

Lea Landolt

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

205
citations

1163117

8
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

370
citing authors

#	ARTICLE	IF	CITATIONS
1	Fibrosis and cancer: shared features and mechanisms suggest common targeted therapeutic approaches. <i>Nephrology Dialysis Transplantation</i> , 2022, 37, 1024-1032.	0.7	18
2	Axl inhibitor bemcentinib alleviates mitochondrial dysfunction in the unilateral ureter obstruction murine model. <i>Journal of Cellular and Molecular Medicine</i> , 2021, 25, 7407-7417.	3.6	11
3	Ultrasound and Microbubbles Enhance Uptake of Doxorubicin in Murine Kidneys. <i>Pharmaceutics</i> , 2021, 13, 2038.	4.5	3
4	AXL targeting reduces fibrosis development in experimental unilateral ureteral obstruction. <i>Physiological Reports</i> , 2019, 7, e14091.	1.7	13
5	Transcriptome-proteome integration of archival human renal cell carcinoma biopsies enables identification of molecular mechanisms. <i>American Journal of Physiology - Renal Physiology</i> , 2019, 316, F1053-F1067.	2.7	15
6	Expanding the Utilization of Formalin-Fixed, Paraffin-Embedded Archives: Feasibility of miR-Seq for Disease Exploration and Biomarker Development from Biopsies with Clear Cell Renal Cell Carcinoma. <i>International Journal of Molecular Sciences</i> , 2018, 19, 803.	4.1	3
7	Clear Cell Renal Cell Carcinoma is linked to Epithelial-to-Mesenchymal Transition and to Fibrosis. <i>Physiological Reports</i> , 2017, 5, e13305.	1.7	36
8	Transcriptome Sequencing (RNAseq) Enables Utilization of Formalin-Fixed, Paraffin-Embedded Biopsies with Clear Cell Renal Cell Carcinoma for Exploration of Disease Biology and Biomarker Development. <i>PLoS ONE</i> , 2016, 11, e0149743.	2.5	50
9	RNA extraction for RNA sequencing of archival renal tissues. <i>Scandinavian Journal of Clinical and Laboratory Investigation</i> , 2016, 76, 426-434.	1.2	38
10	Development and confirmation of potential gene classifiers of human clear cell renal cell carcinoma using next-generation RNA sequencing. <i>Scandinavian Journal of Urology</i> , 2016, 50, 452-462.	1.0	18