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Journal Information

Journal Title	ACTA BIOCHIMICA ET BIOPHYSICA SINICA
Publisher	SCIENCE PRESS
ISSN	1672-9145
E-ISSN	1745-7270
Database	Clarivate
Domain	Health Sciences, Life Sciences

Classification History

Year	A+	A	B	C
2026		✓	✓	
2025		✓	✓	
2024		✓	✓	
2023		✓	✓	
2021/2022		✓	✓	
2020		✓	✓	
2019		✓	✓	
2018		✓		

Classification: The table above shows all journal classifications recorded for each classification year in the Green Journals database. A journal may have more than one classification in the same year.

Recent Articles

MAERL accelerates atherosclerosis by inhibiting macrophage efferocytosis via transglutaminase 2
2026-10-1
Authors: Kun Xiao, Yajing Wang, Qianqian Wu et al.
DOI: 10.3724/abbs.2026092

lncRNA CHROMR promotes DLBCL viability and invasion in an m⁶A-dependent manner
2026-10-1
Authors: Chang Liu, Meina Wu, Chan Zhang et al.
DOI: 10.3724/abbs.2026151

Retraction to: Primate-specific sperm lnc-CLCN7 reveals embryo quality in IVF
2026-10-1
Authors: Xinrong Du, Rui Zheng, Sixian Wu et al.
DOI: 10.3724/abbs.2026194

Biochemical characterization of template-independent and template-dependent activities in DNA polymerase μ from <italic>Neurospora crassa</italic>
2026-10-1
Authors: Zipeng Zhao, Zhi Li, Yan Zhang et al.
DOI: 10.3724/abbs.2026197

HPV16 E6 promotes the proliferation of cervical cancer cells through the CPT1A/c-Myc positive feedback loop
2026-10-1
Authors: Hongyu Zhao, Le Kuai, Zhiwei Zhang et al.
DOI: 10.3724/abbs.2026195

Empagliflozin preserves mitochondrial function and mitigates oxidative stress-associated dysfunction in retinal pigment epithelial cells
2026-10-1
Authors: Haoxin Guo, Meijiao Zhu, Ruiyi Yan et al.
DOI: 10.3724/abbs.2026180

Unlocking PARP1: PCBP1 deacetylation reveals a checkpoint for DNA damage signaling
2026-10-1
Authors: Yuping Chen, Jian Yuan
DOI: 10.3724/abbs.2026189

Plant molecular farming: from gene design to scalable expression of therapeutic recombinant proteins
2026-10-1
Authors: Chunmei Guan, Nikolai V. Ravin, Yuling Jiao
DOI: 10.3724/abbs.2026193

Intracellular COMP drives ovarian cancer progression through AKT/ERK signaling with concomitant pathway-dependent transgelin upregulation
2026-10-1
Authors: Yuxin Zhao, Junyi Zhang, Yingying Hu et al.
DOI: 10.3724/abbs.2026158

Metabolic gene-based clustering identifies the role of acyl-CoA synthetase long-chain family member 1 in sepsis
2026-10-1
Authors: Dehao Zhu, Yinjiaozhi Li, Fan Wang et al.
DOI: 10.3724/abbs.2026148