

# Erin L Crawford

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4524943/publications.pdf>

Version: 2024-02-01

12  
papers

292  
citations

932766

10  
h-index

1199166

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

358  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cross-oncopanel study reveals high sensitivity and accuracy with overall analytical performance depending on genomic regions. <i>Genome Biology</i> , 2021, 22, 109.	3.8	20
2	Advancing NGS quality control to enable measurement of actionable mutations in circulating tumor DNA. <i>Cell Reports Methods</i> , 2021, 1, 100106.	1.4	9
3	Platelet-T cell aggregates in lung cancer patients: Implications for thrombosis. <i>PLoS ONE</i> , 2020, 15, e0236966.	1.1	16
4	Circulating Lactonase Activity but Not Protein Level of PON-1 Predicts Adverse Outcomes in Subjects with Chronic Kidney Disease. <i>Journal of Clinical Medicine</i> , 2019, 8, 1034.	1.0	16
5	Chronic Low Dose Oral Exposure to Microcystin-LR Exacerbates Hepatic Injury in a Murine Model of Non-Alcoholic Fatty Liver Disease. <i>Toxins</i> , 2019, 11, 486.	1.5	30
6	Technical advance in targeted NGS analysis enables identification of lung cancer risk-associated low frequency TP53, PIK3CA, and BRAF mutations in airway epithelial cells. <i>BMC Cancer</i> , 2019, 19, 1081.	1.1	12
7	Control for stochastic sampling variation and qualitative sequencing error in next generation sequencing. <i>Biomolecular Detection and Quantification</i> , 2015, 5, 30-37.	7.0	17
8	Bioinformatics analysis of plant orthologous introns: identification of an intronic tRNA-like sequence. <i>Gene</i> , 2014, 548, 81-90.	1.0	4
9	Targeted RNA-Sequencing with Competitive Multiplex-PCR Amplicon Libraries. <i>PLoS ONE</i> , 2013, 8, e79120.	1.1	30
10	Multiplex standardized RT-PCR for expression analysis of many genes in small samples. <i>Biochemical and Biophysical Research Communications</i> , 2002, 293, 509-516.	1.0	16
11	Expression Measurement of Many Genes Simultaneously by Quantitative RT-PCR Using Standardized Mixtures of Competitive Templates. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 1998, 19, 6-17.	1.4	87
12	The Gene Expression Index <i>c-myc</i> — E2F-1/p21 Is Highly Predictive of Malignant Phenotype in Human Bronchial Epithelial Cells. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 1998, 19, 18-24.	1.4	35