## Michael S Behnke

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Development and application of classical genetics in Toxoplasma gondii. , 2020, , 859-896.		2
2	An Important Role for CD4 <sup>+</sup> T Cells in Adaptive Immunity to Toxoplasma gondii in Mice Lacking the Transcription Factor Batf3. MSphere, 2020, 5, .	1.3	8
3	Evolution of resistance in vitro reveals mechanisms of artemisinin activity in <i>Toxoplasma gondii</i> . Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 26881-26891.	3.3	30
4	WRN conditioned media is sufficient for <i>in vitro</i> propagation of intestinal organoids from large farm and small companion animals. Biology Open, 2017, 6, 698-705.	0.6	88
5	QTL Mapping and CRISPR/Cas9 Editing to Identify a Drug Resistance Gene in <em>Toxoplasma gondii</em> . Journal of Visualized Experiments, 2017, , .	0.2	1
6	The Past, Present, and Future of Genetic Manipulation in Toxoplasma gondii. Trends in Parasitology, 2016, 32, 542-553.	1.5	36
7	Genetic Mapping of Pathogenesis Determinants in <i>Toxoplasma gondii</i> . Annual Review of Microbiology, 2016, 70, 63-81.	2.9	49
8	Local admixture of amplified and diversified secreted pathogenesis determinants shapes mosaic Toxoplasma gondii genomes. Nature Communications, 2016, 7, 10147.	5.8	243
9	Genetic Mapping Reveals that Sinefungin Resistance in Toxoplasma gondii Is Controlled by a Putative Amino Acid Transporter Locus That Can Be Used as a Negative Selectable Marker. Eukaryotic Cell, 2015, 14, 140-148.	3.4	29
10	Rhoptry Proteins ROP5 and ROP18 Are Major Murine Virulence Factors in Genetically Divergent South American Strains of Toxoplasma gondii. PLoS Genetics, 2015, 11, e1005434.	1.5	99
11	NextGen sequencing reveals short double crossovers contribute disproportionately to genetic diversity in Toxoplasma gondii. BMC Genomics, 2014, 15, 1168.	1.2	17
12	Toxoplasma gondii merozoite gene expression analysis with comparison to the life cycle discloses a unique expression state during enteric development. BMC Genomics, 2014, 15, 350.	1.2	80
13	Genotyping Toxoplasma gondii from wildlife in Pennsylvania and identification of natural recombinants virulent to mice. Veterinary Parasitology, 2014, 200, 74-84.	0.7	58
14	miR-146a and miR-155 Delineate a MicroRNA Fingerprint Associated with Toxoplasma Persistence in the Host Brain. Cell Reports, 2014, 6, 928-937.	2.9	96
15	The Polymorphic Pseudokinase ROP5 Controls Virulence in Toxoplasma gondii by Regulating the Active Kinase ROP18. PLoS Pathogens, 2012, 8, e1002992.	2.1	153
16	Compensatory dendritic cell development mediated by BATF–IRF interactions. Nature, 2012, 490, 502-507.	13.7	367
17	Cell cycleâ€dependent, intercellular transmission of <i>Toxoplasma gondii</i> is accompanied by marked changes in parasite gene expression. Molecular Microbiology, 2011, 79, 192-204.	1.2	57
18	Virulence differences in <i>Toxoplasma</i> mediated by amplification of a family of polymorphic pseudokinases. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 9631-9636.	3.3	230

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19	A Systematic Screen to Discover and Analyze Apicoplast Proteins Identifies a Conserved and Essential Protein Import Factor. PLoS Pathogens, 2011, 7, e1002392.	2.1	221
20	A novel multifunctional oligonucleotide microarray for Toxoplasma gondii. BMC Genomics, 2010, 11, 603.	1.2	57
21	T. gondii RP Promoters & Knockdown Reveal Molecular Pathways Associated with Proliferation and Cell-Cycle Arrest. PLoS ONE, 2010, 5, e14057.	1.1	28
22	Phosphorylation of Immunity-Related GTPases by a Toxoplasma gondii-Secreted Kinase Promotes Macrophage Survival and Virulence. Cell Host and Microbe, 2010, 8, 484-495.	5.1	286
23	Coordinated Progression through Two Subtranscriptomes Underlies the Tachyzoite Cycle of Toxoplasma gondii. PLoS ONE, 2010, 5, e12354.	1.1	248
24	Phenotypic and Gene Expression Changes among Clonal Type I Strains of <i>Toxoplasma gondii</i> . Eukaryotic Cell, 2009, 8, 1828-1836.	3.4	76
25	The transcription of bradyzoite genes in <i>Toxoplasma gondii</i> is controlled by autonomous promoter elements. Molecular Microbiology, 2008, 68, 1502-1518.	1.2	91
26	A comprehensive SAGE database for the analysis of $\hat{I}^{3}\hat{I}$ T cells. International Immunology, 2006, 18, 613-626.	1.8	11
27	Changes in the Expression of Human Cell Division Autoantigen-1 Influence Toxoplasma gondii Growth and Development. PLoS Pathogens, 2006, 2, e105.	2.1	81
28	A Secreted Serine-Threonine Kinase Determines Virulence in the Eukaryotic Pathogen Toxoplasma gondii. Science, 2006, 314, 1776-1780.	6.0	520
29	Biochemical and genetic analysis of the distinct proliferating cell nuclear antigens of Toxoplasma gondii. Molecular and Biochemical Parasitology, 2005, 142, 56-65.	0.5	22
30	The transcriptome of Toxoplasma gondii. BMC Biology, 2005, 3, 26.	1.7	167
31	Genetic rescue of a Toxoplasma gondii conditional cell cycle mutant. Molecular Microbiology, 2004, 55, 1060-1071.	1.2	28
32	Serial Analysis of Gene Expression in Circulating γδT Cell Subsets Defines Distinct Immunoregulatory Phenotypes and Unexpected Gene Expression Profiles. Journal of Immunology, 2003, 170, 356-364.	0.4	78