Tim De Meyer

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8614910/publications.pdf

Version: 2024-02-01

107 papers 6,274 citations

39 h-index 74163 75 g-index

121 all docs

121 docs citations

times ranked

121

11469 citing authors

#	Article	IF	CITATIONS
1	Spatiotemporal expression profile of novel and known small RNAs throughout rice plant development focussing on seed tissues. BMC Genomics, 2022, 23, 44.	2.8	4
2	Dehydroascorbate induces plant resistance in rice against rootâ€knot nematode <i>Meloidogyne graminicola</i> . Molecular Plant Pathology, 2022, 23, 1303-1319.	4.2	13
3	Transcript―and annotationâ€guided genome assembly of the European starling. Molecular Ecology Resources, 2022, 22, 3141-3160.	4.8	9
4	Genomeâ€wide shifts in histone modifications at early stage of rice infection with <i>Meloidogyne graminicola</i> . Molecular Plant Pathology, 2021, 22, 440-455.	4.2	14
5	DNA Methylation Regulates Transcription Factor-Specific Neurodevelopmental but Not Sexually Dimorphic Gene Expression Dynamics in Zebra Finch Telencephalon. Frontiers in Cell and Developmental Biology, 2021, 9, 583555.	3.7	8
6	Identification of DNA methylation markers for early detection of CRC indicates a role for nervous system-related genes in CRC. Clinical Epigenetics, 2021, 13, 80.	4.1	22
7	A Hypomorphic Mutant of PHD Domain Protein Male Meiocytes Death 1. Genes, 2021, 12, 516.	2.4	4
8	Clinically relevant aberrant Filip1l DNA methylation detected in a murine model of cutaneous squamous cell carcinoma. EBioMedicine, 2021, 67, 103383.	6.1	4
9	The RNA Atlas expands the catalog of human non-coding RNAs. Nature Biotechnology, 2021, 39, 1453-1465.	17.5	75
10	Non-coding RNAs in the interaction between rice and Meloidogyne graminicola. BMC Genomics, 2021, 22, 560.	2.8	12
11	The genome of the extremophile Artemia provides insight into strategies to cope with extreme environments. BMC Genomics, 2021, 22, 635.	2.8	20
12	Muscle strength is a major determinant of the blood pressure response to isometric stress testing: the Asklepios population study. Journal of Hypertension, 2020, 38, 224-234.	0.5	4
13	Chorismate mutase and isochorismatase, two potential effectors of the migratory nematode <i>Hirschmanniella oryzae</i> , increase host susceptibility by manipulating secondary metabolite content of rice. Molecular Plant Pathology, 2020, 21, 1634-1646.	4.2	12
14	Molecular insights into the compatible and incompatible interactions between sugar beet and the beet cyst nematode. BMC Plant Biology, 2020, 20, 483.	3.6	21
15	Genomeâ€wide DNA hypomethylation shapes nematode patternâ€triggered immunity in plants. New Phytologist, 2020, 227, 545-558.	7.3	44
16	Underestimated effect of intragenic HIV-1 DNA methylation on viral transcription in infected individuals. Clinical Epigenetics, 2020, 12, 36.	4.1	13
17	Molecular correlates of hypothalamic development in songbird ontogeny in comparison with the telencephalon. FASEB Journal, 2020, 34, 4997-5015.	0.5	0
18	Selection of miRNA reference genes for plant defence studies in rice (Oryza sativa). Planta, 2019, 250, 2101-2110.	3.2	9

#	Article	IF	CITATIONS
19	MEXPRESS update 2019. Nucleic Acids Research, 2019, 47, W561-W565.	14.5	179
20	Targeted RNAâ€seq successfully identifies normal and pathogenic splicing events in breast/ovarian cancer susceptibility and Lynch syndrome genes. International Journal of Cancer, 2019, 145, 401-414.	5.1	27
21	Letter by De Meyer Regarding Article, "Short Leukocyte Telomere Length Precedes Clinical Expression of Atherosclerosis: The Blood-and-Muscle Model― Circulation Research, 2018, 122, e71-e72.	4.5	2
22	Analysis of DNA methylation in cancer: location revisited. Nature Reviews Clinical Oncology, 2018, 15, 459-466.	27.6	486
23	Leukocyte telomere length and diet in the apparently healthy, middle-aged Asklepios population. Scientific Reports, 2018, 8, 6540.	3.3	22
24	A comprehensive overview of genomic imprinting in breast and its deregulation in cancer. Nature Communications, 2018, 9, 4120.	12.8	47
25	Genome-wide analyses identify a role for SLC17A4 and AADAT in thyroid hormone regulation. Nature Communications, 2018, 9, 4455.	12.8	181
26	Exploratory analysis of the human breast DNA methylation profile upon soymilk exposure. Scientific Reports, 2018, 8, 13617.	3.3	3
27	Body mass index is negatively associated with telomere length: a collaborative cross-sectional meta-analysis of 87 observational studies. American Journal of Clinical Nutrition, 2018, 108, 453-475.	4.7	137
28	Telomere Length as CardiovascularÂAgingÂBiomarker. Journal of the American College of Cardiology, 2018, 72, 805-813.	2.8	105
29	Reversal of Agingâ€Induced Increases in Aortic Stiffness by Targeting Cytoskeletal Proteinâ€Protein Interfaces. Journal of the American Heart Association, 2018, 7, .	3.7	17
30	Epigenetic sampling effects: nephrectomy modifies the clear cell renal cell cancer methylome. Cellular Oncology (Dordrecht), 2017, 40, 293-297.	4.4	2
31	Below-Ground Attack by the Root Knot Nematode <i>>Meloidogyne graminicola</i> > Predisposes Rice to Blast Disease. Molecular Plant-Microbe Interactions, 2017, 30, 255-266.	2.6	28
32	Telomeres and Atherosclerosis. Hypertension, 2017, 70, 243-244.	2.7	2
33	Methylome analysis of extreme chemoresponsive patients identifies novel markers of platinum sensitivity in high-grade serous ovarian cancer. BMC Medicine, 2017, 15, 116.	5.5	44
34	A Four-Gene Promoter Methylation Marker Panel Consisting of <i>GREM1, NEURL, LAD1,</i> and <i>NEFH</i> Predicts Survival of Clear Cell Renal Cell Cancer Patients. Clinical Cancer Research, 2017, 23, 2006-2018.	7.0	51
35	Microbial Community Dynamics and Response to Plant Growth-Promoting Microorganisms in the Rhizosphere of Four Common Food Crops Cultivated in Hydroponics. Microbial Ecology, 2017, 73, 378-393.	2.8	43
36	Decoy receptor 1 (DCR1) promoter hypermethylation and response to irinotecan in metastatic colorectal cancer. Oncotarget, 2017, 8, 63140-63154.	1.8	19

#	Article	lF	Citations
37	RAB25 expression is epigenetically downregulated in oral and oropharyngeal squamous cell carcinoma with lymph node metastasis. Epigenetics, 2016 , 11 , $653-663$.	2.7	18
38	A genome-wide search for epigenetically regulated genes in zebra finch using MethylCap-seq and RNA-seq. Scientific Reports, 2016, 6, 20957.	3.3	9
39	Molecular and epigenetic features of melanomas and tumor immune microenvironment linked to durable remission to ipilimumab-based immunotherapy in metastatic patients. Journal of Translational Medicine, 2016, 14, 232.	4.4	27
40	Telomeres and Atherosclerosis. Journal of the American College of Cardiology, 2016, 67, 2477-2479.	2.8	8
41	Discovery of new methylation markers to improve screening for cervical intraepithelial neoplasia grade 2/3. Clinical Epigenetics, 2016, 8, 29.	4.1	53
42	Identification and validation of <scp><i>WISP</i></scp> <i>111<i>1<i>5<i>4metastasis in oral squamous cell carcinoma. Genes Chromosomes and Cancer, 2016, 55, 45-59.</i></i></i></i>	2.8	28
43	Identification of long non-coding RNAs involved in neuronal development and intellectual disability. Scientific Reports, 2016, 6, 28396.	3.3	41
44	Locally advanced basal cell carcinoma has a distinct methylation and transcriptomic profile. Experimental Dermatology, 2016, 25, 316-318.	2.9	4
45	Genome-wide methylome analysis using MethylCap-seq uncovers 4 hypermethylated markers with high sensitivity for both adeno- and squamous-cell cervical carcinoma. Oncotarget, 2016, 7, 80735-80750.	1.8	15
46	Dynamic epigenetic changes to <i>VHL</i> occur with sunitinib in metastatic clear cell renal cancer. Oncotarget, 2016, 7, 25241-25250.	1.8	14
47	Decreasing initial telomere length in humans intergenerationally understates ageâ€associated telomere shortening. Aging Cell, 2015, 14, 669-677.	6.7	24
48	Genome-wide DNA methylation detection by MethylCap-seq and Infinium HumanMethylation450 BeadChips: an independent large-scale comparison. Scientific Reports, 2015, 5, 15375.	3.3	17
49	Reproducibility of telomere length assessment: Authors' Response to Damjan Krstajic and Ljubomir Buturovic. International Journal of Epidemiology, 2015, 44, 1739-1741.	1.9	8
50	MEXPRESS: visualizing expression, DNA methylation and clinical TCGA data. BMC Genomics, 2015, 16, 636.	2.8	257
51	Mining for viral fragments in methylation enriched sequencing data. Frontiers in Genetics, 2015, 6, 16.	2.3	5
52	PROTEOFORMER: deep proteome coverage through ribosome profiling and MS integration. Nucleic Acids Research, 2015, 43, e29-e29.	14.5	132
53	Is Southern blotting necessary to measure telomere length reproducibly? Authors' Response to: Commentary: The reliability of telomere length measurements. International Journal of Epidemiology, 2015, 44, 1686-1687.	1.9	8
54	Possible technical and biological explanations for the †parental telomere length inheritance discrepancy' enigma. European Journal of Human Genetics, 2015, 23, 3-4.	2.8	6

#	Article	IF	CITATIONS
55	Reproducibility of telomere length assessment: an international collaborative study. International Journal of Epidemiology, 2015, 44, 1673-1683.	1.9	133
56	Effect of sunitinib treatment on mutations and methylation in metastatic renal cancer Journal of Clinical Oncology, 2015, 33, 492-492.	1.6	18
57	Systemic Suppression of the Shoot Metabolism upon Rice Root Nematode Infection. PLoS ONE, 2014, 9, e106858.	2.5	13
58	ViVar: A Comprehensive Platform for the Analysis and Visualization of Structural Genomic Variation. PLoS ONE, 2014, 9, e113800.	2.5	45
59	SNP-guided identification of monoallelic DNA-methylation events from enrichment-based sequencing data. Nucleic Acids Research, 2014, 42, e157-e157.	14.5	6
60	Identification of Novel Genetic Loci Associated with Thyroid Peroxidase Antibodies and Clinical Thyroid Disease. PLoS Genetics, 2014, 10, e1004123.	3 . 5	150
61	Bacterial Diversity Assessment in Antarctic Terrestrial and Aquatic Microbial Mats: A Comparison between Bidirectional Pyrosequencing and Cultivation. PLoS ONE, 2014, 9, e97564.	2.5	60
62	Gender and telomere length: Systematic review and meta-analysis. Experimental Gerontology, 2014, 51, 15-27.	2.8	394
63	Rapid genetic adaptation precedes the spread of an exotic plant species. Molecular Ecology, 2014, 23, 2157-2164.	3.9	111
64	Arterial stiffness and influences of the metabolic syndrome: A cross-countries study. Atherosclerosis, 2014, 233, 654-660.	0.8	116
65	Nextâ€generation technologies and data analytical approaches for epigenomics. Environmental and Molecular Mutagenesis, 2014, 55, 155-170.	2.2	55
66	A non-genetic, epigenetic-like mechanism of telomere length inheritance?. European Journal of Human Genetics, 2014, 22, 10-11.	2.8	27
67	On Cross-Sectional Associations of Leukocyte Telomere Length with Cardiac Systolic, Diastolic and Vascular Function: The Asklepios Study. PLoS ONE, 2014, 9, e115071.	2.5	19
68	The CpG Island Methylator Phenotype: What's in a Name?. Cancer Research, 2013, 73, 5858-5868.	0.9	154
69	The impact of extensive clonal growth on fine-scale mating patterns: a full paternity analysis of a lily-of-the-valley population (Convallaria majalis). Annals of Botany, 2013, 111, 623-628.	2.9	11
70	Staphylococcal enterotoxin B influences the DNA methylation pattern in nasal polyp tissue: a preliminary study. Allergy, Asthma and Clinical Immunology, 2013, 9, 48.	2.0	13
71	Associations of rs4704397 in Phosphodiesterase 8B with Thyrotropin and Thyroid Hormone Concentrations. Thyroid, 2013, 23, 376-377.	4.5	2
72	The Effect of VEGF-Targeted Therapy on Biomarker Expression in Sequential Tissue from Patients with Metastatic Clear Cell Renal Cancer. Clinical Cancer Research, 2013, 19, 6924-6934.	7.0	62

#	Article	IF	CITATIONS
73	Associations between single nucleotide polymorphisms in thyroid hormone transporter genes (MCT8,) Tj ETQq1 1	0.784314	l rgBT /Ove
74	Family History of Cardiovascular Disease and Offspring Echocardiographic Left Ventricular Structure and Function: The Asklepios Study. Journal of the American Society of Echocardiography, 2013, 26, 1290-1297.e2.	2.8	4
75	SNP discovery using Pairedâ€End RAD â€ŧag sequencing on pooled genomic DNA of Sisymbrium austriacum (Brassicaceae). Molecular Ecology Resources, 2013, 13, 269-275.	4.8	24
76	Transcriptional analysis through RNA sequencing of giant cells induced by Meloidogyne graminicola in rice roots. Journal of Experimental Botany, 2013, 64, 3885-3898.	4.8	128
77	Quality Evaluation of Methyl Binding Domain Based Kits for Enrichment DNA-Methylation Sequencing. PLoS ONE, 2013, 8, e59068.	2.5	50
78	Addition of a Novel, Protective Family History Category Allows Better Profiling of Cardiovascular Risk and Atherosclerotic Burden in the General Population. The Asklepios Study. PLoS ONE, 2013, 8, e63185.	2.5	7
79	Coronary spasm after the topical use of cocaine in nasal surgery. American Journal of Case Reports, 2013, 14, 76-79.	0.8	9
80	Alterations of immune response of non-small cell lung cancer with Azacytidine. Oncotarget, 2013, 4, 2067-2079.	1.8	336
81	Abstract 3458: An epigenomic next-generation sequencing approach to identify predictive markers for PARP inhibitor response in breast cancer cells , 2013, , .		O
82	Transcriptome analysis of rice mature root tissue and root tips in early development by massive parallel sequencing. Journal of Experimental Botany, 2012, 63, 2141-2157.	4.8	41
83	No Shorter Telomeres in Subjects With a Family History of Cardiovascular Disease in the Asklepios Study. Arteriosclerosis, Thrombosis, and Vascular Biology, 2012, 32, 3076-3081.	2.4	16
84	Common Genetic Variation in the 3′- <i>BCL11B</i> Gene Desert Is Associated With Carotid-Femoral Pulse Wave Velocity and Excess Cardiovascular Disease Risk. Circulation: Cardiovascular Genetics, 2012, 5, 81-90.	5.1	90
85	Transcriptional reprogramming by root knot and migratory nematode infection in rice. New Phytologist, 2012, 196, 887-900.	7.3	157
86	Screening of soy and milk protein hydrolysates for their ability to activate the CCK1 receptor. Peptides, 2012, 34, 226-231.	2.4	13
87	Low dose irradiation of thyroid cells reveals a unique transcriptomic and epigenetic signature in RET/PTC-positive cells. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2012, 731, 27-40.	1.0	19
88	Telomere length and cardiovascular aging: The means to the ends?. Ageing Research Reviews, 2011, 10, 297-303.	10.9	68
89	Telomere Length Integrates Psychological Factors in the Successful Aging Story, But What About the Biology?. Psychosomatic Medicine, 2011, 73, 524-527.	2.0	10
90	Response to low-dose X-irradiation is p53-dependent in a papillary thyroid carcinoma model system. International Journal of Oncology, 2011, 39, 1429-41.	3.3	2

#	Article	IF	Citations
91	A variant at chromosome 9p21 is associated with recurrent myocardial infarction and cardiac death after acute coronary syndrome: The GRACE Genetics Study. European Heart Journal, 2010, 31, 1132-1141.	2.2	50
92	Evaluation of standard and advanced preprocessing methods for the univariate analysis of blood serum 1H-NMR spectra. Analytical and Bioanalytical Chemistry, 2010, 398, 1781-1790.	3.7	40
93	Patterns of sex ratio variation and genetic diversity in the dioecious forest perennial Mercurialis perennis. Plant Ecology, 2010, 206, 105-114.	1.6	39
94	Systemic telomere length and preclinical atherosclerosis: the Asklepios Study. European Heart Journal, 2009, 30, 3074-3081.	2.2	67
95	<i>E2F</i> s mediate a fundamental cellâ€cycle deregulation in highâ€grade serous ovarian carcinomas. Journal of Pathology, 2009, 217, 14-20.	4.5	35
96	High Content Analysis of Human Fibroblast Cell Cultures after Exposure to Space Radiation. Radiation Research, 2009, 172, 423-436.	1.5	19
97	Telomere biology in giant cell tumour of bone. Journal of Pathology, 2008, 214, 555-563.	4.5	33
98	NMR-Based Characterization of Metabolic Alterations in Hypertension Using an Adaptive, Intelligent Binning Algorithm. Analytical Chemistry, 2008, 80, 3783-3790.	6.5	217
99	Lower red blood cell counts in middleâ€aged subjects with shorter peripheral blood leukocyte telomere length. Aging Cell, 2008, 7, 700-705.	6.7	23
100	lonizing radiation-induced gene modulations, cytokine content changes and telomere shortening in mouse fetuses exhibiting forelimb defects. Developmental Biology, 2008, 322, 302-313.	2.0	21
101	Studying telomeres in a longitudinal population based study. Frontiers in Bioscience - Landmark, 2008, 13, 2960.	3.0	38
102	PubMeth: a cancer methylation database combining text-mining and expert annotation. Nucleic Acids Research, 2007, 36, D842-D846.	14.5	144
103	Paternal age at birth is an important determinant of offspring telomere length. Human Molecular Genetics, 2007, 16, 3097-3102.	2.9	146
104	Telomere length and cardiovascular risk factors in a middleâ€aged population free of overt cardiovascular disease. Aging Cell, 2007, 6, 639-647.	6.7	309
105	Telomere length versus hormonal and bone mineral status in healthy elderly men. Mechanisms of Ageing and Development, 2005, 126, 1115-1122.	4.6	75
106	Telomere shortening is associated with malformation in p53-deficient mice after irradiation during specific stages of development. DNA Repair, 2005, 4, 1028-1037.	2.8	14
107	Telomere attrition as ageing biomarker. Anticancer Research, 2005, 25, 3011-21.	1.1	111