Stefano Lise

List of Publications by Year in descending order

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STEENNO LISE

#	Article	IF	CITATIONS
1	Clinical <i>BRCA1/2</i> Reversion Analysis Identifies Hotspot Mutations and Predicted Neoantigens Associated with Therapy Resistance. Cancer Discovery, 2020, 10, 1475-1488.	9.4	109
2	The Immunogenic Potential of Recurrent Cancer Drug Resistance Mutations: An In Silico Study. Frontiers in Immunology, 2020, 11, 524968.	4.8	7
3	Extreme intratumour heterogeneity and driver evolution in mismatch repair deficient gastro-oesophageal cancer. Nature Communications, 2020, 11, 139.	12.8	44
4	Genome-wide plasma DNA methylation features of metastatic prostate cancer. Journal of Clinical Investigation, 2020, 130, 1991-2000.	8.2	68
5	Identification of single nucleotide variants using position-specific error estimation in deep sequencing data. BMC Medical Genomics, 2019, 12, 115.	1.5	10
6	Genomic and Transcriptomic Determinants of Therapy Resistance and Immune Landscape Evolution during Anti-EGFR Treatment in Colorectal Cancer. Cancer Cell, 2019, 36, 35-50.e9.	16.8	179
7	<i>In Vivo</i> Modeling of Chemoresistant Neuroblastoma Provides New Insights into Chemorefractory Disease and Metastasis. Cancer Research, 2019, 79, 5382-5393.	0.9	42
8	Mutations in MAST1 Cause Mega-Corpus-Callosum Syndrome with Cerebellar Hypoplasia and Cortical Malformations. Neuron, 2018, 100, 1354-1368.e5.	8.1	35
9	Ultra-Sensitive Mutation Detection and Genome-Wide DNA Copy Number Reconstruction by Error-Corrected Circulating Tumor DNA Sequencing. Clinical Chemistry, 2018, 64, 1626-1635.	3.2	46
10	Biallelic Mutation of ARHGEF18, Involved in the Determination of Epithelial Apicobasal Polarity, Causes Adult-Onset Retinal Degeneration. American Journal of Human Genetics, 2017, 100, 334-342.	6.2	26
11	A Novel Strategy Combining Array-CGH, Whole-exome Sequencing and In Utero Electroporation in Rodents to Identify Causative Genes for Brain Malformations. Journal of Visualized Experiments, 2017, , .	0.3	0
12	Making the most of RNA-seq: Pre-processing sequencing data with Opossum for reliable SNP variant detection. Wellcome Open Research, 2017, 2, 6.	1.8	36
13	Combination of Whole Genome Sequencing, Linkage, and Functional Studies Implicates a Missense Mutation in Titin as a Cause of Autosomal Dominant Cardiomyopathy With Features of Left Ventricular Noncompaction. Circulation: Cardiovascular Genetics, 2016, 9, 426-435.	5.1	67
14	Autosomal dominant osteopetrosis associated with renal tubular acidosis is due to a CLCN7 mutation. American Journal of Medical Genetics, Part A, 2016, 170, 2988-2992.	1.2	10
15	Mutations in REEP6 Cause Autosomal-Recessive Retinitis Pigmentosa. American Journal of Human Genetics, 2016, 99, 1305-1315.	6.2	121
16	Premalignant SOX2 overexpression in the fallopian tubes of ovarian cancer patients: Discovery and validation studies. EBioMedicine, 2016, 10, 137-149.	6.1	34
17	Biallelic Mutations in the Autophagy Regulator DRAM2 Cause Retinal Dystrophy with Early Macular Involvement. American Journal of Human Genetics, 2015, 96, 948-954.	6.2	42
18	Factors influencing success of clinical genome sequencing across a broad spectrum of disorders. Nature Genetics, 2015, 47, 717-726.	21.4	310

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19	Next-generation sequencing (NGS) as a diagnostic tool for retinal degeneration reveals a much higher detection rate in early-onset disease. European Journal of Human Genetics, 2013, 21, 274-280.	2.8	119
20	Periventricular heterotopia in 6q terminal deletion syndrome: role of the C6orf70 gene. Brain, 2013, 136, 3378-3394.	7.6	85
21	Next generation sequencing for molecular diagnosis of neurological disorders using ataxias as a model. Brain, 2013, 136, 3106-3118.	7.6	146
22	Recessive Mutations in SPTBN2 Implicate β-III Spectrin in Both Cognitive and Motor Development. PLoS Genetics, 2012, 8, e1003074.	3.5	94
23	PepSite: prediction of peptide-binding sites from protein surfaces. Nucleic Acids Research, 2012, 40, W423-W427.	14.5	174
24	Exome sequencing can detect pathogenic mosaic mutations present at low allele frequencies. Journal of Human Genetics, 2012, 57, 70-72.	2.3	58
25	Next-generation sequencing in health-care delivery: lessons from the functional analysis of rhodopsin. Genetics in Medicine, 2012, 14, 891-899.	2.4	28
26	Predictions of Hot Spot Residues at Protein-Protein Interfaces Using Support Vector Machines. PLoS ONE, 2011, 6, e16774.	2.5	78
27	Prediction of hot spot residues at protein-protein interfaces by combining machine learning and energy-based methods. BMC Bioinformatics, 2009, 10, 365.	2.6	100
28	Docking protein domains in contact space. BMC Bioinformatics, 2006, 7, 310.	2.6	14
29	The CATH Domain Structure Database and related resources Gene3D and DHS provide comprehensive domain family information for genome analysis. Nucleic Acids Research, 2004, 33, D247-D251.	14.5	226
30	Sequence patterns associated with disordered regions in proteins. Proteins: Structure, Function and Bioinformatics, 2004, 58, 144-150.	2.6	72
31	Nonconservative Earthquake Model of Self-Organized Criticality on a Random Graph. Physical Review Letters, 2002, 88, 228301.	7.8	42
32	Self-organization to criticality in a system without conservation law. Journal of Physics A, 2002, 35, 4641-4649.	1.6	9
33	Bethe approximation for self-interacting lattice trees. Europhysics Letters, 2001, 53, 176-182.	2.0	10
34	Scaling in a nonconservative earthquake model of self-organized criticality. Physical Review E, 2001, 64, 046111.	2.1	27
35	Comment on "Self-Organized Criticality in the Olami-Feder-Christensen Model― Physical Review Letters, 2001, 87, 039801	7.8	25
36	Self-organized criticality and universality in a nonconservative earthquake model. Physical Review E, 2001, 63, 036111.	2.1	57

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37	Langevin equations coupled through correlated noises. Journal of Physics A, 1999, 32, 5251-5260.	1.6	5
38	A non-equilibrium percolation transition in random Ising ferromagnets. Journal of Physics A, 1998, 31, L713-L719.	1.6	0
39	Phase diagram and critical behaviour of homopolymers with steric frustration. Journal of Physics A, 1998, 31, 6183-6188.	1.6	0
40	Bethe approximation for a semiflexible polymer chain. Physical Review E, 1998, 58, R5241-R5244.	2.1	28
41	Boundary effects in a random neighbor model of earthquakes. Physical Review E, 1998, 57, 3633-3636.	2.1	4
42	Nonconservative sandpile models. Physical Review E, 1997, 56, 6702-6709.	2.1	23
43	Transitions in Nonconserving Models of Self-Organized Criticality. Physical Review Letters, 1996, 76, 2326-2329.	7.8	45
44	Interfacial Properties of Interacting Surfaces. Europhysics Letters, 1995, 32, 735-740.	2.0	7