

# Dimitrios Spentzos

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1655683/publications.pdf>

Version: 2024-02-01

13  
papers

769  
citations

933447

10  
h-index

1199594

12  
g-index

17  
all docs

17  
docs citations

17  
times ranked

1140  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Gene Expression Signature With Independent Prognostic Significance in Epithelial Ovarian Cancer. <i>Journal of Clinical Oncology</i> , 2004, 22, 4700-4710.  | 1.6  | 236       |
| 2  | Unique Gene Expression Profile Based on Pathologic Response in Epithelial Ovarian Cancer. <i>Journal of Clinical Oncology</i> , 2005, 23, 7911-7918.   | 1.6  | 133       |
| 3  | MicroRNA paraffin-based studies in osteosarcoma reveal reproducible independent prognostic profiles at 14q32. <i>Genome Medicine</i> , 2013, 5, 2.   | 8.2  | 96        |
| 4  | A network model for angiogenesis in ovarian cancer. <i>BMC Bioinformatics</i> , 2015, 16, 115.   | 2.6  | 60        |
| 5  | IGF axis gene expression patterns are prognostic of survival in epithelial ovarian cancer. <i>Endocrine-Related Cancer</i> , 2007, 14, 781-790.  | 3.1  | 55        |
| 6  | An imprinted non-coding genomic cluster at 14q32 defines clinically relevant molecular subtypes in osteosarcoma across multiple independent datasets. <i>Journal of Hematology and Oncology</i> , 2017, 10, 107.       | 17.0 | 38        |
| 7  | MicroRNA-155 expression is independently predictive of outcome in chordoma. <i>Oncotarget</i> , 2015, 6, 9125-9139.  | 1.8  | 38        |
| 8  | Clinical and Molecular Analysis of Pathologic Fracture-associated Osteosarcoma: MicroRNA profile Is Different and Correlates with Prognosis. <i>Clinical Orthopaedics and Related Research</i> , 2019, 477, 2114-2126. | 1.5  | 19        |
| 9  | EWSR1-ATF1 dependent 3D connectivity regulates oncogenic and differentiation programs in Clear Cell Sarcoma. <i>Nature Communications</i> , 2022, 13, 2267.  | 12.8 | 18        |
| 10 | Genome-wide DNA methylation patterns reveal clinically relevant predictive and prognostic subtypes in human osteosarcoma. <i>Communications Biology</i> , 2022, 5, 213.  | 4.4  | 10        |
| 11 | MicroRNA-mRNA networks define translatable molecular outcome phenotypes in osteosarcoma. <i>Scientific Reports</i> , 2020, 10, 4409.   | 3.3  | 9         |
| 12 | A Collision Tumor Involving a Primary Leiomyosarcoma of the Lower Extremity and a Metastatic Medullary Thyroid Carcinoma. <i>JBJS Case Connector</i> , 2017, 7, e90-e90.   | 0.3  | 1         |
| 13 | Age-Related Transcription Levels of KU70 and BIK in CD34+ Hematopoietic Stem and Progenitor Cells. <i>Blood</i> , 2005, 106, 4205-4205.  | 1.4  | 0         |