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 p21Cip1 Expression
Daniel P. Jackson, Hui Li, Kristen A. Mitchell, Aditya D. Joshi, and Cornelis J. Elferink

β-Arcresin1 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 Ca2+ 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

Gα12 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 α3β3
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 β2-Adrenoceptor
David A. Sykes, Cheryl Parry, John Reilly, Penny Wright, Robin A. Fairhurst, and Steven J. Charlton

Regulation of GluA1 α-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 Ca$^{2+}$ 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 sGC$\alpha_1$$\beta_1$: 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).