26	Compound heterozygous <i>SLC19A3</i> mutations further refine the critical promoter region for biotin-thiamine-responsive basal ganglia disease. Journal of Physical Education and Sports Management, 2017, 3, a001909.	1.2	20
27	Evaluation of the performance of copy number variant prediction tools for the detection of deletions from whole genome sequencing data. Journal of Biomedical Informatics, 2019, 94, 103174.	4.3	20
28	Interaction of monocytoid cells with the mucosal addressin MAdCAM-1 via the integrins VLA-4 and LPAM-1. Immunology and Cell Biology, 1996, 74, 383-393.	2.3	18
29	All Patients With Common Variable Immunodeficiency Disorders (CVID) Should Be Routinely Offered Diagnostic Genetic Testing. Frontiers in Immunology, 2019, 10, 2678.	4.8	15
30	Perspective: Application of the American College of Medical Genetics Variant Interpretation Criteria to Common Variable Immunodeficiency Disorders. Clinical Reviews in Allergy and Immunology, 2021, 61, 226-235.	6.5	15
31	Phenotypic population screen identifies a new mutation in bovine DGAT1 responsible for unsaturated milk fat. Scientific Reports, 2015, 5, 8484.	3.3	14
32	Genetic parameters for total lactation yields of milk, fat, protein, and somatic cell score in New Zealand dairy goats. Animal Science Journal, 2020, 91, e13310.	1.4	14
33	An Immunological Approach to Increase the Brain's Resilience to Insults. ISRN Neuroscience, 2014, 2014, 1-10.	1.5	13
34	Induction of systemic antitumor immunity by gene transfer of mammalian heat shock protein 70.1 into tumors in situ. Cancer Gene Therapy, 2001, 8, 974-981.	4.6	12
35	Penetrance and expressivity of the R858H <i>CACNA1C</i> variant in a fiveâ€generation pedigree segregating an arrhythmogenic channelopathy. Molecular Genetics & Denomic Medicine, 2019, 7, e00476.	1.2	11
36	Perspective: the nose and the stomach play a critical role in the NZACE2-PÄŧari* (modified ACE2) drug treatment project of SARS-CoV-2 infection. Expert Review of Clinical Immunology, 2021, 17, 553-560.	3.0	10

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37	Severe COVID-19 is a T cell immune dysregulatory disorder triggered by SARS-CoV-2. Expert Review of Clinical Immunology, 2022, 18, 557-565.	3.0	10
38	Mouse M290 is the functional homologue of the human mucosal lymphocyte integrin HML-1: Antagonism between the integrin ligands E-cadherin and RGD tripeptide. Immunology and Cell Biology, 1999, 77, 337-344.	2.3	9
39	A Triad of Highly Divergent Polymeric Immunoglobulin Receptor (PIGR) Haplotypes with Major Effect on IgA Concentration in Bovine Milk. PLoS ONE, 2013, 8, e57219.	2.5	9
40	The (apparent) antibody paradox in COVID-19. Expert Review of Clinical Immunology, 2022, 18, 335-345.	3.0	9
41	Modelling brain dopamine-serotonin vesicular transport disease in <i>Caenorhabditis elegans</i> DMM Disease Models and Mechanisms, 2018, 11, .	2.4	8
42	Copy number variants implicate cardiac function and development pathways in earthquake-induced stress cardiomyopathy. Scientific Reports, 2018, 8, 7548.	3.3	8
43	Inhaled modified angiotensin converting enzyme 2 (ACE2) as a decoy to mitigate SARS-CoV-2 infection. New Zealand Medical Journal, 2020, 133, 112-118.	0.5	7
44	Novel PRMT7 mutation in a rare case of dysmorphism and intellectual disability. Journal of Human Genetics, 2021, , .	2.3	6
45	Are All Primary Immunodeficiency Disorders Inborn Errors of Immunity?. Frontiers in Immunology, 2021, 12, 706796.	4.8	6
46	Advantage of including Genomic Information to Predict Breeding Values for Lactation Yields of Milk, Fat, and Protein or Somatic Cell Score in a New Zealand Dairy Goat Herd. Animals, 2021, 11, 24.	2.3	6
47	Common Variable Immunodeficiency Disorders as a Model for Assessing COVID-19 Vaccine Responses in Immunocompromised Patients. Frontiers in Immunology, 2021, 12, 798389.	4.8	6
48	Response to letter to the editor: the clinical utility of diagnostic T cell assays for COVID-19. Expert Review of Clinical Immunology, 2021, 17, 1159-1161.	3.0	5
49	The New Zealand minds for minds autism spectrum disorder self-reported cohort. Research in Autism Spectrum Disorders, 2017, 36, 1-7.	1.5	4
50	Screening for phenotypic outliers identifies an unusually low concentration of a \hat{l}^2 -lactoglobulin B protein isoform in bovine milk caused by a synonymous SNP. Genetics Selection Evolution, 2022, 54, 22.	3.0	4
51	Whole Exome Sequencing Reveals Compound Heterozygosity for Ethnically Distinct PEX7 Mutations Responsible for Rhizomelic Chondrodysplasia Punctata, Type 1. Case Reports in Genetics, 2015, 2015, 1-4.	0.2	3
52	A Capra hircus chromosome 19 locus linked to milk production influences mammary conformation. Journal of Animal Science and Biotechnology, 2022, 13, 4.	5.3	3
53	MAdCAM-1 costimulates T cell proliferation exclusively through integrin $\hat{l}\pm4\hat{l}^27$, whereas VCAM-1 and CS-1 peptide use $\hat{l}\pm4\hat{l}^21$: evidence for "remote" costimulation and induction of hyperresponsiveness to B7 molecules. European Journal of Immunology, 1998, 28, 3605-3615.	2.9	1
54	RBV: Read balance validator, a tool for prioritising copy number variations in germline conditions. Scientific Reports, 2019, 9, 16934.	3.3	0

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55	Under-ascertainment of breast cancer susceptibility gene carriers in a cohort of New Zealand female breast cancer patients. Breast Cancer Research and Treatment, 2021, 185, 583-590.	2.5	O