## CONTENTS

**VOLUME 59 NUMBER 2**

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### REVIEW

**Abnormalities in alternative splicing in diabetes: therapeutic targets**

Zodwa Dlamini, Fortunate Mokoena & Rodney Hull

### RESEARCH

**Serum estradiol levels in controlled ovarian stimulation directly affect the endometrium**

Kamran Ullah, Tanzil Ur Rahman, Hai-Tao Pan, Meng-Xi Guo, Xin-Yan Dong, Juan Liu, Lu-Yang Jin, Yi Cheng, Zhang-Hong Ke, Jun Ren, Xian-Hua Lin, Xiao-Xiao Qiu, Ting-Ting Wang, He-Feng Huang & Jian-Zhong Sheng

**5-ALA ameliorates hepatic steatosis through AMPK signaling pathway**

Haoyong Yu, Mingliang Zhang, Yunqin Ma, Junyi Lu, Jiejin Pan, Pan Pan, Haibing Chen & Weiping Jia

**IGF1 potentiates the pro-inflammatory response in human peripheral blood mononuclear cells via MAPK**

Thalijn Liliana Catharina Wolters, Miha Gheorghe Netea, Adrianus Rudolfus Marinus Maria Hermus, Johannes Willem Adriaan Smit & Romana Teodora Netea-Maier

**Age-associated gene expression changes in the arcuate nucleus of male rhesus macaques**

Dominique H Eghlid, Vasilios T Gargfallou, Steven G Kohama & Henryk F Urbanski

**DMT efficiently inhibits hepatic gluconeogenesis by regulating the Gαq signaling pathway**

Ting-Ting Zhou, Fei Ma, Xiao-Fan Shi, Xin Xu, Te Du, Xiao-Dan Guo, Gai-Hong Wang, Liang Yu, Vatcharin Rukachaisirikul, Li-Hong Hu, Jing Chen & Xu Shen

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Content continued on the inside back cover
GPER modulates tone and coronary vascular reactivity in male and female rats 171–180
Angelina Rafaela Debortoli, Wender do Nascimento Rouver, Nathalie Tristão Banhos Delgado, Vinicius Mengal, Erick Roberto Gonçalves Claudio, Laena Pernomian, Lusiane Maria Bendhack, Margareth Ribeiro Moysés & Roger Lyrio dos Santos

miR-30e targets GLIPR-2 to modulate diabetic nephropathy: in vitro and in vivo experiments 181–190
Dong Zhao, Jinhua Jia & Hong Shao

Contents continued from outside back cover

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This image depicts glycoprotein hormone receptor glycosylation site candidates (in red) mapped on molecular surface representations of a model of the structure of the TSHR extracellular domain (ECD) in complex with the human monoclonal autoantibodies M22 (left and second left), and K1-70 (second right and right). Antibody light chains are shown in green, heavy chains in blue, and the TSHR ECD in yellow. (left and second left) are different points of view related by rotations of ~180° along a vertical axis as are (second right and right). From Núñez Miguel et al. 58 25–41.

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