## Tage Thorstensen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/684294/publications.pdf

Version: 2024-02-01

623734 940533 1,119 16 14 16 citations g-index h-index papers 16 16 16 1493 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Over 2000-Fold Increased Production of the Leaderless Bacteriocin Garvicin KS by Increasing Gene Dose and Optimization of Culture Conditions. Frontiers in Microbiology, 2019, 10, 389.	3.5	12
2	Scandinavian perspectives on plant gene technology: applications, policies and progress. Physiologia Plantarum, 2018, 162, 219-238.	5.2	24
3	The ASH1-RELATED3 SET-Domain Protein Controls Cell Division Competence of the Meristem and the Quiescent Center of the Arabidopsis Primary Root  Â. Plant Physiology, 2014, 166, 632-643.	4.8	35
4	The <i>Arabidopsis</i> Histone Methyltransferase SUVR4 Binds Ubiquitin via a Domain with a Four-Helix Bundle Structure. Biochemistry, 2014, 53, 2091-2100.	2.5	7
5	SET domain proteins in plant development. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2011, 1809, 407-420.	1.9	99
6	The SUVR4 Histone Lysine Methyltransferase Binds Ubiquitin and Converts H3K9me1 to H3K9me3 on Transposon Chromatin in Arabidopsis. PLoS Genetics, 2011, 7, e1001325.	3.5	49
7	The CW domain, a new histone recognition module in chromatin proteins. EMBO Journal, 2011, 30, 1939-1952.	7.8	105
8	The ASH1 HOMOLOG 2 (ASHH2) Histone H3 Methyltransferase Is Required for Ovule and Anther Development in Arabidopsis. PLoS ONE, 2009, 4, e7817.	2.5	110
9	The Arabidopsis SET-domain protein ASHR3 is involved in stamen development and interacts with the bHLH transcription factor ABORTED MICROSPORES (AMS). Plant Molecular Biology, 2008, 66, 47-59.	3.9	69
10	Microarray-based method for detection of unknown genetic modifications. BMC Biotechnology, 2007, 7, 91.	3.3	38
11	Equal Performance of TaqMan, MGB, Molecular Beacon, and SYBR Green-Based Detection Assays in Detection and Quantification of Roundup Ready Soybean. Journal of Agricultural and Food Chemistry, 2006, 54, 9658-9663.	5.2	58
12	HIF- $\hat{\Pi}$ and iNOS levels in crucian carp gills during hypoxia-induced transformation. Journal of Comparative Physiology B: Biochemical, Systemic, and Environmental Physiology, 2006, 176, 359-369.	1.5	58
13	The Arabidopsis SUVR4 protein is a nucleolar histone methyltransferase with preference for monomethylated H3K9. Nucleic Acids Research, 2006, 34, 5461-5470.	14.5	55
14	ABI3 mediates expression of the peroxiredoxin antioxidant AtPER1 gene and induction by oxidative stress. Plant Molecular Biology, 2003, 53, 313-326.	3.9	45
15	Ten members of the Arabidopsis gene family encoding methyl-CpG-binding domain proteins are transcriptionally active and at least one, AtMBD11, is crucial for normal development. Nucleic Acids Research, 2003, 31, 5291-5304.	14.5	56
16	The Arabidopsis thaliana genome contains at least 29 active genes encoding SET domain proteins that can be assigned to four evolutionarily conserved classes. Nucleic Acids Research, 2001, 29, 4319-4333.	14.5	299