Lucy Glover

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcription Dependent Loss of an Ectopically Expressed Variant Surface Glycoprotein during Antigenic Variation in Trypanosoma brucei. MBio, 2022, 13, e0384721.	4.1	1
2	VEX1 Influences mVSG Expression During the Transition to Mammalian Infectivity in Trypanosoma brucei. Frontiers in Cell and Developmental Biology, 2022, 10, 851475.	3.7	2
3	The MRN complex promotes DNA repair by homologous recombination and restrains antigenic variation in African trypanosomes. Nucleic Acids Research, 2021, 49, 1436-1454.	14.5	11
4	Spatial integration of transcription and splicing in a dedicated compartment sustains monogenic antigen expression in African trypanosomes. Nature Microbiology, 2021, 6, 289-300.	13.3	50
5	DNA double strand break position leads to distinct gene expression changes and regulates VSG switching pathway choice. PLoS Pathogens, 2021, 17, e1010038.	4.7	6
6	mSphere of Influence: Expanding the CRISPR Sphere with Single-Locus Proteomics. MSphere, 2020, 5, .	2.9	3
7	Forward Genetics in African Trypanosomes. Methods in Molecular Biology, 2020, 2116, 339-352.	0.9	0
8	Monoallelic expression and epigenetic inheritance sustained by a Trypanosoma brucei variant surface glycoprotein exclusion complex. Nature Communications, 2019, 10, 3023.	12.8	73
9	Persistent DNA Damage Foci and DNA Replication with a Broken Chromosome in the African Trypanosome. MBio, 2019, 10, .	4.1	24
10	A post-transcriptional respiratome regulon in trypanosomes. Nucleic Acids Research, 2019, 47, 7063-7077.	14.5	14
11	CRISPR in Parasitology: Not Exactly Cut and Dried!. Trends in Parasitology, 2019, 35, 409-422.	3.3	43
12	Escaping the immune system by DNA repair and recombination in African trypanosomes. Open Biology, 2019, 9, 190182.	3.6	22
13	Insights into antitrypanosomal drug mode-of-action from cytology-based profiling. PLoS Neglected Tropical Diseases, 2018, 12, e0006980.	3.0	41
14	Genome-wide RNAi selection identifies a regulator of transmission stage-enriched gene families and cell-type differentiation in Trypanosoma brucei. PLoS Pathogens, 2017, 13, e1006279.	4.7	30
15	High-resolution analysis of multi-copy variant surface glycoprotein gene expression sites in African trypanosomes. BMC Genomics, 2016, 17, 806.	2.8	23
16	VEX1 controls the allelic exclusion required for antigenic variation in trypanosomes. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 7225-7230.	7.1	90
17	Genome-scale RNAi screens for high-throughput phenotyping in bloodstream-form African trypanosomes. Nature Protocols, 2015, 10, 106-133.	12.0	49
18	Locus-specific control of DNA resection and suppression of subtelomeric VSG recombination by HAT3 in the African trypanosome. Nucleic Acids Research, 2014, 42, 12600-12613.	14.5	22

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19	PPL2 Translesion Polymerase Is Essential for the Completion of Chromosomal DNA Replication in the African Trypanosome. Molecular Cell, 2013, 52, 554-565.	9.7	54
20	Antigenic variation in A frican trypanosomes: the importance of chromosomal and nuclear context in VSG expression control. Cellular Microbiology, 2013, 15, 1984-1993.	2.1	55
21	DNA Break Site at Fragile Subtelomeres Determines Probability and Mechanism of Antigenic Variation in African Trypanosomes. PLoS Pathogens, 2013, 9, e1003260.	4.7	92
22	Aquaglyceroporin 2 controls susceptibility to melarsoprol and pentamidine in African trypanosomes. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 10996-11001.	7.1	134
23	High-throughput decoding of antitrypanosomal drug efficacy and resistance. Nature, 2012, 482, 232-236.	27.8	276
24	Trypanosomal histone γH2A and the DNA damage response. Molecular and Biochemical Parasitology, 2012, 183, 78-83.	1.1	94
25	Microhomology-mediated deletion and gene conversion in African trypanosomes. Nucleic Acids Research, 2011, 39, 1372-1380.	14.5	68
26	High-throughput phenotyping using parallel sequencing of RNA interference targets in the African trypanosome. Genome Research, 2011, 21, 915-924.	5.5	404
27	Site-specific DNA double-strand breaks greatly increase stable transformation efficiency in Trypanosoma brucei. Molecular and Biochemical Parasitology, 2009, 166, 194-197.	1.1	38
28	Sequence homology and microhomology dominate chromosomal double-strand break repair in African trypanosomes. Nucleic Acids Research, 2008, 36, 2608-2618.	14.5	103
29	Deletion of a trypanosome telomere leads to loss of silencing and progressive loss of terminal DNA in the absence of cell cycle arrest. Nucleic Acids Research, 2007, 35, 872-880.	14.5	45
30	Repression of polymerase lâ€mediated gene expression at Trypanosoma brucei telomeres. EMBO Reports, 2006, 7, 93-99.	4.5	46
31	Tagging a T. brucei RRNA locus improves stable transfection efficiency and circumvents inducible expression position effects. Molecular and Biochemical Parasitology, 2005, 144, 142-148.	1.1	135