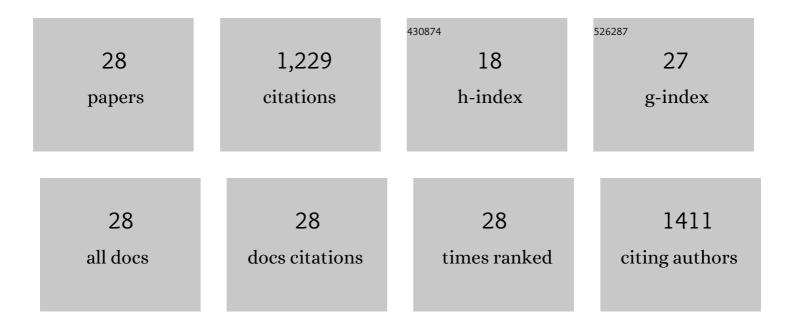
Theodore Morgan

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Cold adaptation does not alter ATP homeostasis during cold exposure in <i>Drosophila melanogaster</i> . Integrative Zoology, 2018, 13, 471-481.	2.6	7
2	Costs of cold acclimation on survival and reproductive behavior in Drosophila melanogaster. PLoS ONE, 2018, 13, e0197822.	2.5	20
3	Speed of exposure to rapid cold hardening and genotype drive the level of acclimation response in Drosophila melanogaster. Journal of Thermal Biology, 2018, 76, 21-28.	2.5	8
4	Genetic Decoupling of Thermal Hardiness across Metamorphosis in Drosophila melanogaster. Integrative and Comparative Biology, 2017, 57, 999-1009.	2.0	37
5	Seasonal variation in basal and plastic cold tolerance: Adaptation is influenced by both long―and shortâ€ŧerm phenotypic plasticity. Ecology and Evolution, 2017, 7, 5248-5257.	1.9	33
6	Adaptation to Low Temperature Exposure Increases Metabolic Rates Independently of Growth Rates. Integrative and Comparative Biology, 2016, 56, 62-72.	2.0	39
7	Artificial selection on chill-coma recovery time in Drosophila melanogaster: Direct and correlated responses to selection. Journal of Thermal Biology, 2016, 59, 77-85.	2.5	27
8	First Report of Spotted Wing Drosophila, <i>Drosophila suzukii</i> Matsumura (Diptera:) Tj ETQq0 0 0 rgBT /Over	rlock 10 Tf	50 462 Td (
9	Constraints, independence, and evolution of thermal plasticity: Probing genetic architecture of long- and short-term thermal acclimation. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 4399-4404.	7.1	144
10	Ecotypes of an ecologically dominant prairie grass (<i><scp>A</scp>ndropogon gerardii</i>) exhibit genetic divergence across the <scp>U</scp> . <scp>S</scp> . Midwest grasslands' environmental gradient. Molecular Ecology, 2014, 23, 6011-6028.	3.9	50
11	Developmental thermal plasticity among <i><scp>D</scp>rosophila melanogaster</i> populations. Journal of Evolutionary Biology, 2014, 27, 557-564.	1.7	30

11	Journal of Evolutionary Biology, 2014, 27, 557-564.	1.7	30
12	Natural selection drives clinal life history patterns in the perennial sunflower species, Helianthus maximiliani. Molecular Ecology, 2011, 20, 2318-2328.	3.9	64
13	Molecular phylogeny of the subfamily Amphistichinae (Teleostei: Embiotocidae) reveals parallel divergent evolution of red pigmentation in two rapidly evolving lineages of sand-dwelling surfperch. Journal of Fish Biology, 2011, 79, no-no.	1.6	4
14	Genetic variation in heat-stress tolerance among South American Drosophila populations. Genetica, 2011, 139, 1331-1337.	1.1	14
15	Heritable Variation in Garter Snake Color Patterns in Postglacial Populations. PLoS ONE, 2011, 6, e24199.	2.5	7
15 16		2.5 0.9	7 21
	e24199. Genetic variation in senescence marker protein-30 is associated with natural variation in cold		

Evolutionary Ecology, 2010, 24, 631-655.

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19	Quantitative genomics of locomotor behavior in Drosophila melanogaster. Genome Biology, 2007, 8, R172.	9.6	68
20	Pleiotropic fitness effects of the Tre1-Gr5a region in Drosophila melanogaster. Nature Genetics, 2006, 38, 824-829.	21.4	27
21	Quantitative trait loci for thermotolerance phenotypes in Drosophila melanogaster. Heredity, 2006, 96, 232-242.	2.6	191
22	Phenotypic Variation and Natural Selection at Catsup, a Pleiotropic Quantitative Trait Gene in Drosophila. Current Biology, 2006, 16, 912-919.	3.9	92
23	Quantitative Trait Loci for Locomotor Behavior in Drosophila melanogaster. Genetics, 2006, 174, 271-284.	2.9	66
24	Quantitative Genomics of Aggressive Behavior in Drosophila melanogaster. PLoS Genetics, 2006, 2, e154.	3.5	165
25	High-Resolution Mapping of Quantitative Trait Loci Affecting Increased Life Span in Drosophila melanogaster. Genetics, 2006, 173, 1455-1463.	2.9	30
26	Molecular and quantitative genetic divergence among populations of house mice with known evolutionary histories. Heredity, 2005, 94, 518-525.	2.6	22
27	The Mode of Evolution of Molecular Markers in Populations of House Mice Under Artificial Selection for Locomotor Behavior. , 2003, 94, 236-242.		4
28	Molecular analysis of the chromosomal region encoding thenifAandnifBgenes ofAcetobacter diazotrophicus. FEMS Microbiology Letters, 1999, 176, 301-309.	1.8	22