Ting He

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3770084/publications.pdf

Version: 2024-02-01

13	330	1307594 7	1281871 11
papers	citations	h-index	g-index
13 all docs	13 docs citations	13 times ranked	387 citing authors

#	Article	IF	CITATIONS
1	Nanobody-Based Enzyme Immunoassay for Aflatoxin in Agro-Products with High Tolerance to Cosolvent Methanol. Analytical Chemistry, 2014, 86, 8873-8880.	6.5	126
2	Safety analysis of edible oil products via Raman spectroscopy. Talanta, 2019, 191, 324-332.	5.5	56
3	CRISPR-Cas12a <i>trans</i> -cleaves DNA G-quadruplexes. Chemical Communications, 2020, 56, 12526-12529.	4.1	40
4	Nanobody Technology for Mycotoxin Detection in the Field of Food Safety: Current Status and Prospects. Toxins, 2018, 10, 180.	3.4	35
5	Development of a biotinylated nanobody for sensitive detection of aflatoxin B1 in cereal via ELISA. Talanta, 2022, 239, 123125.	5.5	28
6	Determination of Aspergillus pathogens in agricultural products by a specific nanobody-polyclonal antibody sandwich ELISA. Scientific Reports, 2017, 7, 4348.	3.3	27
7	Enhancing the detection sensitivity of nanobody against aflatoxin B1 through structure-guided modification. International Journal of Biological Macromolecules, 2022, 194, 188-197.	7.5	8
8	Structural Insight into the Binding of TGIF1 to SIN3A PAH2 Domain through a C-Terminal Amphipathic Helix. International Journal of Molecular Sciences, 2021, 22, 12631.	4.1	5
9	Chemical shift assignments of a camelid nanobody against aflatoxin B1. Biomolecular NMR Assignments, 2019, 13, 75-78.	0.8	3
10	Chemical shift assignments of CHU_1110: an AHSA1-like protein from Cytophaga hutchinsonii. Biomolecular NMR Assignments, 2018, 12, 155-158.	0.8	1
11	Solution NMR structure of CHU_1110 from <i>Cytophaga hutchinsonii</i> , an AHSA1 protein potentially involved in metal ion stress response. Proteins: Structure, Function and Bioinformatics, 2019, 87, 91-95.	2.6	1
12	Chemical shift assignments of Mb1858 (24-155), a FHA domain-containing protein from Mycobacterium bovis. Biomolecular NMR Assignments, 2018, 12, 1-4.	0.8	0
13	Solution NMR structure and ligand identification of human Gas7 SH3 domain reveal a typical SH3 fold but a non-canonical ligand-binding mode. Biochemical and Biophysical Research Communications, 2019, 516, 1190-1195.	2.1	0