

pmid	title	authors	journal	volume	issue	citation	pub_date	pages	doi	pubc	using_cxcr4	url
191000	How West is the West? Evolution of <i>Salix</i> B. Hultine et al.	Journal of Biogeography	35	1	2010	2010	1-12			TRUE		https://doi.org/10.1111/j.1365-3113.2009.04398.x
191001	Newly identified transmembrane from <i>Mus musculus</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	200-204			TRUE		https://doi.org/10.1016/j.jmb.2009.11.001
191002	Population history of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	205-210			TRUE		https://doi.org/10.1016/j.jmb.2009.11.002
191003	Arise! Invasive species! <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	211-216			TRUE		https://doi.org/10.1016/j.jmb.2009.11.003
191004	Recapitulating the <i>Caenorhabditis elegans</i> <i>Wago-1</i> pathway. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	217-222			TRUE		https://doi.org/10.1016/j.jmb.2009.11.004
191005	Genome-wide association of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	223-228			TRUE		https://doi.org/10.1016/j.jmb.2009.11.005
191006	Comparative genomics of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	229-234			TRUE		https://doi.org/10.1016/j.jmb.2009.11.006
191007	Adult variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	235-240			TRUE		https://doi.org/10.1016/j.jmb.2009.11.007
191008	Shaded genetic diversity in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	241-246			TRUE		https://doi.org/10.1016/j.jmb.2009.11.008
191009	<i>Caenorhabditis elegans</i> <i>Wago-1</i> pathway. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	247-252			TRUE		https://doi.org/10.1016/j.jmb.2009.11.009
191010	Protein-protein interaction network in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	253-258			TRUE		https://doi.org/10.1016/j.jmb.2009.11.010
191011	Linking genes and metabolites in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	259-264			TRUE		https://doi.org/10.1016/j.jmb.2009.11.011
191012	Metabolic pathway analysis in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	265-270			TRUE		https://doi.org/10.1016/j.jmb.2009.11.012
191013	A model for evolutionary ecology of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	271-276			TRUE		https://doi.org/10.1016/j.jmb.2009.11.013
191014	An integrative systems biology of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	277-282			TRUE		https://doi.org/10.1016/j.jmb.2009.11.014
191015	Woodwide 2011: a new look at <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	283-288			TRUE		https://doi.org/10.1016/j.jmb.2009.11.015
191016	Polymorphism and genetic diversity in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	289-294			TRUE		https://doi.org/10.1016/j.jmb.2009.11.016
191017	Why do individuals differ? <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	295-300			TRUE		https://doi.org/10.1016/j.jmb.2009.11.017
191018	Natural variation in the distribution of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	301-306			TRUE		https://doi.org/10.1016/j.jmb.2009.11.018
191019	Regulatory changes in the distribution of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	307-312			TRUE		https://doi.org/10.1016/j.jmb.2009.11.019
191020	Correlations of genotype with climate in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	313-318			TRUE		https://doi.org/10.1016/j.jmb.2009.11.020
191021	Evolutionary divergence of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	319-324			TRUE		https://doi.org/10.1016/j.jmb.2009.11.021
191022	Fast genetic divergence of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	325-330			TRUE		https://doi.org/10.1016/j.jmb.2009.11.022
191023	Long-range dispersal of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	331-336			TRUE		https://doi.org/10.1016/j.jmb.2009.11.023
191024	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	337-342			TRUE		https://doi.org/10.1016/j.jmb.2009.11.024
191025	Change in leaflet morphology in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	343-348			TRUE		https://doi.org/10.1016/j.jmb.2009.11.025
191026	From 'New World' to the 'Old World' <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	349-354			TRUE		https://doi.org/10.1016/j.jmb.2009.11.026
191027	A novel novel of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	355-360			TRUE		https://doi.org/10.1016/j.jmb.2009.11.027
191028	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	361-366			TRUE		https://doi.org/10.1016/j.jmb.2009.11.028
191029	The genetic basis of natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	367-372			TRUE		https://doi.org/10.1016/j.jmb.2009.11.029
191030	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	373-378			TRUE		https://doi.org/10.1016/j.jmb.2009.11.030
191031	A genome-wide association study of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	379-384			TRUE		https://doi.org/10.1016/j.jmb.2009.11.031
191032	The structure of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	385-390			TRUE		https://doi.org/10.1016/j.jmb.2009.11.032
191033	Long-range dispersal of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	391-396			TRUE		https://doi.org/10.1016/j.jmb.2009.11.033
191034	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	397-402			TRUE		https://doi.org/10.1016/j.jmb.2009.11.034
191035	Change in leaflet morphology in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	403-408			TRUE		https://doi.org/10.1016/j.jmb.2009.11.035
191036	From 'New World' to the 'Old World' <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	409-414			TRUE		https://doi.org/10.1016/j.jmb.2009.11.036
191037	A novel novel of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	415-420			TRUE		https://doi.org/10.1016/j.jmb.2009.11.037
191038	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	421-426			TRUE		https://doi.org/10.1016/j.jmb.2009.11.038
191039	The genetic basis of natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	427-432			TRUE		https://doi.org/10.1016/j.jmb.2009.11.039
191040	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	433-438			TRUE		https://doi.org/10.1016/j.jmb.2009.11.040
191041	A genome-wide association study of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	439-444			TRUE		https://doi.org/10.1016/j.jmb.2009.11.041
191042	The structure of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	445-450			TRUE		https://doi.org/10.1016/j.jmb.2009.11.042
191043	Long-range dispersal of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	451-456			TRUE		https://doi.org/10.1016/j.jmb.2009.11.043
191044	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	457-462			TRUE		https://doi.org/10.1016/j.jmb.2009.11.044
191045	Change in leaflet morphology in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	463-468			TRUE		https://doi.org/10.1016/j.jmb.2009.11.045
191046	From 'New World' to the 'Old World' <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	469-474			TRUE		https://doi.org/10.1016/j.jmb.2009.11.046
191047	A novel novel of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	475-480			TRUE		https://doi.org/10.1016/j.jmb.2009.11.047
191048	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	481-486			TRUE		https://doi.org/10.1016/j.jmb.2009.11.048
191049	The genetic basis of natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	487-492			TRUE		https://doi.org/10.1016/j.jmb.2009.11.049
191050	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	493-498			TRUE		https://doi.org/10.1016/j.jmb.2009.11.050
191051	A genome-wide association study of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	499-504			TRUE		https://doi.org/10.1016/j.jmb.2009.11.051
191052	The structure of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	505-510			TRUE		https://doi.org/10.1016/j.jmb.2009.11.052
191053	Long-range dispersal of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	511-516			TRUE		https://doi.org/10.1016/j.jmb.2009.11.053
191054	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	517-522			TRUE		https://doi.org/10.1016/j.jmb.2009.11.054
191055	Change in leaflet morphology in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	523-528			TRUE		https://doi.org/10.1016/j.jmb.2009.11.055
191056	From 'New World' to the 'Old World' <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	529-534			TRUE		https://doi.org/10.1016/j.jmb.2009.11.056
191057	A novel novel of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	535-540			TRUE		https://doi.org/10.1016/j.jmb.2009.11.057
191058	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	541-546			TRUE		https://doi.org/10.1016/j.jmb.2009.11.058
191059	The genetic basis of natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	547-552			TRUE		https://doi.org/10.1016/j.jmb.2009.11.059
191060	Natural variation in <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	553-558			TRUE		https://doi.org/10.1016/j.jmb.2009.11.060
191061	A genome-wide association study of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	559-564			TRUE		https://doi.org/10.1016/j.jmb.2009.11.061
191062	The structure of <i>Salix glauca</i> L. J. Wang et al.	Journal of Molecular Biology	29	19	2010	2010	565-570					

31569401	How Weird is The Worm? Evolution of the Developmental Gene Toolkit in <i>Caenorhabditis elegans</i> .			
27866149	Correlations of Genotype with Climate Parameters Suggest <i>Caenorhabditis elegans</i> Niche Adaptations.			
27893361	Regulatory changes in two chemoreceptor genes contribute to a <i>Caenorhabditis elegans</i> QTL for foraging behavior.			
28486636	Natural Variation in the Distribution and Abundance of Transposable Elements Across the <i>Caenorhabditis elegans</i> Species.			
28495877	A maternal-effect selfish genetic element in <i>Caenorhabditis elegans</i> .			
28700616	Natural variation in a single amino acid substitution underlies physiological responses to topoisomerase II poisons.			
28819099	The genetic basis of natural variation in a phoretic behavior.			
28827289	MIP-MAP: High-Throughput Mapping of <i>Caenorhabditis elegans</i> Temperature-Sensitive Mutants via Molecular Inversion Probes.			
28892780	Natural diversity facilitates the discovery of conserved chemotherapeutic response mechanisms.			
28973976	Why do Individuals Differ in Viral Susceptibility? A Story Told by Model Organisms.			
29066469	Polygenicity and Epistasis Underlie Fitness-Proximal Traits in the <i>Caenorhabditis elegans</i> Multiparental Experimental Evolution (CeMEE) Panel.			
29069413	WormBase 2017: molting into a new stage.			
29199021	<i>An Integrated Systems Genetics and Omics Toolkit to Probe Gene Function.</i>			
29353923	A Model for Evolutionary Ecology of Disease: The Case for <i>Caenorhabditis</i> Nematodes and Their Natural Parasites.			
29603298	<i>Hawaiian Drosophila as an Evolutionary Model Clade: Days of Future Past.</i>			
29779955	Linking Genomic and Metabolomic Natural Variation Uncovers Nematode Pheromone Biosynthesis.			
30078564	Natural Genetic Variation in a Multigenerational Phenotype in <i>C. elegans</i> .			
30086719	Proteomic and evolutionary analyses of sperm activation identify uncharacterized genes in <i>Caenorhabditis</i> nematodes.			
30091255	<i>Caenorhabditis elegans</i> as an emerging model system in environmental epigenetics.			
30171204	An open-source platform for analyzing and sharing worm-behavior data.			
30258230	A natural variant and engineered mutation in a GPCR promote DEET resistance in <i>C. elegans</i> .			
30287515	From "the Worm" to "the Worms" and Back Again: The Evolutionary Developmental Biology of Nematodes.			
30328811	Changes to social feeding behaviors are not sufficient for fitness gains of the <i>Caenorhabditis elegans</i> N2 reference strain.			
30341085	Shared Genomic Regions Underlie Natural Variation in Diverse Toxin Responses.			
30372484	Extreme allelic heterogeneity at a <i>Caenorhabditis elegans</i> beta-tubulin locus explains natural resistance to benzimidazoles.			
30513394	Adult Influence on Juvenile Phenotypes by Stage-Specific Pheromone Production.			
30713076	Comparative Epigenomics Reveals that RNA Polymerase II Pausing and Chromatin Domain Organization Control Nematode piRNA Biogenesis.			
30733094	<i>Genome-wide Approaches to Investigate Anthelmintic Resistance.</i>			
30958264	Natural variation in <i>C. elegans</i> arsenic toxicity is explained by differences in branched chain amino acid metabolism.			
31007946	Comparative genomics of 10 new <i>Caenorhabditis</i> species.			
31123080	Recompleting the <i>Caenorhabditis elegans</i> genome.			
31123081	Long-read sequencing reveals intra-species tolerance of substantial structural variations and new subtelomere formation in <i>C. elegans</i> .			
31171655	A Novel Gene Underlies Bleomycin-Response Variation in <i>Caenorhabditis elegans</i> .			
31213597	Fast genetic mapping of complex traits in <i>C. elegans</i> using millions of individuals in bulk.			
31395653	Natural Variation and Genetic Determinants of <i>Caenorhabditis elegans</i> Sperm Size.			
31406358	<i>A new reference genome sequence for Caenorhabditis elegans?</i>			
31414984	Extensive intraspecies cryptic variation in an ancient embryonic gene regulatory network.			
31444297	Population Selection and Sequencing of <i>Caenorhabditis elegans</i> Wild Isolates Identifies a Region on Chromosome III Affecting Starvation Resistance.			
31548647	Selection and gene flow shape niche-associated variation in pheromone response.			
31552506	De novo genome sequencing and comparative stage-specific transcriptomic analysis of <i>Dirofilaria repens</i> .			
31564490	<i>Newly Identified Nematodes from Mono Lake Exhibit Extreme Arsenic Resistance.</i>			
31636085	Accurate Allele Frequencies from Ultra-low Coverage Pool-Seq Samples in Evolve-and-Resequence Experiments.			
31636939	Tightly linked antagonistic-effect loci underlie polygenic phenotypic variation in <i>C. elegans</i> .			
31642470	WormBase: a modern Model Organism Information Resource.			
31672732	<i>Defects in mating behavior and tail morphology are the primary cause of sterility in Caenorhabditis elegans males at high temperature.</i>			
31704915	A fin-2 mutation affects lethal pathology and lifespan in <i>C. elegans</i> .			
31752671	Transcriptional variation and divergence of host-finding behaviour in <i>Steinernema carpocapsae</i> infective juveniles.			
31757604	Natural Variation in a Dendritic Scaffold Protein Remodels Experience-Dependent Plasticity by Altering Neuropeptide Expression.			
31793880	Deep sampling of Hawaiian <i>Caenorhabditis elegans</i> reveals high genetic diversity and admixture with global populations.			
31825311	Diversification of the <i>Caenorhabditis</i> heat shock response by Helitron transposable elements.			
31960906	WormQTL2: an interactive platform for systems genetics in <i>Caenorhabditis elegans</i> .			
32072668	<i>Interspecific Variation in Nematode Responses to Metals.</i>			
32086852	The evolution of parasite host range in heterogeneous host populations.			
32093691	Unexpected cell type-dependent effects of autophagy on polyglutamine aggregation revealed by natural genetic variation in <i>C. elegans</i> .			
32258041	Evolution and Developmental System Drift in the Endoderm Gene Regulatory Network of <i>Caenorhabditis</i> and Other Nematodes.			
32344661	Genetic Variation in <i>Caenorhabditis elegans</i> Responses to Pathogenic Microbiota.			
32385045	The Gene <i>scb-1</i> Underlies Variation in <i>Caenorhabditis elegans</i> Chemotherapeutic Responses.			
32423919	The Ancestral <i>Caenorhabditis elegans</i> Cuticle Suppresses <i>rol-1</i> .			
32482879	Natural cryptic variation in epigenetic modulation of an embryonic gene regulatory network.			
32489524	Predicting gene essentiality in <i>Caenorhabditis elegans</i> by feature engineering and machine-learning			
32550506	The plant terpenoid carvone is a chemotaxis repellent for <i>C. elegans</i> .			
32589676	The nematode <i>Caenorhabditis elegans</i> and the terrestrial isopod <i>Porcellio scaber</i> likely interact opportunistically.			
32767821	<i>Remembering your enemies: mechanisms of within-generation and multigenerational immune priming in Caenorhabditis elegans.</i>			
32840479	Nongenetic inheritance and multigenerational plasticity in the nematode <i>C. elegans</i> .			
32851977	A broad mutational target explains a fast rate of phenotypic evolution.			
32857789	Natural variation in a glucuronosyltransferase modulates propionate sensitivity in a <i>C. elegans</i> propionic acidemia model.			
32868407	PhenoMIP: High-Throughput Phenotyping of Diverse <i>Caenorhabditis elegans</i> Populations via Molecular Inversion Probes.			
32888477	<i>A Natural Mutational Event Uncovers a Life History Trade-Off via Hormonal Pleiotropy.</i>			
32966209	Stoichiometric interactions explain spindle dynamics and scaling across 100 million years of nematode evolution.			
33005351	<i>Caenorhabditis elegans</i> dauers vary recovery in response to bacteria from natural habitat.			

33175833	Natural variation in the sequestosome-related gene, sqst-5, underlies zinc homeostasis in <i>Caenorhabditis elegans</i>.								
33199611	A microbial metabolite synergizes with endogenous serotonin to trigger <i>C. elegans</i> reproductive behavior.								
33249235	<i>Caenorhabditis elegans</i> in anthelmintic research - Old model, new perspectives.								
33317926	<i>Complementary Approaches with Free-living and Parasitic Nematodes to Understanding Anthelmintic Resistance.</i>								
<p>italics = found through Google Scholar, but not pubmed bold = found on Google Scholar and pubmed red = found on Google scholar, but did not actually cite CeNDR no italics or bold = found on pubmed, but not Google Scholar</p>									

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3422987	From QTL to gene Evans KB, van V Trends in genetics - TIC		2021 Jul 3			TRUE	
3415727	Gaining an understanding Reiss AP, Rankin Journal of neurogenetics		2021 Jun 20	1-13		FALSE	
3401945	Behavioral finger McChernoff-Rouss Molecular system 17	5	2021 May	e10287		FALSE	https://www.ncbi.nlm.nih.gov/pmc/articles/PMC8363630/
3376754	Natural variation Witt J, Rodriguez International journal of		2021 Apr 17	1-8		FALSE	https://www.nature.com/articles/s41467-021-19077-7
3382069	Balancing selection Lee D, Zsuzsanna Nature ecology & evolution	6	2021 Jun	794-807		FALSE	https://www.nature.com/articles/s41467-021-19077-7
3404707	Natural genetic Zhang F, Weich Current biology - CB	12	2021 Jun 21	2803-2818.e9		FALSE	https://www.sciencedirect.com/science/article/pii/S0969596221005937
	Natural variation Zhang G, Moradpour (Bethesda, Md)		2021 May 13			FALSE	https://academic.oup.com/g3/advance-article-abstract/doi/10.1093/g3/abaa016/6275292
3398439	Cryptic genetic Koneva SL, Hirt Nature communications	12	2021 May 31	3283		FALSE	https://www.nature.com/articles/s41467-021-23587-1
	Fast genetic map Huang HY, Wang Scientific reports	11	2021 May 26	11017		FALSE	https://www.nature.com/articles/s41598-021-90190-x
	Revealing the role of Lin Q, Wu Food & function	12	2021 Apr 21	3296-3306		FALSE	https://pubs.rsc.org/en/content/articlelanding/2021/fo/cd00248c
3405984	Large genetic diversity Ma F, Lau CY, Zh Genome biology	13	2021 May 7			FALSE	https://academic.oup.com/gb/advance-article-abstract/doi/10.1093/gb/ggab016/6275292

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Genomic architecture of 5S rDNA cluster and its variations within and between species									
Natural variation in the 11d gene family affects insulin/IGF signaling and starvation resistance									
3450264	Natural variation	Lim J, Kim J, Lee OJ (Bethesda, M D)	12		2021 Dec 8			FALSE	https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3854432/
3451820	Variation in copy number	Haston A, Milga Chemosphere	207	P11	2022 Jan	131883		FALSE	https://www.ncbi.nlm.nih.gov/pmc/articles/PMC8610495/
3483058	Return and Auto	Nasser A, M. S. International Jour	22		2021 Nov 12			FALSE	https://www.ncbi.nlm.nih.gov/pmc/articles/PMC8119500/
34613959	easyFutucin	An Di, Bernardo M, C. PflS one	16		2021	60254293		FALSE	https://journals.jku.at/journals/10.1371/journal.pone.0254293
3388511	Comprehensive	Durham T.J, Dazr Genome research	31		2021 Oct	1950-1969		FALSE	https://genome.cshlp.org/content/31/10/1952.full

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Natural variation in the Irf3 gene family affects insulin/IGF signaling and starvation resistance									
35329571	Isolating Caserof-Gimond C, PoulikMethods in mol 2468			2022	283-292		FALSE	https://link.springer.com/journal/10.1007/978-1-4939-2181-3_15	
35322256	Megawati carni Barker L, PanariCommunications 5	1		2022 Mar 23	253		FALSE	https://www.tandfonline.com/doi/full/10.1080/20918474.2022.2060631	
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35156649	Rapid Isolation of Triton IBC, Sibat Journal of visualized experiments 179			2022 Jan 31			FALSE	https://www.ncbi.nlm.nih.gov/pmc/articles/PMC8857960/	
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35134197	Natural genetic v Andersen EC, Rl Genetics 220	1		2022 Jan 4			FALSE	https://academic.oup.com/genetics/advance-article-abstract/doi/10.1093/genetics/iaab220/1/1/6497728	

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Natural variation in the Irf gene family affects insulin/IGF signaling and starvation resistance									
An atlas of gene expression variation across the <i>Caenorhabditis elegans</i> species									
An anchored experimental design and meta-analysis approach to address batch effects in large-scale metabolomics									
Bergian Strains of <i>C. elegans</i> Revealed Expansion of TCI Elements Expose a Significant Genomic and Fitness Cost									
Accurate detection of structural variation is hard									
Dynamic evolution of recently duplicated genes in <i>Caenorhabditis elegans</i>									
<i>Caenorhabditis remanei</i> colonizes ephemeral resource patches in neotropical forests									
35647500	An automated sgCRISPR TA, Ch3-microPublication	2022				FALSE	https://www.ncbi.nlm.nih.gov/pmc/articles/PMC6934690/		
35536194	Evaluating the pcWidmayer S.J. Ev G3 (Bethesda, Md.)	2022 May 10				FALSE	https://academic.oup.com/iadw/article/127/3/kae114/6581107/egp/9846		
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Metabolic Trapping Fungi and Caenorhabditis elegans as a Model System for Predator-Prey Interactions - Vidal-Cheiz de Utzurum et al (chapter in a book called Fungal Associations)									
3827045	Limitations of the inference of the distribution of fitness effects of new mutations in partially selfing populations with linkage - Daigle A, John P. (published online ahead of print, so should be able to add next line)								
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3848806	Role of UDP-Glycyl Auid MZ, Nucleic Chemical reasen 37	4		2024 Apr 15	590-599	https://pubs.acs.org/doi/10.1021/acs.chemrestv.3c00410			
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