MINIREVIEW
Adrenaline Rush: The Role of Adrenergic Receptors in Stimulant-Induced Behaviors
Karl T. Schmidt and David Weinshenker

ARTICLES
Ah Receptor–Mediated Suppression of Liver Regeneration through NC-XRE–Driven \( \text{p21}^{Cip1} \) Expression
Daniel P. Jackson, Hui Li, Kristen A. Mitchell, Aditya D. Joshi, and Cornelis J. Elferink

\( \beta \)- Arrestin1 and Distinct CXCR4 Structures Are Required for Stromal Derived Factor-1 to Downregulate CXCR4 Cell-Surface Levels in Neuroblastoma
Ian C. Clift, Adebowale O. Bamidele, Christie Rodriguez-Ramirez, Kimberly N. Kremer, and Karen E. Hedin

Allosteric Noncompetitive Small Molecule Selective Inhibitors of CD45 Tyrosine Phosphatase Suppress T-Cell Receptor Signals and Inflammation In Vivo
Michael D. Perron, Shafinaz Chowdhury, Isabelle Aubry, Enrico Purisima, Michel L. Tremblay, and H. Uri Saragovi

Eudistomin D and Penaresin Derivatives as Modulators of Ryanodine Receptor Channels and Sarcoplasmic Reticulum Ca\(^{2+}\) ATPase in Striated Muscle
Paula L. Diaz-Sylvester, Maura Porta, Vanessa V. Juettner, Yuanzhao Lv, Sidney Fleischer, and Julio A. Copello

The Novel Arsenical Darinaparsin Is Transported by Cystine Importing Systems

G\( \alpha_{12} \) Structural Determinants of Hsp90 Interaction Are Necessary for Serum Response Element–Mediated Transcriptional Activation

Structure/Activity Relationships of (M)ANT- and TNP-Nucleotides for Inhibition of Rat Soluble Guanylyl Cyclase \( \alpha_{3\beta_1} \)
Stefan Dove, Kerstin Yvonne Danker, Johannes-Peter Stasch, Volkhard Kaever, and Roland Seifert

Observed Drug-Receptor Association Rates Are Governed by Membrane Affinity: The Importance of Establishing “Micro-Pharmacokinetic/Pharmacodynamic Relationships” at the \( \beta_2 \)-Adrenoceptor
David A. Sykes, Cheryl Parry, John Reilly, Penny Wright, Robin A. Fairhurst, and Steven J. Charlton

Regulation of GluA1 \( \alpha \)-Amino-3-Hydroxy-5-Methyl-4-Isoxazolepropionic Acid Receptor Function by Protein Kinase C at Serine-818 and Threonine-840
Meagan A. Jenkins, Gordon Wells, Julia Bachman, James P. Snyder, Andrew Jenkins, Richard L. Huganir, Robert E. Oswald, and Stephen F. Traynelis
Nanomolar Bifenthrin Alters Synchronous Ca²⁺ Oscillations and Cortical Neuron Development Independent of Sodium Channel Activity
Zhengyu Cao, Yanjun Cui, Hai M. Nguyen, David Paul Jenkins, Heike Wulff, and Isaac N. Pessah

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About the cover: Models of the interactions of inhibitors with sGCα₁β₁: binding site of bis-MANT-ITP, represented by the lipophilic potential mapped onto a MOLCAD Connolly surface. See the article by Dove et al. (dx.doi.org/10.1124/mol.113.091017).