

# Stephan Werner

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11625838/publications.pdf>

Version: 2024-02-01

10  
papers

250  
citations

1163117

8  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

319  
citing authors

#	ARTICLE	IF	CITATIONS
1	NOseq: amplicon sequencing evaluation method for RNA m6A sites after chemical deamination. <i>Nucleic Acids Research</i> , 2021, 49, e23-e23.	14.5	25
2	Manganese Ions Individually Alter the Reverse Transcription Signature of Modified Ribonucleosides. <i>Genes</i> , 2020, 11, 950.	2.4	15
3	Machine learning of reverse transcription signatures of variegated polymerases allows mapping and discrimination of methylated purines in limited transcriptomes. <i>Nucleic Acids Research</i> , 2020, 48, 3734-3746.	14.5	45
4	Creating Auditory Illusions with Binaural Technology. <i>Modern Acoustics and Signal Processing</i> , 2020, , 623-663.	0.8	5
5	Graphical Workflow System for Modification Calling by Machine Learning of Reverse Transcription Signatures. <i>Frontiers in Genetics</i> , 2019, 10, 876.	2.3	10
6	A Vastly Increased Chemical Variety of RNA Modifications Containing a Thioacetal Structure. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 7893-7897.	13.8	44
7	Engineering of a DNA Polymerase for Direct m <sup>6</sup> A Sequencing. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 417-421.	13.8	66
8	Influences of Training on Externalization of Binaural Synthesis in Situations of Room Divergence. <i>AES: Journal of the Audio Engineering Society</i> , 2017, 65, 178-187.	1.0	15
9	Auditory Adaptation to Non-Individual HRTF Cues in Binaural Audio Reproduction. <i>AES: Journal of the Audio Engineering Society</i> , 2016, 64, 45-54.	1.0	9
10	CoverageAnalyzer (CA <sub>n</sub> ): A Tool for Inspection of Modification Signatures in RNA Sequencing Profiles. <i>Biomolecules</i> , 2016, 6, 42.	4.0	16