

# Vilma-Lotta Lehtokari

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6355836/publications.pdf>

Version: 2024-02-01

22  
papers

1,296  
citations

567247

15  
h-index

677123

22  
g-index

24  
all docs

24  
docs citations

24  
times ranked

1146  
citing authors

#	ARTICLE	IF	CITATIONS
1	Nemaline Myopathy with Minicores Caused by Mutation of the CFL2 Gene Encoding the Skeletal Muscle Actin- $\alpha$ -Binding Protein, Cofilin-2. <i>American Journal of Human Genetics</i> , 2007, 80, 162-167.	6.2	213
2	Mutations in KLHL40 Are a Frequent Cause of Severe Autosomal-Recessive Nemaline Myopathy. <i>American Journal of Human Genetics</i> , 2013, 93, 6-18.	6.2	186
3	Distal myopathy caused by homozygous missense mutations in the nebulin gene. <i>Brain</i> , 2007, 130, 1465-1476.	7.6	130
4	Identification of 45 novel mutations in the nebulin gene associated with autosomal recessive nemaline myopathy. <i>Human Mutation</i> , 2006, 27, 946-956.	2.5	112
5	Mutation Update: The Spectra of Nebulin Variants and Associated Myopathies. <i>Human Mutation</i> , 2014, 35, 1418-1426.	2.5	107
6	Complete genomic structure of the human nebulin gene and identification of alternatively spliced transcripts. <i>European Journal of Human Genetics</i> , 2004, 12, 744-751.	2.8	101
7	Cap disease caused by heterozygous deletion of the $\beta$ -tropomyosin gene TPM2. <i>Neuromuscular Disorders</i> , 2007, 17, 433-442.	0.6	90
8	Nemaline myopathy caused by mutations in the nebulin gene may present as a distal myopathy. <i>Neuromuscular Disorders</i> , 2011, 21, 556-562.	0.6	56
9	The exon 55 deletion in the nebulin gene - One single founder mutation with world-wide occurrence. <i>Neuromuscular Disorders</i> , 2009, 19, 179-181.	0.6	54
10	Mutation-specific effects on thin filament length in thin filament myopathy. <i>Annals of Neurology</i> , 2016, 79, 959-969.	5.3	54
11	Identification of a founder mutation in TPM3 in nemaline myopathy patients of Turkish origin. <i>European Journal of Human Genetics</i> , 2008, 16, 1055-1061.	2.8	36
12	A recurrent copy number variation of the NEB triplicate region: only revealed by the targeted nemaline myopathy CGH array. <i>European Journal of Human Genetics</i> , 2016, 24, 574-580.	2.8	32
13	Dominantly inherited distal nemaline/cap myopathy caused by a large deletion in the nebulin gene. <i>Neuromuscular Disorders</i> , 2019, 29, 97-107.	0.6	31
14	An Extended Targeted Copy Number Variation Detection Array Including 187 Genes for the Diagnostics of Neuromuscular Disorders. <i>Journal of Neuromuscular Diseases</i> , 2018, 5, 307-314.	2.6	17
15	MoBiDiC Prioritization Algorithm, a Free, Accessible, and Efficient Pipeline for Single-Nucleotide Variant Annotation and Prioritization for Next-Generation Sequencing Routine Molecular Diagnosis. <i>Journal of Molecular Diagnostics</i> , 2018, 20, 465-473.	2.8	13
16	Bilateral foot-drop as predominant symptom in nebulin (NEB) gene related $\alpha$ -core-rod-congenital myopathy. <i>European Journal of Medical Genetics</i> , 2015, 58, 556-561.	1.3	12
17	A nebulin super-repeat panel reveals stronger actin binding toward the ends of the super-repeat region. <i>Muscle and Nerve</i> , 2019, 59, 116-121.	2.2	10
18	Congenital asymmetric distal myopathy with hemifacial weakness caused by a heterozygous large de novo mosaic deletion in nebulin. <i>Neuromuscular Disorders</i> , 2021, 31, 539-545.	0.6	9

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19	Clinically variable nemaline myopathy in a three-generation family caused by mutation of the skeletal muscle alpha-actin gene. <i>Neuromuscular Disorders</i> , 2018, 28, 323-326.	0.6	6
20	A custom ddPCR method for the detection of copy number variations in the nebulin triplicate region. <i>PLoS ONE</i> , 2022, 17, e0267793.	2.5	2
21	A commentary on identification of the rare compound heterozygous variants in the NEB gene in a Korean family with intellectual disability, epilepsy and early-childhood-onset generalized muscle weakness. <i>Journal of Human Genetics</i> , 2015, 60, 161-162.	2.3	1
22	Array Comparative Genomic Hybridisation and Droplet Digital PCR Uncover Recurrent Copy Number Variation of the TTN Segmental Duplication Region. <i>Genes</i> , 2022, 13, 905.	2.4	0