# Michael Anthony J Ferguson

# List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

293 papers

14,424 citations

61 h-index 108 g-index

311 ext. papers

15,504 ext. citations

**6.1** avg, IF

6.19 L-index

#	Paper	IF	Citations
293	Visualisation of experimentally determined and predicted protein N-glycosylation and predicted glycosylphosphatidylinositol anchor addition in <i>Wellcome Open Research</i> , <b>2022</b> , 7, 33	4.8	O
292	Visualisation of proteome-wide ordered protein abundances in Wellcome Open Research, 2022, 7, 34	4.8	0
291	The Leishmania donovani Ortholog of the Glycosylphosphatidylinositol Anchor Biosynthesis Cofactor PBN1 Is Essential for Host Infection <i>MBio</i> , <b>2022</b> , e0043322	7.8	1
290	Proteomic identification of the UDP-GlcNAc: PI 🛭 - 6 GlcNAc-transferase subunits of the glycosylphosphatidylinositol biosynthetic pathway of Trypanosoma brucei. <i>PLoS ONE</i> , <b>2021</b> , 16, e02446	59 <sup>3</sup> 9 <sup>7</sup>	2
289	Multiple unbiased approaches identify oxidosqualene cyclase as the molecular target of a promising anti-leishmanial. <i>Cell Chemical Biology</i> , <b>2021</b> , 28, 711-721.e8	8.2	4
288	Nucleotide sugar biosynthesis occurs in the glycosomes of procyclic and bloodstream form Trypanosoma brucei. <i>PLoS Neglected Tropical Diseases</i> , <b>2021</b> , 15, e0009132	4.8	6
287	Elimination of GPI2 suppresses glycosylphosphatidylinositol GlcNAc transferase activity and alters GPI glycan modification in Trypanosoma brucei. <i>Journal of Biological Chemistry</i> , <b>2021</b> , 297, 100977	5.4	2
286	A broadly active fucosyltransferase LmjFUT1 whose mitochondrial localization and activity are essential in parasitic. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2021</b> , 118,	11.5	3
285	An essential, kinetoplastid-specific GDP-Fuc: ED-Gal El ,2-fucosyltransferase is located in the mitochondrion of. <i>ELife</i> , <b>2021</b> , 10,	8.9	4
284	A Trypanosoma brucei B glycosyltransferase superfamily gene encodes a 🛭 -6 GlcNAc-transferase mediating N-glycan and GPI anchor modification. <i>Journal of Biological Chemistry</i> , <b>2021</b> , 297, 101153	5.4	1
283	CAP-MAP: cap analysis protocol with minimal analyte processing, a rapid and sensitive approach to analysing mRNA cap structures. <i>Open Biology</i> , <b>2020</b> , 10, 190306	7	15
282	A mechanism-inspired UDPacetylglucosamine pyrophosphorylase inhibitor. <i>RSC Chemical Biology</i> , <b>2020</b> , 1, 13-25	3	6
281	Setting Our Sights on Infectious Diseases. <i>ACS Infectious Diseases</i> , <b>2020</b> , 6, 3-13	5.5	9
280	Preclinical candidate for the treatment of visceral leishmaniasis that acts through proteasome inhibition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2019</b> , 116, 9318-9323	11.5	65
279	Reprogramming of Trypanosoma cruzi metabolism triggered by parasite interaction with the host cell extracellular matrix. <i>PLoS Neglected Tropical Diseases</i> , <b>2019</b> , 13, e0007103	4.8	19
278	Phosphomannomutase and Guanosine Diphosphate-Mannose Pyrophosphorylase Ligandability Assessment. <i>Antimicrobial Agents and Chemotherapy</i> , <b>2019</b> , 63,	5.9	5
277	Proteome turnover in the bloodstream and procyclic forms of measured by quantitative proteomics. <i>Wellcome Open Research</i> , <b>2019</b> , 4, 152	4.8	11

276	Pharmacological Validation of N-Myristoyltransferase as a Drug Target in Leishmania donovani. <i>ACS Infectious Diseases</i> , <b>2019</b> , 5, 111-122	5.5	31
275	Proteomic Analysis of the Cell Cycle of Procylic Form. <i>Molecular and Cellular Proteomics</i> , <b>2018</b> , 17, 1184-	-1/1/95	17
274	Cyclin-dependent kinase 12 is a drug target for visceral leishmaniasis. <i>Nature</i> , <b>2018</b> , 560, 192-197	50.4	73
273	African trypanosomes evade immune clearance by O-glycosylation of the VSG surface coat. <i>Nature Microbiology</i> , <b>2018</b> , 3, 932-938	26.6	30
272	N-glycan microheterogeneity regulates interactions of plasma proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2018</b> , 115, 8763-8768	11.5	62
271	Gluconeogenesis using glycerol as a substrate in bloodstream-form Trypanosoma brucei. <i>PLoS Pathogens</i> , <b>2018</b> , 14, e1007475	7.6	14
270	The mRNA cap methyltransferase gene TbCMT1 is not essential in vitro but is a virulence factor in vivo for bloodstream form Trypanosoma brucei. <i>PLoS ONE</i> , <b>2018</b> , 13, e0201263	3.7	2
269	Anti-trypanosomatid drug discovery: an ongoing challenge and a continuing need. <i>Nature Reviews Microbiology</i> , <b>2017</b> , 15, 217-231	22.2	225
268	Fluorescent mannosides serve as acceptor substrates for glycosyltransferase and sugar-1-phosphate transferase activities in Euglena gracilis membranes. <i>Carbohydrate Research</i> , <b>2017</b> , 438, 26-38	2.9	12
267	Single-subunit oligosaccharyltransferases of display different and predictable peptide acceptor specificities. <i>Journal of Biological Chemistry</i> , <b>2017</b> , 292, 20328-20341	5.4	10
266	Genetic metabolic complementation establishes a requirement for GDP-fucose in. <i>Journal of Biological Chemistry</i> , <b>2017</b> , 292, 10696-10708	5.4	9
265	Prediction of Protein Complexes in by Protein Correlation Profiling Mass Spectrometry and Machine Learning. <i>Molecular and Cellular Proteomics</i> , <b>2017</b> , 16, 2254-2267	7.6	20
264	Proteomic Identification of Immunodiagnostic Antigens for Trypanosoma vivax Infections in Cattle and Generation of a Proof-of-Concept Lateral Flow Test Diagnostic Device. <i>PLoS Neglected Tropical Diseases</i> , <b>2016</b> , 10, e0004977	4.8	15
263	A Gene of the B-Glycosyltransferase Family Encodes N-Acetylglucosaminyltransferase II Function in Trypanosoma brucei. <i>Journal of Biological Chemistry</i> , <b>2016</b> , 291, 13834-45	5.4	9
262	Global Membrane Protein Interactome Analysis using In vivo Crosslinking and Mass Spectrometry-based Protein Correlation Profiling. <i>Molecular and Cellular Proteomics</i> , <b>2016</b> , 15, 2476-90	7.6	46
261	Leishmania major UDP-sugar pyrophosphorylase salvages galactose for glycoconjugate biosynthesis. <i>International Journal for Parasitology</i> , <b>2015</b> , 45, 783-90	4.3	8
260	Identification of a glycosylphosphatidylinositol anchor-modifying <b>1</b> -3 galactosyltransferase in Trypanosoma brucei. <i>Glycobiology</i> , <b>2015</b> , 25, 438-47	5.8	12
259	Molecular control of irreversible bistability during trypanosome developmental commitment.  Journal of Cell Biology, 2015, 211, 455-68	7.3	35

258	TrypanoCyc: a community-led biochemical pathways database for Trypanosoma brucei. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, D637-44	20.1	28
257	Parasite Glycobiology: A Bittersweet Symphony. <i>PLoS Pathogens</i> , <b>2015</b> , 11, e1005169	7.6	32
256	Depletion of UDP-Glucose and UDP-Galactose Using a Degron System Leads to Growth Cessation of Leishmania major. <i>PLoS Neglected Tropical Diseases</i> , <b>2015</b> , 9, e0004205	4.8	14
255	Serum proteome of nonalcoholic fatty liver disease: a multimodal approach to discovery of biomarkers of nonalcoholic steatohepatitis. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , <b>2014</b> , 29, 1839-47	4	29
254	Lead optimization of a pyrazole sulfonamide series of Trypanosoma brucei N-myristoyltransferase inhibitors: identification and evaluation of CNS penetrant compounds as potential treatments for stage 2 human African trypanosomiasis. <i>Journal of Medicinal Chemistry</i> , <b>2014</b> , 57, 9855-69	8.3	42
253	Probing the substrate specificity of Trypanosoma brucei GlcNAc-PI de-N-acetylase with synthetic substrate analogues. <i>Organic and Biomolecular Chemistry</i> , <b>2014</b> , 12, 1919-34	3.9	6
252	High-confidence glycosome proteome for procyclic form Trypanosoma brucei by epitope-tag organelle enrichment and SILAC proteomics. <i>Journal of Proteome Research</i> , <b>2014</b> , 13, 2796-806	5.6	72
251	TbGT8 is a bifunctional glycosyltransferase that elaborates N-linked glycans on a protein phosphatase AcP115 and a GPI-anchor modifying glycan in Trypanosoma brucei. <i>Parasitology International</i> , <b>2014</b> , 63, 513-8	2.1	7
250	Fragment screening reveals salicylic hydroxamic acid as an inhibitor of Trypanosoma brucei GPI GlcNAc-PI de-N-acetylase. <i>Carbohydrate Research</i> , <b>2014</b> , 387, 54-8	2.9	10
249	Evaluation of the diagnostic accuracy of prototype rapid tests for human African trypanosomiasis. <i>PLoS Neglected Tropical Diseases</i> , <b>2014</b> , 8, e3373	4.8	29
248	Proteomic selection of immunodiagnostic antigens for Trypanosoma congolense. <i>PLoS Neglected Tropical Diseases</i> , <b>2014</b> , 8, e2936	4.8	8
247	Identification of sVSG117 as an immunodiagnostic antigen and evaluation of a dual-antigen lateral flow test for the diagnosis of human African trypanosomiasis. <i>PLoS Neglected Tropical Diseases</i> , <b>2014</b> , 8, e2976	4.8	12
246	Identification and functional characterization of a highly divergent N-acetylglucosaminyltransferase I (TbGnTI) in Trypanosoma brucei. <i>Journal of Biological Chemistry</i> , <b>2014</b> , 289, 9328-39	5.4	17
245	Global quantitative SILAC phosphoproteomics reveals differential phosphorylation is widespread between the procyclic and bloodstream form lifecycle stages of Trypanosoma brucei. <i>Journal of Proteome Research</i> , <b>2013</b> , 12, 2233-44	5.6	139
244	Genetic and structural validation of Aspergillus fumigatus UDP-N-acetylglucosamine pyrophosphorylase as an antifungal target. <i>Molecular Microbiology</i> , <b>2013</b> , 89, 479-93	4.1	17
243	A novel allosteric inhibitor of the uridine diphosphate N-acetylglucosamine pyrophosphorylase from Trypanosoma brucei. <i>ACS Chemical Biology</i> , <b>2013</b> , 8, 1981-7	4.9	22
242	Proteomic selection of immunodiagnostic antigens for human African trypanosomiasis and generation of a prototype lateral flow immunodiagnostic device. <i>PLoS Neglected Tropical Diseases</i> , <b>2013</b> , 7, e2087	4.8	35
241	Exploring the Trypanosoma brucei Hsp83 potential as a target for structure guided drug design. <i>PLoS Neglected Tropical Diseases</i> , <b>2013</b> , 7, e2492	4.8	25

## (2011-2013)

240	Genetic and structural validation of Aspergillus fumigatus N-acetylphosphoglucosamine mutase as an antifungal target. <i>Bioscience Reports</i> , <b>2013</b> , 33,	4.1	15	
239	Structure of a complex phosphoglycan epitope from gp72 of Trypanosoma cruzi. <i>Journal of Biological Chemistry</i> , <b>2013</b> , 288, 11093-105	5.4	19	
238	Creation and characterization of glycosyltransferase mutants of Trypanosoma brucei. <i>Methods in Molecular Biology</i> , <b>2013</b> , 1022, 249-75	1.4	6	
237	Phosphoglucomutase is absent in Trypanosoma brucei and redundantly substituted by phosphomannomutase and phospho-N-acetylglucosamine mutase. <i>Molecular Microbiology</i> , <b>2012</b> , 85, 513-34	4.1	19	
236	Chemical proteomic analysis reveals the drugability of the kinome of Trypanosoma brucei. <i>ACS Chemical Biology</i> , <b>2012</b> , 7, 1858-65	4.9	47	
235	The de novo and salvage pathways of GDP-mannose biosynthesis are both sufficient for the growth of bloodstream-form Trypanosoma brucei. <i>Molecular Microbiology</i> , <b>2012</b> , 84, 340-51	4.1	20	
234	Discovery of a novel class of orally active trypanocidal N-myristoyltransferase inhibitors. <i>Journal of Medicinal Chemistry</i> , <b>2012</b> , 55, 140-52	8.3	88	
233	Comparative SILAC proteomic analysis of Trypanosoma brucei bloodstream and procyclic lifecycle stages. <i>PLoS ONE</i> , <b>2012</b> , 7, e36619	3.7	122	
232	Inhibitors incorporating zinc-binding groups target the GlcNAc-PI de-N-acetylase in Trypanosoma brucei, the causative agent of African sleeping sickness. <i>Chemical Biology and Drug Design</i> , <b>2012</b> , 79, 27	70 <sup>2</sup> 8 <sup>9</sup>	7	
231	Modeling of the N-glycosylated transferrin receptor suggests how transferrin binding can occur within the surface coat of Trypanosoma brucei. <i>PLoS Pathogens</i> , <b>2012</b> , 8, e1002618	7.6	34	
230	The lipid-linked oligosaccharide donor specificities of Trypanosoma brucei oligosaccharyltransferases. <i>Glycobiology</i> , <b>2012</b> , 22, 696-703	5.8	18	
229	Deep evolutionary conservation of an intramolecular protein kinase activation mechanism. <i>PLoS ONE</i> , <b>2012</b> , 7, e29702	3.7	17	
228	Systematic review of performance of non-invasive biomarkers in the evaluation of non-alcoholic fatty liver disease. <i>Liver International</i> , <b>2011</b> , 31, 461-73	7.9	32	
227	. Tetrahedron Letters, <b>2011</b> , 52, 7091-7094	2	10	
226	Probing elongating and branching ED-galactosyltransferase activities in Leishmania parasites by making use of synthetic phosphoglycans. <i>ACS Chemical Biology</i> , <b>2011</b> , 6, 648-57	4.9	5	
225	Synthesis of potential metal-binding group compounds to examine the zinc dependency of the GPI de-N-acetylase metalloenzyme in Trypanosoma brucei. <i>Carbohydrate Research</i> , <b>2011</b> , 346, 708-14	2.9	7	
224	Protein O-GlcNAcylation is required for fibroblast growth factor signaling in Drosophila. <i>Science Signaling</i> , <b>2011</b> , 4, ra89	8.8	21	
223	Characterization, localization, essentiality, and high-resolution crystal structure of glucosamine 6-phosphate N-acetyltransferase from Trypanosoma brucei. <i>Eukaryotic Cell</i> , <b>2011</b> , 10, 985-97		27	

222	Chemical structure of Trichomonas vaginalis surface lipoglycan: a role for short galactose (1-4/3) N-acetylglucosamine repeats in host cell interaction. <i>Journal of Biological Chemistry</i> , <b>2011</b> , 286, 40494-	50 <sup>5</sup> 8 <sup>4</sup>	33
221	N-myristoyltransferase inhibitors as new leads to treat sleeping sickness. <i>Nature</i> , <b>2010</b> , 464, 728-32	50.4	213
220	Identification, subcellular localization, biochemical properties, and high-resolution crystal structure of Trypanosoma brucei UDP-glucose pyrophosphorylase. <i>Glycobiology</i> , <b>2010</b> , 20, 1619-30	5.8	25
219	Application of electrospray mass spectrometry to the structural determination of glycosylphosphatidylinositol membrane anchors. <i>Glycobiology</i> , <b>2010</b> , 20, 576-85	5.8	8
218	Prophossi: automating expert validation of phosphopeptide-spectrum matches from tandem mass spectrometry. <i>Bioinformatics</i> , <b>2010</b> , 26, 2153-9	7.2	17
217	A multidimensional strategy to detect polypharmacological targets in the absence of structural and sequence homology. <i>PLoS Computational Biology</i> , <b>2010</b> , 6, e1000648	5	61
216	Stoichiometric quantification of Akt phosphorylation using LC-MS/MS. <i>Journal of Proteome Research</i> , <b>2010</b> , 9, 743-51	5.6	21
215	Computer-aided identification of Trypanosoma brucei uridine diphosphate galactose 4Fepimerase inhibitors: toward the development of novel therapies for African sleeping sickness. <i>Journal of Medicinal Chemistry</i> , <b>2010</b> , 53, 5025-32	8.3	45
214	Myristoyl-CoA:protein N-myristoyltransferase depletion in trypanosomes causes avirulence and endocytic defects. <i>Molecular and Biochemical Parasitology</i> , <b>2010</b> , 169, 55-8	1.9	44
213	Glycotyping of Trypanosoma brucei variant surface glycoprotein MITat1.8. <i>Molecular and Biochemical Parasitology</i> , <b>2010</b> , 174, 74-7	1.9	13
212	Trypanosoma brucei UDP-glucose:glycoprotein glucosyltransferase has unusual substrate specificity and protects the parasite from stress. <i>Eukaryotic Cell</i> , <b>2009</b> , 8, 230-40		40
211	Fate of glycosylphosphatidylinositol (GPI)-less procyclin and characterization of sialylated non-GPI-anchored surface coat molecules of procyclic-form Trypanosoma brucei. <i>Eukaryotic Cell</i> , <b>2009</b> , 8, 1407-17		21
210	The phosphoproteome of bloodstream form Trypanosoma brucei, causative agent of African sleeping sickness. <i>Molecular and Cellular Proteomics</i> , <b>2009</b> , 8, 1527-38	7.6	129
209	Chapter 3 The GlcNAc-PI de-N-acetylase. <i>The Enzymes</i> , <b>2009</b> , 49-64	2.3	4
208	Identification and specific localization of tyrosine-phosphorylated proteins in Trypanosoma brucei. <i>Eukaryotic Cell</i> , <b>2009</b> , 8, 617-26		33
207	Proteomic scale high-sensitivity analyses of GPI membrane anchors. <i>Glycoconjugate Journal</i> , <b>2009</b> , 26, 915-21	3	8
206	Identification of a glycosylphosphatidylinositol anchor-modifying beta1-3 N-acetylglucosaminyl transferase in Trypanosoma brucei. <i>Molecular Microbiology</i> , <b>2009</b> , 71, 478-91	4.1	30
205	Distinct donor and acceptor specificities of Trypanosoma brucei oligosaccharyltransferases. <i>EMBO Journal</i> , <b>2009</b> , 28, 2650-61	13	87

### (2006-2008)

204	Probing enzymes late in the trypanosomal glycosylphosphatidylinositol biosynthetic pathway with synthetic glycosylphosphatidylinositol analogues. <i>ACS Chemical Biology</i> , <b>2008</b> , 3, 625-34	4.9	26
203	The synthesis of UDP-N-acetylglucosamine is essential for bloodstream form trypanosoma brucei in vitro and in vivo and UDP-N-acetylglucosamine starvation reveals a hierarchy in parasite protein glycosylation. <i>Journal of Biological Chemistry</i> , <b>2008</b> , 283, 16147-61	5.4	46
202	Phosphatidylethanolamine in Trypanosoma brucei is organized in two separate pools and is synthesized exclusively by the Kennedy pathway. <i>Journal of Biological Chemistry</i> , <b>2008</b> , 283, 23636-44	5.4	44
201	Deletion of the TbALG3 gene demonstrates site-specific N-glycosylation and N-glycan processing in Trypanosoma brucei. <i>Glycobiology</i> , <b>2008</b> , 18, 367-83	5.8	54
200	Synthesis of 1-D-6-O-[2-(N-hydroxyaminocarbonyl)amino-2-deoxy-alpha-D-glucopyranosyl]-myo-inositol 1-(n-octadecyl phosphate): a potential metalloenzyme inhibitor of glycosylphosphatidylinositol biosynthesis. Carbohydrate Research, 2008, 343, 1478-81	2.9	6
199	Probing Trypanosoma brucei glycosylphosphatidylinositol biosynthesis using novel precursor-analogues. <i>Chemical Biology and Drug Design</i> , <b>2008</b> , 72, 127-32	2.9	8
198	Structure of the glycosylphosphatidylinositol anchor of the Trypanosoma brucei transferrin receptor. <i>Molecular and Biochemical Parasitology</i> , <b>2007</b> , 151, 220-3	1.9	17
197	The Chemical Synthesis of Glycosylphosphatidylinositol Anchors from Trypanosoma cruzi Trypomastigote Mucins. <i>ACS Symposium Series</i> , <b>2007</b> , 285-306	0.4	2
196	The de novo synthesis of GDP-fucose is essential for flagellar adhesion and cell growth in Trypanosoma brucei. <i>Journal of Biological Chemistry</i> , <b>2007</b> , 282, 28853-28863	5.4	37
195	Sugar nucleotide pools of Trypanosoma brucei, Trypanosoma cruzi, and Leishmania major. <i>Eukaryotic Cell</i> , <b>2007</b> , 6, 1450-63		109
195 194		1.9	109 45
	Eukaryotic Cell, 2007, 6, 1450-63  The suppression of galactose metabolism in Trypanosoma cruzi epimastigotes causes changes in cell surface molecular architecture and cell morphology. Molecular and Biochemical Parasitology,	1.9	
194	The suppression of galactose metabolism in Trypanosoma cruzi epimastigotes causes changes in cell surface molecular architecture and cell morphology. <i>Molecular and Biochemical Parasitology</i> , <b>2006</b> , 147, 126-36  Characterization of the glycosylphosphatidylinositol anchor of the immunodominant		45
194	The suppression of galactose metabolism in Trypanosoma cruzi epimastigotes causes changes in cell surface molecular architecture and cell morphology. <i>Molecular and Biochemical Parasitology</i> , <b>2006</b> , 147, 126-36  Characterization of the glycosylphosphatidylinositol anchor of the immunodominant Cryptosporidium parvum 17-kDa antigen. <i>Molecular and Biochemical Parasitology</i> , <b>2006</b> , 149, 108-12  The chemical synthesis of bioactive glycosylphosphatidylinositols from Trypanosoma cruzi containing an unsaturated fatty acid in the lipid. <i>Angewandte Chemie - International Edition</i> , <b>2006</b> ,	1.9	45
194 193 192	The suppression of galactose metabolism in Trypanosoma cruzi epimastigotes causes changes in cell surface molecular architecture and cell morphology. <i>Molecular and Biochemical Parasitology</i> , 2006, 147, 126-36  Characterization of the glycosylphosphatidylinositol anchor of the immunodominant Cryptosporidium parvum 17-kDa antigen. <i>Molecular and Biochemical Parasitology</i> , 2006, 149, 108-12  The chemical synthesis of bioactive glycosylphosphatidylinositols from Trypanosoma cruzi containing an unsaturated fatty acid in the lipid. <i>Angewandte Chemie - International Edition</i> , 2006, 45, 468-74  Outer chain N-glycans are required for cell wall integrity and virulence of Candida albicans. <i>Journal</i>	1.9	45 13 49
194 193 192	Eukaryotic Cell, 2007, 6, 1450-63  The suppression of galactose metabolism in Trypanosoma cruzi epimastigotes causes changes in cell surface molecular architecture and cell morphology. Molecular and Biochemical Parasitology, 2006, 147, 126-36  Characterization of the glycosylphosphatidylinositol anchor of the immunodominant Cryptosporidium parvum 17-kDa antigen. Molecular and Biochemical Parasitology, 2006, 149, 108-12  The chemical synthesis of bioactive glycosylphosphatidylinositols from Trypanosoma cruzi containing an unsaturated fatty acid in the lipid. Angewandte Chemie - International Edition, 2006, 45, 468-74  Outer chain N-glycans are required for cell wall integrity and virulence of Candida albicans. Journal of Biological Chemistry, 2006, 281, 90-8  Synthetic glycovaccine protects against the bite of leishmania-infected sand flies. Journal of	1.9 16.4 5.4	45 13 49 182
194 193 192 191	The suppression of galactose metabolism in Trypanosoma cruzi epimastigotes causes changes in cell surface molecular architecture and cell morphology. <i>Molecular and Biochemical Parasitology</i> , 2006, 147, 126-36  Characterization of the glycosylphosphatidylinositol anchor of the immunodominant Cryptosporidium parvum 17-kDa antigen. <i>Molecular and Biochemical Parasitology</i> , 2006, 149, 108-12  The chemical synthesis of bioactive glycosylphosphatidylinositols from Trypanosoma cruzi containing an unsaturated fatty acid in the lipid. <i>Angewandte Chemie - International Edition</i> , 2006, 45, 468-74  Outer chain N-glycans are required for cell wall integrity and virulence of Candida albicans. <i>Journal of Biological Chemistry</i> , 2006, 281, 90-8  Synthetic glycovaccine protects against the bite of leishmania-infected sand flies. <i>Journal of Infectious Diseases</i> , 2006, 194, 512-8	1.9 16.4 5.4	45 13 49 182 48

186	Recombinant human PPAR-beta/delta ligand-binding domain is locked in an activated conformation by endogenous fatty acids. <i>Journal of Molecular Biology</i> , <b>2006</b> , 356, 1005-13	6.5	73
185	Reevaluation of the PPAR-beta/delta ligand binding domain model reveals why it exhibits the activated form. <i>Molecular Cell</i> , <b>2006</b> , 21, 1-2	17.6	50
184	Trypanosoma brucei UDP-galactose-4Fepimerase in ternary complex with NAD+ and the substrate analogue UDP-4-deoxy-4-fluoro-alpha-D-galactose. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , <b>2006</b> , 62, 829-34		15
183	Hypomorphic promoter mutation in PIGM causes inherited glycosylphosphatidylinositol deficiency. <i>Nature Medicine</i> , <b>2006</b> , 12, 846-51	50.5	164
182	Identification of novel inhibitors of UDP-Glc 4Tepimerase, a validated drug target for african sleeping sickness. <i>Bioorganic and Medicinal Chemistry Letters</i> , <b>2006</b> , 16, 5744-7	2.9	32
181	The Chemical Synthesis of Bioactive Glycosylphosphatidylinositols from Trypanosoma cruzi Containing an Unsaturated Fatty Acid in the Lipid. <i>Angewandte Chemie</i> , <b>2006</b> , 118, 482-488	3.6	8
180	The proteome of Trypanosoma cruzi shed vesicles involved in host immunomodulation and cell invasion. <i>FASEB Journal</i> , <b>2006</b> , 20, A514	0.9	O
179	A robust and selective method for the quantification of glycosylphosphatidylinositols in biological samples. <i>Glycobiology</i> , <b>2005</b> , 15, 131-8	5.8	17
178	Synthesis of a cell-permeable analogue of a glycosylphosphatidylinositol (GPI) intermediate that is toxic to the living bloodstream form of Trypanosoma brucei. <i>Tetrahedron Letters</i> , <b>2005</b> , 46, 7419-7421	2	7
177	Mannosamine can replace glucosamine in glycosylphosphatidylinositols of Plasmodium falciparum in vitro. <i>Molecular and Biochemical Parasitology</i> , <b>2005</b> , 142, 12-24	1.9	4
176	Synthetic fragments of antigenic lipophosphoglycans from Leishmania major and Leishmania mexicana and their use for characterisation of the Leishmania elongating alpha-D-mannopyranosylphosphate transferase. <i>Chemistry - A European Journal</i> , <b>2005</b> , 11, 2019-30	4.8	24
175	The identification of isoprenoids that bind in the intersubunit cavity of Escherichia coli 2C-methyl-D-erythritol-2,4-cyclodiphosphate synthase by complementary biophysical methods. <i>Acta Crystallographica Section D: Biological Crystallography</i> , <b>2005</b> , 61, 45-52		36
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44	Glycoinositol-phospholipid profiles of four serotypically distinct Old World Leishmania strains. <i>Biochemical Journal</i> , <b>1994</b> , 304 ( Pt 2), 603-9	3.8	31
43	Parasite glycoconjugates. Part 1. The synthesis of some early and related intermediates in the biosynthetic pathway of glycosyl-phosphatidylinositol membrane anchors. <i>Journal of the Chemical Society Perkin Transactions 1</i> , <b>1993</b> , 2945		34

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41	The structure, biosynthesis and function of glycosylated phosphatidylinositols in the parasitic protozoa and higher eukaryotes. <i>Biochemical Journal</i> , <b>1993</b> , 294 (Pt 2), 305-24	3.8	824
40	A simple purification of procyclic acidic repetitive protein and demonstration of a sialylated glycosyl-phosphatidylinositol membrane anchor. <i>Biochemical Journal</i> , <b>1993</b> , 291 ( Pt 1), 51-5	3.8	108
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32	Mucin-like glycoproteins linked to the membrane by glycosylphosphatidylinositol anchor are the major acceptors of sialic acid in a reaction catalyzed by trans-sialidase in metacyclic forms of Trypanosoma cruzi. <i>Molecular and Biochemical Parasitology</i> , <b>1993</b> , 59, 293-303	1.9	182
31	Biosynthesis of Glycosyl-Phosphatidylinositol Protein Anchors in African Trypanosomes <b>1993</b> , 275-286		1
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28	Inhibition of the GlcNAc transferase of the glycosylphosphatidylinositol anchor biosynthesis in African trypanosomes. <i>FEBS Journal</i> , <b>1992</b> , 208, 309-14		40
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26	Structural characterization of a novel glycosyl-phosphatidylinositol from the protozoan Tetrahymena mimbres. <i>Biochemical Journal</i> , <b>1991</b> , 279 ( Pt 2), 605-8	3.8	12
25	Evolutionary aspects of GPI metabolism in kinetoplastid parasites. <i>Cell Biology International Reports</i> , <b>1991</b> , 15, 991-1005		15

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22	Glycosyl-Phosphatidylinositol Membrane Anchors <b>1991</b> , 331-348		
21	Characterisation of the asparagine-linked oligosaccharides from Trypanosoma brucei type-I variant surface glycoproteins. <i>FEBS Journal</i> , <b>1990</b> , 187, 657-63		56
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19	Characterization of the cross-reacting determinant (CRD) of the glycosyl-phosphatidylinositol membrane anchor of Trypanosoma brucei variant surface glycoprotein. <i>FEBS Journal</i> , <b>1988</b> , 176, 527-34		129
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1	A broadly active fucosyltransferase LmjFUT1 whose mitochondrial localization and catalytic activity is essential in the parasitic protozoan Leishmania		1