

# Elena S Martens-Uzunova

## List of Publications by Year in descending order

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Version: 2024-02-01

26  
papers

9,208  
citations

430442

18  
h-index

580395

25  
g-index

26  
all docs

26  
docs citations

26  
times ranked

15214  
citing authors

#	ARTICLE	IF	CITATIONS
1	Extracellular vesicles as a source of prostate cancer biomarkers in liquid biopsies: a decade of research. <i>British Journal of Cancer</i> , 2022, 126, 331-350.	2.9	39
2	Extracellular Vesicles as Novel Players in Kidney Disease. <i>Journal of the American Society of Nephrology: JASN</i> , 2022, 33, 467-471.	3.0	6
3	CRISPRs in the human genome are differentially expressed between malignant and normal adjacent to tumor tissue. <i>Communications Biology</i> , 2022, 5, 338.	2.0	2
4	Urinary extracellular vesicles: A position paper by the Urine Task Force of the International Society for Extracellular Vesicles. <i>Journal of Extracellular Vesicles</i> , 2021, 10, e12093.	5.5	182
5	Androgens alter the heterogeneity of small extracellular vesicles and the small RNA cargo in prostate cancer. <i>Journal of Extracellular Vesicles</i> , 2021, 10, e12136.	5.5	15
6	Urinary Extracellular Vesicles in Urology: Current Successes and Challenges Ahead. <i>European Urology</i> , 2021, 81, 127-127.	0.9	0
7	The role of OncoSnoRNAs and Ribosomal RNA 2â€™-O-methylation in Cancer. <i>RNA Biology</i> , 2021, 18, 61-74.	1.5	21
8	VIRMA-Dependent N6-Methyladenosine Modifications Regulate the Expression of Long Non-Coding RNAs CCAT1 and CCAT2 in Prostate Cancer. <i>Cancers</i> , 2020, 12, 771.	1.7	59
9	Minimal information for studies of extracellular vesicles 2018 (MISEV2018): a position statement of the International Society for Extracellular Vesicles and update of the MISEV2014 guidelines. <i>Journal of Extracellular Vesicles</i> , 2018, 7, 1535750.	5.5	6,961
10	Summary of the ISEV workshop on extracellular vesicles as disease biomarkers, held in Birmingham, UK, during December 2017. <i>Journal of Extracellular Vesicles</i> , 2018, 7, 1473707.	5.5	60
11	The Non-Coding Transcriptome of Prostate Cancer: Implications for Clinical Practice. <i>Molecular Diagnosis and Therapy</i> , 2017, 21, 385-400.	1.6	18
12	Systematic Identification of MicroRNAs That Impact on Proliferation of Prostate Cancer Cells and Display Changed Expression in Tumor Tissue. <i>European Urology</i> , 2016, 69, 1120-1128.	0.9	53
13	A comprehensive repertoire of tRNA-derived fragments in prostate cancer. <i>Oncotarget</i> , 2016, 7, 24766-24777.	0.8	144
14	Identification and Diagnostic Performance of a Small RNA within the PCA3 and BMCC1 Gene Locus That Potentially Targets mRNA. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 268-275.	1.1	10
15	FlaiMapper: computational annotation of small ncRNA-derived fragments using RNA-seq high-throughput data. <i>Bioinformatics</i> , 2015, 31, 665-673.	1.8	28
16	C/D-box snoRNA-derived RNA production is associated with malignant transformation and metastatic progression in prostate cancer. <i>Oncotarget</i> , 2015, 6, 17430-17444.	0.8	80
17	Long Noncoding RNA in Prostate, Bladder, and Kidney Cancer. <i>European Urology</i> , 2014, 65, 1140-1151.	0.9	601
18	miQâ€™A novel microRNA based diagnostic and prognostic tool for prostate cancer. <i>International Journal of Cancer</i> , 2013, 132, 2867-2875.	2.3	79

#	ARTICLE	IF	CITATIONS
19	Beyond microRNA " Novel RNAs derived from small non-coding RNA and their implication in cancer. <i>Cancer Letters</i> , 2013, 340, 201-211.	3.2	169
20	Diagnostic and prognostic signatures from the small non-coding RNA transcriptome in prostate cancer. <i>Oncogene</i> , 2012, 31, 978-991.	2.6	239
21	miR-183-96-182 Cluster Is Overexpressed in Prostate Tissue and Regulates Zinc Homeostasis in Prostate Cells. <i>Journal of Biological Chemistry</i> , 2011, 286, 44503-44511.	1.6	120
22	An inventory of the <i>Aspergillus niger</i> secretome by combining in silico predictions with shotgun proteomics data. <i>BMC Genomics</i> , 2010, 11, 584.	1.2	74
23	Assessment of the pectin degrading enzyme network of <i>Aspergillus niger</i> by functional genomics. <i>Fungal Genetics and Biology</i> , 2009, 46, S170-S179.	0.9	102
24	An evolutionary conserved d-galacturonic acid metabolic pathway operates across filamentous fungi capable of pectin degradation. <i>Fungal Genetics and Biology</i> , 2008, 45, 1449-1457.	0.9	74
25	Carboxy-Terminal Extension Effects on Crystal Formation and Insecticidal Properties of Colorado Potato Beetle-Active <i>Bacillus thuringiensis</i> $\delta$ -Endotoxins. <i>Molecular Biotechnology</i> , 2006, 32, 185-196.	1.3	10
26	A new group of exo-acting family 28 glycoside hydrolases of <i>Aspergillus niger</i> that are involved in pectin degradation. <i>Biochemical Journal</i> , 2006, 400, 43-52.	1.7	62