## Albert Batushansky

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	An In Vivo Stable Isotope Labeling Method to Investigate Individual Matrix Protein Synthesis, Ribosomal Biogenesis, and Cellular Proliferation in Murine Articular Cartilage. Function, 2022, 3, zqac008.	2.3	8
2	Insulin-like growth factor 1 receptor mediates photoreceptor neuroprotection. Cell Death and Disease, 2022, 13, .	6.3	7
3	Sirt5 Deficiency Causes Posttranslational Protein Malonylation and Dysregulated Cellular Metabolism in Chondrocytes Under Obesity Conditions. Cartilage, 2021, 13, 1185S-1199S.	2.7	16
4	Increasing Glycolysis Protects Cardiac Function Against High Fat Dietâ€Induced Cardiomyopathy. FASEB Journal, 2021, 35, .	0.5	0
5	PFKFB3â€dependent glucose metabolism regulates 3T3â€L1 adipocyte development. FASEB Journal, 2021, 35, e21728.	0.5	3
6	Correlation network analysis shows divergent effects of a long-term, high-fat diet and exercise on early stage osteoarthritis phenotypes in mice. Journal of Sport and Health Science, 2020, 9, 119-131.	6.5	17
7	The complex response of free and bound amino acids to water stress during the seed setting stage in Arabidopsis. Plant Journal, 2020, 102, 838-855.	5.7	9
8	Proximal colon–derived O-glycosylated mucus encapsulates and modulates the microbiota. Science, 2020, 370, 467-472.	12.6	122
9	A Shift in Glycerolipid Metabolism Defines the Follicular Fluid of IVF Patients with Unexplained Infertility. Biomolecules, 2020, 10, 1135.	4.0	9
10	Can metabolic tightening and expansion of co-expression network play a role in stress response and tolerance?. Plant Science, 2020, 293, 110409.	3.6	11
11	Enhancing cardiac glycolysis causes an increase in PDK4 content in response to short-term high-fat diet. Journal of Biological Chemistry, 2019, 294, 16831-16845.	3.4	13
12	GC–MS metabolic profiling reveals fructose-2,6-bisphosphate regulates branched chain amino acid metabolism in the heart during fasting. Metabolomics, 2019, 15, 18.	3.0	18
13	Adaptive responses of amino acid metabolism to the combination of desiccation and low nitrogen availability in Sporobolus stapfianus. Planta, 2019, 249, 1535-1549.	3.2	4
14	Topological Data Analysis as a Morphometric Method: Using Persistent Homology to Demarcate a Leaf Morphospace. Frontiers in Plant Science, 2018, 9, 553.	3.6	62
15	Independent effects of dietary fat and sucrose content on chondrocyte metabolism and osteoarthritis pathology in mice. DMM Disease Models and Mechanisms, 2018, 11, .	2.4	20
16	Network-Guided GWAS Improves Identification of Genes Affecting Free Amino Acids. Plant Physiology, 2017, 173, 872-886.	4.8	52
17	A combination of stomata deregulation and a distinctive modulation of amino acid metabolism are associated with enhanced tolerance of wheat varieties to transient drought. Metabolomics, 2017, 13, 1.	3.0	6
18	Correlation-Based Network Generation, Visualization, and Analysis as a Powerful Tool in Biological Studies: A Case Study in Cancer Cell Metabolism. BioMed Research International, 2016, 2016, 1-9.	1.9	68

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19	Environmental and genetic effects on tomato seed metabolic balance and its association with germination vigor. BMC Genomics, 2016, 17, 1047.	2.8	28
20	Paclobutrazol induces tolerance in tomato to deficit irrigation through diversified effects on plant morphology, physiology and metabolism. Scientific Reports, 2016, 6, 39321.	3.3	47
21	Effects of Parental Temperature and Nitrate on Seed Performance are Reflected by Partly Overlapping Genetic and Metabolic Pathways. Plant and Cell Physiology, 2016, 57, 473-487.	3.1	37
22	Metabolic and Physiological Responses of Shiraz and Cabernet Sauvignon (Vitis vinifera L.) to Near Optimal Temperatures of 25 and 35 °C. International Journal of Molecular Sciences, 2015, 16, 24276-24294.	4.1	52
23	Sulfite Oxidase Activity Is Essential for Normal Sulfur, Nitrogen and Carbon Metabolism in Tomato Leaves. Plants, 2015, 4, 573-605.	3.5	22
24	The transporter GAT1 plays an important role in GABA-mediated carbon-nitrogen interactions in Arabidopsis. Frontiers in Plant Science, 2015, 6, 785.	3.6	30
25	Combined correlationâ€based network and <scp>mQTL</scp> analyses efficiently identified loci for branchedâ€chain amino acid, serine to threonine, and proline metabolism in tomato seeds. Plant Journal, 2015, 81, 121-133.	5.7	55
26	Combined Transcriptomics and Metabolomics of Arabidopsis thaliana Seedlings Exposed to Exogenous GABA Suggest Its Role in Plants Is Predominantly Metabolic. Molecular Plant, 2014, 7, 1065-1068.	8.3	56
27	Metabolite and transcript profiling of berry skin during fruit development elucidates differential regulation between Cabernet Sauvignon and Shiraz cultivars at branching points in the polyphenol pathway. BMC Plant Biology, 2014, 14, 188.	3.6	135
28	Metabolite Profiling and Integrative Modeling Reveal Metabolic Constraints for Carbon Partitioning under Nitrogen Starvation in the Green Algae Haematococcus pluvialis. Journal of Biological Chemistry, 2014, 289, 30387-30403.	3.4	103
29	Growth, lipid production and metabolic adjustments in the euryhaline eustigmatophyte Nannochloropsis oceanica CCALA 804 in response to osmotic downshift. Applied Microbiology and Biotechnology, 2013, 97, 8291-8306.	3.6	65
30	Growth Platform-Dependent and -Independent Phenotypic and Metabolic Responses of Arabidopsis and Its Halophytic Relative, Eutrema salsugineum, to Salt Stress. Plant Physiology, 2013, 162, 1583-1598.	4.8	50
31	The Investment in Scent: Time-Resolved Metabolic Processes in Developing Volatile-Producing Nigella sativa L. Seeds. PLoS ONE, 2013, 8, e73061.	2.5	5