## Amir Bitran

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2048881/publications.pdf

Version: 2024-02-01

2258059 2550090 64 8 3 3 citations h-index g-index papers 11 11 11 96 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	Cotranslational folding allows misfolding-prone proteins to circumvent deep kinetic traps.  Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 1485-1495.	7.1	44
2	Validation of DBFOLD: An efficient algorithm for computing folding pathways of complex proteins. PLoS Computational Biology, 2020, 16, e1008323.	3.2	11
3	Mechanisms of fast and stringent search in homologous pairing of double-stranded DNA. PLoS Computational Biology, 2017, 13, e1005421.	3.2	7
4	Validation of DBFOLD: An efficient algorithm for computing folding pathways of complex proteins. , 2020, 16, e1008323.		0
5	Validation of DBFOLD: An efficient algorithm for computing folding pathways of complex proteins. , 2020, 16, e1008323.		O
6	Validation of DBFOLD: An efficient algorithm for computing folding pathways of complex proteins. , 2020, 16, e1008323.		0
7	Validation of DBFOLD: An efficient algorithm for computing folding pathways of complex proteins. , 2020, 16, e1008323.		O
8	Validation of DBFOLD: An efficient algorithm for computing folding pathways of complex proteins. , 2020, 16, e1008323.		0