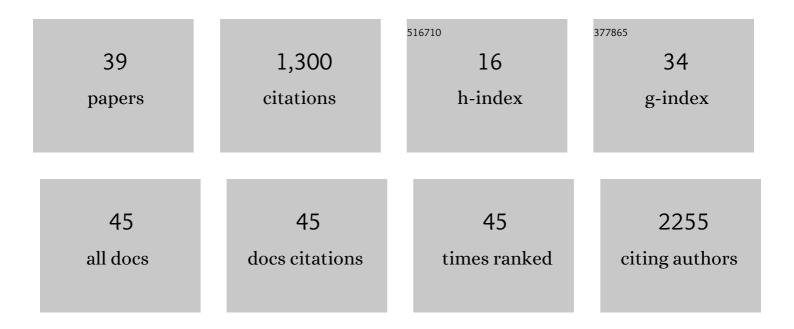
Godfrey Grech

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9337301/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	A randomized, controlled trial on the effect of anesthesia on chronic pain after total knee arthroplasty. Pain Management, 2022, 12, 711-723.	1.5	4
2	P234 The effect of calcitriol on the expression of interferon signature genes in dendritic cells and macrophages in systemic lupus erythematosus. Rheumatology, 2022, 61, .	1.9	0
3	Differential Expression of the Sphingolipid Pathway Is Associated with Sensitivity to the PP2A Activator FTY720 in Colorectal Cancer Cell Lines. Journal of Clinical Medicine, 2021, 10, 4999.	2.4	1
4	A loop involving NRF2, miRâ€29bâ€1â€5p and AKT, regulates cell fate of MDAâ€MBâ€231 tripleâ€negative breast cancer cells. Journal of Cellular Physiology, 2020, 235, 629-637.	4.1	34
5	Somatic copy number aberrations in metastatic patients: The promise of liquid biopsies. Seminars in Cancer Biology, 2020, 60, 302-310.	9.6	11
6	The landscape of genomic copy number alterations in colorectal cancer and their consequences on gene expression levels and disease outcome. Molecular Aspects of Medicine, 2019, 69, 48-61.	6.4	40
7	Aspirin impairs acetyl-coenzyme A metabolism in redox-compromised yeast cells. Scientific Reports, 2019, 9, 6152.	3.3	5
8	Loss of MCL1 function sensitizes the MDAâ€MBâ€231 breast cancer cells to rhâ€TRAIL by increasing DR4 levels. Journal of Cellular Physiology, 2019, 234, 18432-18447.	4.1	7
9	Bead-based RNA multiplex panels for biomarker detection in oncology samples. Methods, 2019, 158, 86-91.	3.8	4
10	Acquired and Intrinsic Resistance to Colorectal Cancer Treatment. , 2018, , .		6
11	Optimization of a Multiplex RNA-based Expression Assay Using Breast Cancer Archival Material. Journal of Visualized Experiments, 2018, , .	0.3	3
12	A novel PAX9 mutation causing oligodontia. Archives of Oral Biology, 2017, 84, 100-105.	1.8	8
13	CIP2A expression predicts recurrences of tamoxifen-treated breast cancer. Tumor Biology, 2017, 39, 101042831772206.	1.8	2
14	Suppressive role exerted by microRNA-29b-1-5p in triple negative breast cancer through SPIN1 regulation. Oncotarget, 2017, 8, 28939-28958.	1.8	57
15	Molecular Classification of Breast Cancer Patients Using Formalin-fixed Paraffin-embedded Derived RNA Samples. Journal of Molecular Biomarkers & Diagnosis, 2016, 01, .	0.4	1
16	A European Spectrum of Pharmacogenomic Biomarkers: Implications for Clinical Pharmacogenomics. PLoS ONE, 2016, 11, e0162866.	2.5	96
17	Deregulation of the protein phosphatase 2A, PP2A in cancer: complexity and therapeutic options. Tumor Biology, 2016, 37, 11691-11700.	1.8	46
18	Breast cancer epidemic in the early twenty-first century: evaluation of risk factors, cumulative questionnaires and recommendations for preventive measures. Tumor Biology, 2016, 37, 12941-12957.	1.8	108

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19	The effect of turmeric (Curcumin) supplementation on cytokine and inflammatory marker responses following 2 hours of endurance cycling. Journal of the International Society of Sports Nutrition, 2015, 12, 5.	3.9	51
20	Actionable pharmacogenetic markers for prediction and prognosis in breast cancer. EPMA Journal, 2015, 6, 15.	6.1	17
21	250 Large Deletion in the EPCAM Gene Responsible for the Milder Phenotype of Congenital Tufting Enteropathy. Gastroenterology, 2015, 148, S-57.	1.3	0
22	EPMA position paper in cancer: current overview and future perspectives. EPMA Journal, 2015, 6, 9.	6.1	86
23	Preventive and Predictive Genetics: Towards Personalised Medicine. Advances in Predictive, Preventive and Personalised Medicine, 2015, , .	0.6	6
24	Implementation of Genomic Medicine: Tools and Challenges. Advances in Predictive, Preventive and Personalised Medicine, 2015, , 329-347.	0.6	0
25	Abstract B22: Identification of novel drug combinations to target molecular pathways involved in breast cancer. , 2015, , .		0
26	Deregulation of the phosphatase, PP2A is a common event in breast cancer, predicting sensitivity to FTY720. EPMA Journal, 2014, 5, 3.	6.1	39
27	Expression of different functional isoforms in haematopoiesis. International Journal of Hematology, 2014, 99, 4-11.	1.6	7
28	Grsf1-Induced Translation of the SNARE Protein Use1 Is Required for Expansion of the Erythroid Compartment. PLoS ONE, 2014, 9, e104631.	2.5	22
29	PWE-107â€Osteoporosis in Crohn'S Disease Patients Expressing an Autophagy-Related Atg16L1 Gene Variant: Abstract PWE-107 Table. Gut, 2013, 62, A174.1-A174.	12.1	0
30	Conference Scene: Golden Helix Pharmacogenomics Days: educational activities on pharmacogenomics and personalized medicine. Pharmacogenomics, 2012, 13, 525-528.	1.3	7
31	The Role of Translation Initiation Regulation in Haematopoiesis. Comparative and Functional Genomics, 2012, 2012, 1-10.	2.0	10
32	Does Quantitative Heterogeneity of Human Fetal Hemoglobin (Hb F) Reveal Friends or Foes of KLF1 in Globin Gene Switching ?. Blood, 2011, 118, 1092-1092.	1.4	1
33	Haploinsufficiency for the erythroid transcription factor KLF1 causes hereditary persistence of fetal hemoglobin. Nature Genetics, 2010, 42, 801-805.	21.4	323
34	Tissue-specific splicing factor gene expression signatures. Nucleic Acids Research, 2008, 36, 4823-4832.	14.5	172
35	lgbp1 is part of a positive feedback loop in stem cell factor–dependent, selective mRNA translation initiation inhibiting erythroid differentiation. Blood, 2008, 112, 2750-2760.	1.4	36
36	Translation Initiation Factor 4E Inhibits Differentiation of Erythroid Progenitors. Molecular and Cellular Biology, 2005, 25, 8496-8506.	2.3	42

#	Article	IF	CITATIONS
37	Translation of IGBP1 mRNA Contributes to the Regulation of Expansion and Differentiation of Erythroid Progenitors Blood, 2005, 106, 72-72.	1.4	0
38	The Regulation of mRNA Translation Controls the Balance between Expansion and Differentiation of Erythroid Progenitors Blood, 2004, 104, 462-462.	1.4	4
39	Genetic and Serological Markers in Identifying Unclassified Colitis. , 0, , .		0