

# Suzanne Vanhauwaert

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6113053/publications.pdf>

Version: 2024-02-01

13  
papers

1,191  
citations

1040056

9  
h-index

1125743

13  
g-index

17  
all docs

17  
docs citations

17  
times ranked

3249  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cellular senescence in neuroblastoma. <i>British Journal of Cancer</i> , 2022, 126, 1529-1538.	6.4	5
2	Recurrent chromosomal imbalances provide selective advantage to human embryonic stem cells under enhanced replicative stress conditions. <i>Genes Chromosomes and Cancer</i> , 2021, 60, 272-281.	2.8	3
3	MYCN-induced nucleolar stress drives an early senescence-like transcriptional program in hTERT-immortalized RPE cells. <i>Scientific Reports</i> , 2021, 11, 14454.	3.3	6
4	A novel TLX1-driven T-ALL zebrafish model: comparative genomic analysis with other leukemia models. <i>Leukemia</i> , 2020, 34, 3398-3403.	7.2	12
5	Purification of high-quality RNA from a small number of fluorescence activated cell sorted zebrafish cells for RNA sequencing purposes. <i>BMC Genomics</i> , 2019, 20, 228.	2.8	10
6	ALK positively regulates MYCN activity through repression of HBP1 expression. <i>Oncogene</i> , 2019, 38, 2690-2705.	5.9	17
7	TBX2 is a neuroblastoma core regulatory circuitry component enhancing MYCN/FOXM1 reactivation of DREAM targets. <i>Nature Communications</i> , 2018, 9, 4866.	12.8	91
8	In silico discovery of a FOXM1 driven embryonal signaling pathway in therapy resistant neuroblastoma tumors. <i>Scientific Reports</i> , 2018, 8, 17468.	3.3	11
9	Expressed repetitive elements are broadly applicable reference targets for normalization of reverse transcription-qPCR data in mice. <i>Scientific Reports</i> , 2018, 8, 7642.	3.3	10
10	EV-TRACK: transparent reporting and centralizing knowledge in extracellular vesicle research. <i>Nature Methods</i> , 2017, 14, 228-232.	19.0	886
11	Refinement of the critical 2p25.3 deletion region: the role of MYT1L in intellectual disability and obesity. <i>Genetics in Medicine</i> , 2015, 17, 460-466.	2.4	45
12	Expressed Repeat Elements Improve RT-qPCR Normalization across a Wide Range of Zebrafish Gene Expression Studies. <i>PLoS ONE</i> , 2014, 9, e109091.	2.5	38
13	The Notch driven long non-coding RNA repertoire in T-cell acute lymphoblastic leukemia. <i>Haematologica</i> , 2014, 99, 1808-1816.	3.5	50