## Erin L Crawford

## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Expression Measurement of Many Genes Simultaneously by Quantitative RT-PCR Using Standardized Mixtures of Competitive Templates. American Journal of Respiratory Cell and Molecular Biology, 1998, 19, 6-17.	2.9	87
2	The Gene Expression Index c- <i>myc</i> $\tilde{A}$ — E2F-1/p21 Is Highly Predictive of Malignant Phenotype in Human Bronchial Epithelial Cells. American Journal of Respiratory Cell and Molecular Biology, 1998, 19, 18-24.	2.9	35
3	Chronic Low Dose Oral Exposure to Microcystin-LR Exacerbates Hepatic Injury in a Murine Model of Non-Alcoholic Fatty Liver Disease. Toxins, 2019, 11, 486.	3.4	30
4	Targeted RNA-Sequencing with Competitive Multiplex-PCR Amplicon Libraries. PLoS ONE, 2013, 8, e79120.	2.5	30
5	Cross-oncopanel study reveals high sensitivity and accuracy with overall analytical performance depending on genomic regions. Genome Biology, 2021, 22, 109.	8.8	20
6	Control for stochastic sampling variation and qualitative sequencing error in next generation sequencing. Biomolecular Detection and Quantification, 2015, 5, 30-37.	7.0	17
7	Multiplex standardized RT-PCR for expression analysis of many genes in small samples. Biochemical and Biophysical Research Communications, 2002, 293, 509-516.	2.1	16
8	Circulating Lactonase Activity but Not Protein Level of PON-1 Predicts Adverse Outcomes in Subjects with Chronic Kidney Disease. Journal of Clinical Medicine, 2019, 8, 1034.	2.4	16
9	Platelet-T cell aggregates in lung cancer patients: Implications for thrombosis. PLoS ONE, 2020, 15, e0236966.	2.5	16
10	Technical advance in targeted NGS analysis enables identification of lung cancer risk-associated low frequency TP53, PIK3CA, and BRAF mutations in airway epithelial cells. BMC Cancer, 2019, 19, 1081.	2.6	12
11	Advancing NGS quality control to enable measurement of actionable mutations in circulating tumor DNA. Cell Reports Methods, 2021, 1, 100106.	2.9	9
12	Bioinformatics analysis of plant orthologous introns: identification of an intronic tRNA-like sequence. Gene, 2014, 548, 81-90.	2.2	4