## Giovanni Nigita

List of Publications by Year in descending order

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Version: 2024-02-01

279798 289244 1,685 54 23 40 citations h-index g-index papers 60 60 60 2762 docs citations times ranked citing authors all docs

| #  | Article   | IF          | Citations |
|----|---|-------------|-----------|
| 1  | LEDGF/p75-mediated chemoresistance of mixed-lineage leukemia involves cell survival pathways and super enhancer activators. Cancer Gene Therapy, 2022, 29, 133-140.   | 4.6         | 7         |
| 2  | Synergistic apoptotic effect of miR-183-5p and Polo-Like kinase 1 inhibitor NMS-P937 in breast cancer cells. Cell Death and Differentiation, 2022, 29, 407-419.   | 11.2        | 5         |
| 3  | A large fraction of trisomy 12, 17p $<$ sup $>$ â°' $<$ sup $>$ , and 11q $<$ sup $>$ â°' $<$ sup $>$ CLL cases carry unidentified microdeletions of $<$ i> $>$ miR-15a/16-1 $<$  i $>$ . Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, . | 7.1         | 3         |
| 4  | Disparities in Lung Cancer: miRNA Isoform Characterization in Lung Adenocarcinoma. Cancers, 2022, 14, 773.  | 3.7         | 4         |
| 5  | Loss of expression of both miR-15/16 loci in CML transition to blast crisis. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .  | 7.1         | 6         |
| 6  | Detecting and Characterizing A-To-I microRNA Editing in Cancer. Cancers, 2021, 13, 1699.  | 3.7         | 17        |
| 7  | The MicroRNA Family Gets Wider: The IsomiRs Classification and Role. Frontiers in Cell and Developmental Biology, 2021, 9, 668648.  | 3.7         | 52        |
| 8  | Abstract 474: Extracellular vesicle - MDM2 as liquid biopsy biomarker for disease identification in retroperitoneal liposarcoma. , 2021, , .  |             | 0         |
| 9  | MiREDiBase, a manually curated database of validated and putative editing events in microRNAs.<br>Scientific Data, 2021, 8, 199.  | <b>5.</b> 3 | 18        |
| 10 | Non-Coding RNA Editing in Cancer Pathogenesis. Cancers, 2020, 12, 1845.   | 3.7         | 16        |
| 11 | MicroRNAs in Skeletal Muscle and Hints on Their Potential Role in Muscle Wasting During Cancer<br>Cachexia. Frontiers in Oncology, 2020, 10, 607196.  | 2.8         | 15        |
| 12 | MiR-124a Regulates Extracellular Vesicle Release by Targeting GTPase Rabs in Lung Cancer. Frontiers in Oncology, 2020, 10, 1454.  | 2.8         | 8         |
| 13 | Combined loss of function of two different loci of miR-15/16 drives the pathogenesis of acute myeloid leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 12332-12340.  | 7.1         | 28        |
| 14 | Exosomal miRNA signatures of pancreatic lesions. BMC Gastroenterology, 2020, 20, 137.   | 2.0         | 25        |
| 15 | Pleiotropic tumor suppressor functions of WWOX antagonize metastasis. Signal Transduction and Targeted Therapy, 2020, 5, 43.  | 17.1        | 27        |
| 16 | Abstract 2543: Concurrent profiling of canonical and modified miRNAomes from TCGA and TARGET cohorts leads to enhanced resolution in cancer., 2020,,.   |             | 0         |
| 17 | isoTar: Consensus Target Prediction with Enrichment Analysis for MicroRNAs Harboring Editing Sites and Other Variations. Methods in Molecular Biology, 2019, 1970, 211-235.   | 0.9         | 13        |
| 18 | Investigating miRNA–IncRNA Interactions: Computational Tools and Resources. Methods in Molecular Biology, 2019, 1970, 251-277.  | 0.9         | 22        |

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|----|---|------|-----------|
| 19 | Identification of tRNA-derived ncRNAs in TCGA and NCI-60 panel cell lines and development of the public database tRFexplorer. Database: the Journal of Biological Databases and Curation, 2019, 2019, .           | 3.0  | 36        |
| 20 | WWOX Inhibits Metastasis of Triple-Negative Breast Cancer Cells via Modulation of miRNAs. Cancer Research, 2019, 79, 1784-1798.   | 0.9  | 30        |
| 21 | ncRNA Editing: Functional Characterization and Computational Resources. Methods in Molecular<br>Biology, 2019, 1912, 133-174.   | 0.9  | 20        |
| 22 | Reprogramming miRNAs global expression orchestrates development of drug resistance in BRAF mutated melanoma. Cell Death and Differentiation, 2019, 26, 1267-1282.   | 11.2 | 47        |
| 23 | Circulating Micrornas Predict Survival of Patients with Tumors of Glial Origin. EBioMedicine, 2018, 30, 105-112.  | 6.1  | 27        |
| 24 | miRandola 2017: a curated knowledge base of non-invasive biomarkers. Nucleic Acids Research, 2018, 46, D354-D359.   | 14.5 | 61        |
| 25 | Editorial: Epitranscriptomics: The Novel RNA Frontier. Frontiers in Bioengineering and Biotechnology, 2018, 6, 191.   | 4.1  | 6         |
| 26 | Knockout of both miR-15/16 loci induces acute myeloid leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 13069-13074.  | 7.1  | 39        |
| 27 | miR-125a and miR-34a expression predicts Richter syndrome in chronic lymphocytic leukemia patients.<br>Blood, 2018, 132, 2179-2182.   | 1.4  | 25        |
| 28 | Circulating miR-106b-3p, miR-101-3p and miR-1246 as diagnostic biomarkers of hepatocellular carcinoma. Oncotarget, 2018, 9, 15350-15364.  | 1.8  | 79        |
| 29 | The TLR7/8/9 Antagonist IMO-8503 Inhibits Cancer-Induced Cachexia. Cancer Research, 2018, 78, 6680-6690.  | 0.9  | 33        |
| 30 | Tissue and exosomal miRNA editing in Non-Small Cell Lung Cancer. Scientific Reports, 2018, 8, 10222.  | 3.3  | 38        |
| 31 | RNA Methylation in ncRNA: Classes, Detection, and Molecular Associations. Frontiers in Genetics, 2018, 9, 243.  | 2.3  | 40        |
| 32 | Abstract 473: miR-135b mediates gemcitabine sensitivity in breast cancer cells by modulating epithelial-to-mesenchymal transition and mTOR-signaling. , 2018, , .   |      | 0         |
| 33 | Prognostic and Biologic Significance of Transfer RNA-Derived Small RNAs (tsRNAs) Expression in Younger Adult Patients (Pts) with Cytogenetically Normal Acute Myeloid Leukemia (CN-AML). Blood, 2018, 132, 89-89. | 1.4  | 9         |
| 34 | Immunotherapy Bridge 2016 and Melanoma Bridge 2016: meeting abstracts. Journal of Translational Medicine, 2017, 15, .   | 4.4  | 1         |
| 35 | Selective targeting of point-mutated KRAS through artificial microRNAs. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E4203-E4212.                                  | 7.1  | 38        |
| 36 | tsRNA signatures in cancer. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 8071-8076.  | 7.1  | 202       |

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|----|---|------|-----------|
| 37 | Transcriptomic analysis of collecting duct carcinoma of the kidney. Annals of Oncology, 2016, 27, vi274.  | 1.2  | O         |
| 38 | Dysregulation of a family of short noncoding RNAs, tsRNAs, in human cancer. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 5071-5076.        | 7.1  | 183       |
| 39 | Noncoding RNA: Current Deep Sequencing Data Analysis Approaches and Challenges. Human Mutation, 2016, 37, 1283-1298.  | 2.5  | 74        |
| 40 | microRNA editing in seed region aligns with cellular changes in hypoxic conditions. Nucleic Acids Research, 2016, 44, 6298-6308.  | 14.5 | 41        |
| 41 | MicroRNA fingerprints in juvenile myelomonocytic leukemia (JMML) identified miR-150-5p as a tumor suppressor and potential target for treatment. Oncotarget, 2016, 7, 55395-55408.        | 1.8  | 30        |
| 42 | MAPK15 upregulation promotes cell proliferation and prevents DNA damage in male germ cell tumors. Oncotarget, 2016, 7, 20981-20998.   | 1.8  | 37        |
| 43 | Gene-expression profiling of collecting duct carcinoma of the kidney Journal of Clinical Oncology, 2016, 34, 540-540.   | 1.6  | 0         |
| 44 | Abstract LB-166: miRNA editing in seed region is in synergy with cellular changes in hypoxic conditions. , $2016,$  |      | 0         |
| 45 | Role of Ts-RNAs in CLL. Blood, 2016, 128, 2016-2016.  | 1.4  | 0         |
| 46 | Knowledge in the Investigation of A-to-I RNA Editing Signals. Frontiers in Bioengineering and Biotechnology, 2015, 3, 18.   | 4.1  | 17        |
| 47 | A-to-I RNA Editing: Current Knowledge Sources and Computational Approaches with Special Emphasis on Non-Coding RNA Molecules. Frontiers in Bioengineering and Biotechnology, 2015, 3, 37. | 4.1  | 47        |
| 48 | Computational Approaches for the Analysis of ncRNA through Deep Sequencing Techniques. Frontiers in Bioengineering and Biotechnology, 2015, 3, 77.  | 4.1  | 66        |
| 49 | Microrna-150 Regulates STAT5b Levels in Juvenile Myelomonocytic Leukemia (JMML). Blood, 2015, 126, 2851-2851.   | 1.4  | 1         |
| 50 | Abstract C17: Role of miR-135b in gemcitabine sensitivity for metastatic breast cancer patients. , 2015, , .  |      | 0         |
| 51 | VIRGO: visualization of A-to-I RNA editing sites in genomic sequences. BMC Bioinformatics, 2013, 14, S5.  | 2.6  | 10        |
| 52 | miR-EdiTar: a database of predicted A-to-I edited miRNA target sites. Bioinformatics, 2012, 28, 3166-3168.  | 4.1  | 28        |
| 53 | miRandola: Extracellular Circulating MicroRNAs Database. PLoS ONE, 2012, 7, e47786.   | 2.5  | 142       |
| 54 | An integrated system for mining relations among microRNAs, drugs and phenotypes. EMBnet Journal, 2012, 18, 75.  | 0.6  | 0         |