

Robert E Oswald

List of Publications by Year in descending order

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52
papers

1,781
citations

236925

25
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265206

42
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53
all docs

53
docs citations

53
times ranked

1422
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | In Vitro Effects of (+)MK-801 (dizocilpine) and Memantine on β -Amyloid Peptides Linked to Alzheimer's Disease. <i>Biochemistry</i> , 2020, 59, 4517-4522. | 2.5 | 0 |
| 2 | Surface water and groundwater analysis using aryl hydrocarbon and endocrine receptor biological assays and liquid chromatography-high resolution mass spectrometry in Susquehanna County, PA. <i>Environmental Sciences: Processes and Impacts</i> , 2019, 21, 988-998. | 3.5 | 3 |
| 3 | Comparative screening of recombinant antigen thermostability for improved leptospirosis vaccine design. <i>Biotechnology and Bioengineering</i> , 2019, 116, 260-271. | 3.3 | 6 |
| 4 | Noncompetitive antagonists induce cooperative AMPA receptor channel gating. <i>Journal of General Physiology</i> , 2019, 151, 156-173. | 1.9 | 15 |
| 5 | Modulation of AMPA Receptor Gating by the Anticonvulsant Drug, Perampanel. <i>ACS Medicinal Chemistry Letters</i> , 2019, 10, 237-242. | 2.8 | 18 |
| 6 | 7-Phenoxy-Substituted 3,4-Dihydro-2H-1,2,4-benzothiadiazine 1,1-Dioxides as Positive Allosteric Modulators of α -Amino-3-hydroxy-5-methyl-4-isoxazolepropionic Acid (AMPA) Receptors with Nanomolar Potency. <i>Journal of Medicinal Chemistry</i> , 2018, 61, 251-264. | 6.4 | 41 |
| 7 | New Insights into the Mechanism of Ca^{2+} -Dependent Inactivation of NMDA Receptors. <i>Biophysical Journal</i> , 2017, 113, 2131-2132. | 0.5 | 2 |
| 8 | Extended low-resolution structure of a <i>Leptospira</i> antigen offers high bactericidal antibody accessibility amenable to vaccine design. <i>ELife</i> , 2017, 6, . | 6.0 | 12 |
| 9 | Current Recording and Kinetic Analyses for Single AMPA Receptors. <i>NeuroMethods</i> , 2016, , 257-272. | 0.3 | 2 |
| 10 | NMR Approaches to Functional Dynamics of Genetically Separated iGluR Domains. <i>NeuroMethods</i> , 2016, , 101-118. | 0.3 | 1 |
| 11 | Long-term impacts of unconventional drilling operations on human and animal health. <i>Journal of Environmental Science and Health - Part A Toxic/Hazardous Substances and Environmental Engineering</i> , 2015, 50, 447-459. | 1.7 | 28 |
| 12 | Role of Stoichiometry in the Dimer-Stabilizing Effect of AMPA Receptor Allosteric Modulators. <i>ACS Chemical Biology</i> , 2014, 9, 128-133. | 3.4 | 9 |
| 13 | Thermodynamics and Mechanism of the Interaction of Willardiine Partial Agonists with a Glutamate Receptor: Implications for Drug Development. <i>Biochemistry</i> , 2014, 53, 3790-3795. | 2.5 | 8 |
| 14 | Dynamics of Cleft Closure of the GluA2 Ligand-binding Domain in the Presence of Full and Partial Agonists Revealed by Hydrogen-Deuterium Exchange. <i>Journal of Biological Chemistry</i> , 2013, 288, 27658-27666. | 3.4 | 27 |
| 15 | Impacts of Gas Drilling on Human and Animal Health. <i>New Solutions</i> , 2012, 22, 51-77. | 1.2 | 185 |
| 16 | The Structure of (β)-Kaitocephalin Bound to the Ligand Binding Domain of the (S)- α -Amino-3-hydroxy-5-methyl-4-isoxazolepropionic Acid (AMPA)/Glutamate Receptor, GluA2. <i>Journal of Biological Chemistry</i> , 2012, 287, 41007-41013. | 3.4 | 10 |
| 17 | The Loss of an Electrostatic Contact Unique to AMPA Receptor Ligand Binding Domain 2 Slows Channel Activation. <i>Biochemistry</i> , 2012, 51, 4015-4027. | 2.5 | 9 |
| 18 | Mechanisms of Modal Activation of GluA3 Receptors. <i>Molecular Pharmacology</i> , 2011, 80, 49-59. | 2.3 | 41 |

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|----|---|-----|-----------|
| 19 | Mechanism of AMPA Receptor Activation by Partial Agonists. <i>Journal of Biological Chemistry</i> , 2011, 286, 35257-35266. | 3.4 | 39 |
| 20 | On the Mechanisms of \pm -Amino-3-hydroxy-5-methylisoxazole-4-propionic Acid (AMPA) Receptor Binding to Glutamate and Kainate. <i>Journal of Biological Chemistry</i> , 2010, 285, 12334-12343. | 3.4 | 26 |
| 21 | Hydrophobic Side Chain Dynamics of a Glutamate Receptor Ligand Binding Domain. <i>Journal of Biological Chemistry</i> , 2010, 285, 10154-10162. | 3.4 | 16 |
| 22 | Characterizing Single-Channel Behavior of GluA3 Receptors. <i>Biophysical Journal</i> , 2010, 99, 1437-1446. | 0.5 | 37 |
| 23 | Molecular Mechanism of Flop Selectivity and Subsite Recognition for an AMPA Receptor Allosteric Modulator: Structures of GluA2 and GluA3 in Complexes with PEPA. <i>Biochemistry</i> , 2010, 49, 2843-2850. | 2.5 | 31 |
| 24 | Piracetam Defines a New Binding Site for Allosteric Modulators of \pm -Amino-3-hydroxy-5-methyl-4-isoxazole-propionic Acid (AMPA) Receptors. <i>Journal of Medicinal Chemistry</i> , 2010, 53, 2197-2203. | 6.4 | 58 |
| 25 | Structure of the S1S2 glutamate binding domain of GLuR3. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 75, 628-637. | 2.6 | 44 |
| 26 | Probing the Allosteric Modulator Binding Site of GluR2 with Thiazide Derivatives. <i>Biochemistry</i> , 2009, 48, 8594-8602. | 2.5 | 65 |
| 27 | Mechanisms of Antagonism of the GluR2 AMPA Receptor: Structure and Dynamics of the Complex of Two Willardiine Antagonists with the Glutamate Binding Domain. <i>Biochemistry</i> , 2009, 48, 3894-3903. | 2.5 | 37 |
| 28 | NMR Spectroscopy of the Ligand-Binding Core of Ionotropic Glutamate Receptor 2 Bound to 5-Substituted Willardiine Partial Agonists. <i>Journal of Molecular Biology</i> , 2008, 378, 673-685. | 4.2 | 33 |
| 29 | Mechanism of Partial Agonism at the GluR2 AMPA Receptor: Measurements of Lobe Orientation in Solution. <i>Biochemistry</i> , 2008, 47, 10600-10610. | 2.5 | 38 |
| 30 | Structure of Glutamate Receptors. <i>Current Drug Targets</i> , 2007, 8, 573-582. | 2.1 | 33 |
| 31 | Dynamics of the S1S2 Glutamate Binding Domain of GluR2 Measured Using ^{19}F NMR Spectroscopy. <i>Journal of Biological Chemistry</i> , 2007, 282, 12773-12784. | 3.4 | 44 |
| 32 | Molecular properties of local anesthetics as predictors of affinity for nicotinic acetylcholine receptors. <i>Journal of Neuroscience Research</i> , 2007, 85, 2943-2949. | 2.9 | 5 |
| 33 | Backbone chemical shift assignment of a glutamate receptor ligand binding domain in complexes with five partial agonists. <i>Biomolecular NMR Assignments</i> , 2007, 1, 241-243. | 0.8 | 5 |
| 34 | Ionotropic Glutamate Receptor Recognition and Activation. <i>Advances in Protein Chemistry</i> , 2004, 68, 313-349. | 4.4 | 32 |
| 35 | Emerging structural explanations of ionotropic glutamate receptor function. <i>FASEB Journal</i> , 2004, 18, 428-438. | 0.5 | 54 |
| 36 | cDNA sequence and in vitro folding of GsMTx4, a specific peptide inhibitor of mechanosensitive channels. <i>Toxicon</i> , 2003, 42, 263-274. | 1.6 | 74 |

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|----|--|-----|-----------|
| 37 | Solution Structure of Peptide Toxins That Block Mechanosensitive Ion Channels. <i>Journal of Biological Chemistry</i> , 2002, 277, 34443-34450. | 3.4 | 88 |
| 38 | Structural Mobility of the Extracellular Ligand-Binding Core of an Ionotropic Glutamate Receptor. Analysis of NMR Relaxation Dynamics. <i>Biochemistry</i> , 2002, 41, 10472-10481. | 2.5 | 84 |
| 39 | Semi-automated backbone resonance assignments of the extracellular ligand-binding domain of an ionotropic glutamate receptor. <i>Journal of Biomolecular NMR</i> , 2002, 22, 297-298. | 2.8 | 13 |
| 40 | Concerted Motion of a Protein-Peptide Complex: Backbone Dynamics Studies of an ¹⁵ N-Labeled Peptide Derived from P21-Activated Kinase Bound to Cdc42Hs-GMPPCP. <i>Biochemistry</i> , 2001, 40, 14368-14375. | 2.5 | 16 |
| 41 | Structural Insights Into NMDA Ionotropic Glutamate Receptors via Molecular Modelling. <i>Journal of Molecular Modeling</i> , 2000, 6, 16-25. | 1.8 | 8 |
| 42 | Backbone Dynamics of Inactive, Active, and Effector-Bound Cdc42Hs from Measurements of ¹⁵ N Relaxation Parameters at Multiple Field Strengths. <i>Biochemistry</i> , 1999, 38, 12547-12557. | 2.5 | 49 |
| 43 | Identification of the Binding Surface on Cdc42Hs for p21-Activated Kinase. <i>Biochemistry</i> , 1998, 37, 14030-14037. | 2.5 | 38 |
| 44 | Unraveling the modular design of glutamate-gated ion channels. <i>Trends in Neurosciences</i> , 1995, 18, 161-168. | 8.6 | 264 |
| 45 | Asn-265 of frog kainate binding protein is a functional glycosylation site: implications for the transmembrane topology of glutamate receptors. <i>FEBS Letters</i> , 1995, 368, 230-234. | 2.8 | 7 |
| 46 | Model Building Predicts an Additional Conformational Switch when GTP Binds to the CDC42HS Protein. <i>Protein and Peptide Letters</i> , 1994, 1, 84-91. | 0.9 | 6 |
| 47 | A reevaluation of the mathematical models for simulating single-channel and whole-cell ionic currents. <i>Synapse</i> , 1988, 2, 97-103. | 1.2 | 8 |
| 48 | Effects of pineal factors on the action potentials of sympathetic neurons. <i>Cellular and Molecular Neurobiology</i> , 1986, 6, 381-395. | 3.3 | 3 |
| 49 | Effects of Calcium on the Binding of Phencyclidine to Acetylcholine Receptor-Rich Membrane Fragments from <i>Torpedo californica</i> Electroplaque. <i>Journal of Neurochemistry</i> , 1983, 41, 1077-1084. | 3.9 | 15 |
| 50 | Evidence for a skeleton in acetylcholine receptor-rich membranes from <i>Torpedo marmorata</i> electric organ. <i>FEBS Letters</i> , 1982, 145, 250-257. | 2.8 | 12 |
| 51 | Crosslinking of α -bungarotoxin to the acetylcholine receptor from <i>Torpedo marmorata</i> by ultraviolet light irradiation. <i>FEBS Letters</i> , 1982, 139, 225-229. | 2.8 | 76 |
| 52 | Analysis of α -Bungarotoxin Binding in the Goldfish Central Nervous System. <i>Journal of Neurochemistry</i> , 1981, 37, 1586-1593. | 3.9 | 6 |