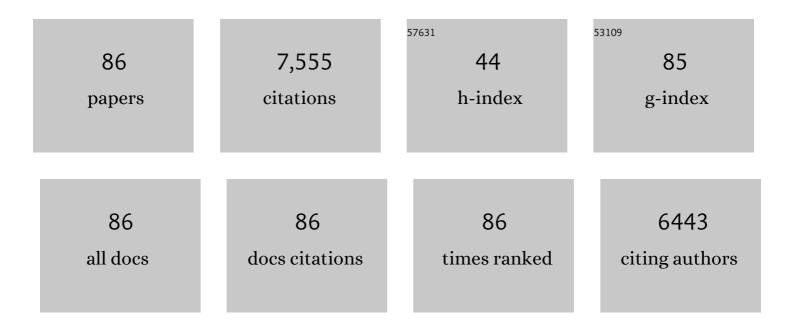
## R Desalle

List of Publications by Year in descending order

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**P DESALLE** 

#	Article	IF	CITATIONS
1	Multiple Sources of Character Information and the Phylogeny of Hawaiian Drosophilids. Systematic Biology, 1997, 46, 654-673.	2.7	533
2	Taking race out of human genetics. Science, 2016, 351, 564-565.	6.0	474
3	Tempo and mode of sequence evolution in mitochondrial DNA of HawaiianDrosophila. Journal of Molecular Evolution, 1987, 26, 157-164.	0.8	463
4	Alignment-Ambiguous Nucleotide Sites and the Exclusion of Systematic Data. Molecular Phylogenetics and Evolution, 1993, 2, 152-157.	1.2	323
5	Patterns of mitochondrial versus nuclear DNA sequence divergence among nymphalid butterflies: the utility of wingless as a source of characters for phylogenetic inference. Insect Molecular Biology, 1998, 7, 73-82.	1.0	272
6	DNA sequences from a fossil termite in Oligo-Miocene amber and their phylogenetic implications. Science, 1992, 257, 1933-1936.	6.0	250
7	Assessing the Relative Contribution of Molecular and Morphological Characters in Simultaneous Analysis Trees. Molecular Phylogenetics and Evolution, 1998, 9, 427-436.	1.2	245
8	Character-based DNA barcoding allows discrimination of genera, species and populations in Odonata. Proceedings of the Royal Society B: Biological Sciences, 2008, 275, 237-247.	1.2	232
9	Class-level relationships in the phylum Cnidaria: molecular and morphological evidence Molecular Biology and Evolution, 1995, 12, 679-89.	3.5	205
10	Nonspecific Adherence by Actinobacillus actinomycetemcomitans Requires Genes Widespread inBacteria and Archaea. Journal of Bacteriology, 2000, 182, 6169-6176.	1.0	194
11	GENEFAMILYEVOLUTION ANDHOMOLOGY: Genomics Meets Phylogenetics. Annual Review of Genomics and Human Genetics, 2000, 1, 41-73.	2.5	193
12	GENE TREES, SPECIES TREES, AND SYSTEMATICS: A Cladistic Perspective. Annual Review of Ecology, Evolution, and Systematics, 1996, 27, 423-450.	6.7	191
13	Phylogeny of genes for secretion NTPases: Identification of the widespread tadA subfamily and development of a diagnostic key for gene classification. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 2503-2508.	3.3	189
14	Molecular Phylogeny of Acipenserinae. Molecular Phylogenetics and Evolution, 1998, 9, 141-155.	1.2	185
15	Molecular evolution of glutamate receptors: a primitive signaling mechanism that existed before plants and animals diverged. Molecular Biology and Evolution, 1999, 16, 826-838.	3.5	185
16	flp-1, the first representative of a new pilin gene subfamily, is required for non-specific adherence of Actinobacillus actinomycetemcomitans. Molecular Microbiology, 2001, 40, 542-554.	1.2	179
17	The Identity of Plant Glutamate Receptors. Science, 2001, 292, 1486b-1487.	6.0	175
18	Elision: A Method for Accommodating Multiple Molecular Sequence Alignments with Alignment-Ambiguous Sites. Molecular Phylogenetics and Evolution, 1995, 4, 1-9.	1.2	145

#	Article	IF	CITATIONS
19	Genes for tight adherence of Actinobacillus actinomycetemcomitans: from plaque to plague to pond scum. Trends in Microbiology, 2001, 9, 429-437.	3.5	135
20	A Cladistic Analysis of Mitochondrial Ribosomal DNA from the Bovidae. Molecular Phylogenetics and Evolution, 1997, 7, 303-319.	1.2	134
21	Discordance of nuclear and mitochondrial DNA phylogenies in Hawaiian Drosophila Proceedings of the United States of America, 1986, 83, 6902-6906.	3.3	111
22	World-wide genetic differentiation ofEubalaena: questioning the number of right whale species. Molecular Ecology, 2000, 9, 1793-1802.	2.0	107
23	Character Congruence of Multiple Data Partitions and the Origin of the Hawaiian Drosophilidae. Molecular Phylogenetics and Evolution, 1998, 9, 225-235.	1.2	101
24	Temporal and Spatial Heterogeneity of mtDNA Polymorphisms in Natural Populations of <i>Drosophila mercatorum</i> . Genetics, 1987, 116, 215-223.	1.2	101
25	Process Partitions, Congruence, and the Independence of Characters: Inferring Relationships among Closely Related Hawaiian Drosophila from Multiple Gene Regions. Systematic Biology, 1997, 46, 751-764.	2.7	100
26	Molluscan engrailed expression, serial organization, and shell evolution. Evolution & Development, 2000, 2, 340-347.	1.1	93
27	The phylogenetic relationships of flies in the family drosophilidae deduced from mtDNA sequences. Molecular Phylogenetics and Evolution, 1992, 1, 31-40.	1.2	90
28	OrthologID: automation of genome-scale ortholog identification within a parsimony framework. Bioinformatics, 2006, 22, 699-707.	1.8	89
29	Calibration of the change in thermal stability of DNA duplexes and degree of base pair mismatch. Journal of Molecular Evolution, 1988, 27, 212-216.	0.8	74
30	Phylogenetic Analysis of the repleta Species Group of the Genus Drosophila Using Multiple Sources of Characters. Molecular Phylogenetics and Evolution, 2000, 16, 296-307.	1.2	72
31	Comparing and combining distanceâ€based and characterâ€based approaches for barcoding turtles. Molecular Ecology Resources, 2011, 11, 956-967.	2.2	72
32	Evolution of MDA-5/RIG-I-dependent innate immunity: Independent evolution by domain grafting. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 17040-17045.	3.3	71
33	Phylogeny of the Bovidae (Artiodactyla, Mammalia), based on mitochondrial ribosomal DNA sequences Molecular Biology and Evolution, 1992, 9, 433-46.	3.5	65
34	PCR jumping in clones of 30-million-year-old DNA fragments from amber preserved termites (Mastotermes electrodominicus). Experientia, 1993, 49, 906-909.	1.2	60
35	Molecular Phylogeny of Acipenseridae: Nonmonophylyof Scaphirhynchinae. Copeia, 2002, 2002, 287-301.	1.4	58
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37	Mitochondrial DNA variability in natural populations of Hawaiian Drosophila. II. Genetic and phylogenetic relationships of natural populations of D. silvestris and D. heteroneura. Heredity, 1986, 56, 87-96.	1.2	56
38	Mitochondrial DNA variability in natural populations of Hawaiian Drosophila. I. Methods and levels of variability in D. silvestris and D. heteroneura populations. Heredity, 1986, 56, 75-85.	1.2	54
39	[14] Isolation and characterization of animal mitochondrial DNA. Methods in Enzymology, 1993, 224, 176-204.	0.4	52
40	THE EVOLUTION AND DEVELOPMENT OF DIPTERAN WING VEINS: A Systematic Approach. Annual Review of Entomology, 1999, 44, 97-129.	5.7	52
41	ON COMBINING PROTEIN SEQUENCES AND NUCLEIC ACID SEQUENCES IN PHYLOGENETIC ANALYSIS: THE HOMEOBOX PROTEIN CASE. Cladistics, 1996, 12, 65-82.	1.5	49
42	Using molecular and ecological data to diagnose endangered populations of the puritan tiger beetle Cicindela puritana. Molecular Ecology, 1993, 2, 375-383.	2.0	48
43	Characteristic attributes in cancer microarrays. Journal of Biomedical Informatics, 2002, 35, 111-122.	2.5	48
44	The origin and possible time of divergence of the Hawaiian Drosophilidae: evidence from DNA sequences Molecular Biology and Evolution, 1992, 9, 905-16.	3.5	46
45	Current problems with the zootype and the early evolution of Hox genes. The Journal of Experimental Zoology, 2001, 291, 169-174.	1.4	43
46	An effective method for isolating DNA from historical specimens of baleen. Molecular Ecology, 1997, 6, 677-681.	2.0	42
47	Homogenization of geographical variants at the nontranscribed spacer of rDNA in Drosophila mercatorum Molecular Biology and Evolution, 1985, 2, 338-46.	3.5	41
48	A Molecular Phylogeny of Costaceae (Zingiberales). Molecular Phylogenetics and Evolution, 2001, 21, 333-345.	1.2	41
49	Molecular genetic analysis among subspecies of two Eurasian sturgeon species, Acipenser baerii and A. stellatus. Molecular Ecology, 1999, 8, S117-S127.	2.0	37
50	Morphological and Molecular Systematics of the Drosophilidae. Annual Review of Ecology, Evolution, and Systematics, 1991, 22, 447-475.	6.7	36
51	The potential of distanceâ€based thresholds and characterâ€based <scp>DNA</scp> barcoding for defining problematic taxonomic entities by <scp>CO</scp> 1 and <scp>ND</scp> 1. Molecular Ecology Resources, 2013, 13, 1069-1081.	2.2	36
52	THE MOLECULAR THROUGH ECOLOGICAL GENETICS OF ABNORMAL ABDOMEN. II. RIBOSOMAL DNA POLYMORPHISM IS ASSOCIATED WITH THE ABNORMAL ABDOMEN SYNDROME IN <i>DROSOPHILA MERCATORUM</i> . Genetics, 1986, 112, 861-875.	1.2	34
53	Homologues of theengrailedgene from five molluscan classes. FEBS Letters, 1995, 365, 71-74.	1.3	32
54	Characterization of microsatellite loci in the endangered St. Vincent Parrot, Amazona guildingii. Molecular Ecology Notes, 2001, 1, 162-164.	1.7	32

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55	Systematic Analysis of DNA Microarray Data: Ordering and Interpreting Patterns of Gene Expression. Genome Research, 2001, 11, 1149-1155.	2.4	31
56	Analysis of Paedomorphosis Using Allometric Characters: The Example of Reduncini Antelopes (Bovidae, Mammalia). Systematic Biology, 1994, 43, 92-116.	2.7	28
57	Evidence of Adaptive Evolution of Accessory Gland Proteins in Closely Related Species of the Drosophila repleta Group. Molecular Biology and Evolution, 2008, 25, 2043-2053.	3.5	28
58	Low <scp>MHC</scp> variation in the polar bear: implications in the face of <scp>A</scp> rctic warming?. Animal Conservation, 2013, 16, 671-683.	1.5	27
59	Molecular evolution in Hawaiian drosophilids. Trends in Ecology and Evolution, 1987, 2, 212-216.	4.2	26
60	Captive breeding, reintroduction, and the conservation genetics of black and white ruffed lemurs, Varecia variegata variegata. Molecular Ecology, 1999, 8, S107-S115.	2.0	26
61	Isolation and characterization of microsatellite loci in Piaractus mesopotamicus and their applicability in other Serrasalminae fish. Molecular Ecology Notes, 2001, 1, 245-247.	1.7	26
62	Polytene chromosomes as indicators of phylogeny in several species groups of Drosophila. BMC Evolutionary Biology, 2001, 1, 6.	3.2	26
63	Implications of ancient DNA for phylogenetic studies. Experientia, 1994, 50, 543-550.	1.2	25
64	Phylogeny and Character Behavior in the Family Lemuridae. Molecular Phylogenetics and Evolution, 2000, 15, 124-134.	1.2	25
65	THE MOLECULAR THROUGH ECOLOGICAL GENETICS OF ABNORMAL ABDOMEN. III. TISSUE-SPECIFIC DIFFERENTIAL REPLICATION OF RIBOSOMAL GENES MODULATES THE ABNORMAL ABDOMEN PHENOTYPE IN <i>DROSOPHILA MERCATORUM</i> . Genetics, 1986, 112, 877-886.	1.2	24
66	The molecular through ecological genetics of abnormal abdomen. IV. Components of genetic variation in a natural population of Drosophila mercatorum Genetics, 1992, 130, 355-366.	1.2	23
67	The mtDNA Genealogy of Closely Related Drosophila silvestris. Journal of Heredity, 1992, 83, 211-216.	1.0	21
68	Phylogenetic Pattern and Developmental Process in Drosophila. Systematic Biology, 1993, 42, 458-475.	2.7	20
69	Speciation and phylogenetic resolution. Trends in Ecology and Evolution, 1994, 9, 297-298.	4.2	20
70	Can we ever identify the Urmetazoan?. Integrative and Comparative Biology, 2007, 47, 670-676.	0.9	20
71	[4] Collection and storage of invertebrate samples. Methods in Enzymology, 1993, 224, 51-65.	0.4	19
72	Characters and the Systematics of Drosophilidae. Journal of Heredity, 1992, 83, 182-188.	1.0	17

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73	Molecular phylogeny of Acipenserinae and black caviar species identification Journal of Applied Ichthyology, 1999, 15, 12-16.	0.3	17
74	Failure to confirm previous identification of two putative museum specimens of the Atlantic sturgeon, Acipenser sturio , as the Adriatic sturgeon, A. naccarii. Marine Biology, 2000, 136, 373-377.	0.7	16
75	DNA Isolation, Manipulation and Characterization from Old Tissues. , 1996, 18, 13-32.		15
76	The Plant Proteome Folding Project: Structure and Positive Selection in Plant Protein Families. Genome Biology and Evolution, 2012, 4, 360-371.	1.1	13
77	Very old DNA. Current Opinion in Genetics and Development, 1994, 4, 810-815.	1.5	12
78	Transformationalism, Taxism, and Developmental Biology in Systematics. Systematic Biology, 2000, 49, 19-27.	2.7	12
79	Computational problems in molecular systematics. Exs, 1994, 69, 353-370.	1.4	9
80	Insect evolution: How the fruit fly changed (some of) its spots. Current Biology, 2000, 10, R75-R77.	1.8	6
81	Hierarchical structure in the Drosophila mojavensis cluster (Diptera: Drosophilidae). Hereditas, 2004, 139, 223-227.	0.5	5
82	Flies and congruence. American Journal of Physical Anthropology, 1994, 94, 125-141.	2.1	4
83	Genetic differentiation and adaptive evolution at reproductive loci in incipient <i>Drosophila</i> species. Journal of Evolutionary Biology, 2017, 30, 524-537.	0.8	4
84	Crossroads, Milestones, amd Landmarks in Insect Development and Evolution: Implications for Systematics. Aliso, 1995, 14, 305-321.	0.4	4
85	Genetic divergence within the Drosophila mayaguana subcluster, a closely related triad of Caribbean species in the repleta species group. Hereditas, 2002, 136, 240-246.	0.5	3
86	The evolution of HOM-C homeoboxes in the Dipteran family Drosophilidae. Insect Molecular Biology, 2003, 12, 345-351.	1.0	1