

# Amali Mallawaarachchi

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11800771/publications.pdf>

Version: 2024-02-01

9  
papers

232  
citations

1478505

6  
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1720034

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9  
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9  
docs citations

9  
times ranked

515  
citing authors

#	ARTICLE	IF	CITATIONS
1	Standardized practices for RNA diagnostics using clinically accessible specimens reclassifies 75% of putative splicing variants. <i>Genetics in Medicine</i> , 2022, 24, 130-145.	2.4	45
2	Attitudes and Practices of Australian Nephrologists Toward Implementation of Clinical Genomics. <i>Kidney International Reports</i> , 2021, 6, 272-283.	0.8	28
3	Australia and New Zealand renal gene panel testing in routine clinical practice of 542 families. <i>Npj Genomic Medicine</i> , 2021, 6, 20.	3.8	11
4	Comprehensive evaluation of a prospective Australian patient cohort with suspected genetic kidney disease undergoing clinical genomic testing: a study protocol. <i>BMJ Open</i> , 2019, 9, e029541.	1.9	6
5	Renal genetics in Australia: Kidney medicine in the genomic age. <i>Nephrology</i> , 2019, 24, 279-286.	1.6	18
6	Nemaline myopathy and distal arthrogryposis associated with an autosomal recessive <i>TNNT3</i> splice variant. <i>Human Mutation</i> , 2018, 39, 383-388.	2.5	48
7	Massively parallel sequencing and targeted exomes in familial kidney disease can diagnose underlying genetic disorders. <i>Kidney International</i> , 2017, 92, 1493-1506.	5.2	74
8	Novel complement factor H gene mutation causing atypical haemolytic uraemic syndrome: early Eculizumab prevents acute dialysis. <i>CKJ: Clinical Kidney Journal</i> , 2017, 10, 263-265.	2.9	0
9	The HIDDEN Protocol: An Australian Prospective Cohort Study to Determine the Utility of Whole Genome Sequencing in Kidney Failure of Unknown Aetiology. <i>Frontiers in Medicine</i> , 0, 9, .	2.6	2