

Bryan N Danforth

List of Publications by Year in descending order

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Version: 2024-02-01

111
papers

10,465
citations

41344

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docs citations

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times ranked

7797
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Phylogenetic relationships and the evolution of host preferences in the largest clade of brood parasitic bees (Apidae: Nomadinae). <i>Molecular Phylogenetics and Evolution</i> , 2022, 166, 107326. | 2.7 | 11 |
| 2 | Phylogeny, biogeography and diversification of the mining bee family Andrenidae. <i>Systematic Entomology</i> , 2022, 47, 283-302. | 3.9 | 33 |
| 3 | Exosymbiotic microbes within fermented pollen provisions are as important for the development of solitary bees as the pollen itself. <i>Ecology and Evolution</i> , 2022, 12, e8788. | 1.9 | 8 |
| 4 | Climate-driven range shifts of a rare specialist bee, <i>Macropis nuda</i> (Melittidae), and its host plant, <i>Lysimachia ciliata</i> (Primulaceae). <i>Global Ecology and Conservation</i> , 2022, 37, e02180. | 2.1 | 1 |
| 5 | Genome of the bee <i>Holcopasites calliopsidis</i> a species showing the common apid trait of brood parasitism. <i>G3: Genes, Genomes, Genetics</i> , 2022, 12, . | 1.8 | 2 |
| 6 | (More than) Hitchhikers through the network: the shared microbiome of bees and flowers. <i>Current Opinion in Insect Science</i> , 2021, 44, 8-15. | 4.4 | 55 |
| 7 | Bees in the trees: Diverse spring fauna in temperate forest edge canopies. <i>Forest Ecology and Management</i> , 2021, 482, 118903. | 3.2 | 43 |
| 8 | Wild insect diversity increases inter-annual stability in global crop pollinator communities. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20210212. | 2.6 | 43 |
| 9 | Gene Tree Estimation Error with Ultraconserved Elements: An Empirical Study on <i>Pseudapis</i> Bees. <i>Systematic Biology</i> , 2021, 70, 803-821. | 5.6 | 25 |
| 10 | Apple grower pollination practices and perceptions of alternative pollinators in New York and Pennsylvania. <i>Renewable Agriculture and Food Systems</i> , 2020, 35, 1-14. | 1.8 | 32 |
| 11 | Phylogenomic and Morphological Reevaluation of the Bee Tribes Biastini, Neolarrini, and Townsendiellini (Hymenoptera: Apidae) With Description of Three New Species of <i>Schwarzia</i> . <i>Insect Systematics and Diversity</i> , 2020, 4, . | 1.7 | 9 |
| 12 | Towards a U.S. national program for monitoring native bees. <i>Biological Conservation</i> , 2020, 252, 108821. | 4.1 | 54 |
| 13 | Pollen defenses negatively impact foraging and fitness in a generalist bee (<i>Bombus impatiens</i> : Apidae). <i>Scientific Reports</i> , 2020, 10, 3112. | 3.3 | 39 |
| 14 | Sources and frequency of brood loss in solitary bees. <i>Apidologie</i> , 2019, 50, 515-525. | 2.0 | 20 |
| 15 | Omnivory in Bees: Elevated Trophic Positions among All Major Bee Families. <i>American Naturalist</i> , 2019, 194, 414-421. | 2.1 | 47 |
| 16 | The diversification of neopasiphaeine bees during the Cenozoic (Hymenoptera: Colletidae). <i>Zoologica Scripta</i> , 2019, 48, 226-242. | 1.7 | 27 |
| 17 | Agriculturally dominated landscapes reduce bee phylogenetic diversity and pollination services. <i>Science</i> , 2019, 363, 282-284. | 12.6 | 183 |
| 18 | Combining transcriptomes and ultraconserved elements to illuminate the phylogeny of Apidae. <i>Molecular Phylogenetics and Evolution</i> , 2019, 130, 121-131. | 2.7 | 127 |

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|----|--|-----|-----------|
| 19 | Viral transmission in honey bees and native bees, supported by a global black queen cell virus phylogeny. <i>Environmental Microbiology</i> , 2019, 21, 972-983. | 3.8 | 38 |
| 20 | On the universality of targeted enrichment baits for phylogenomic research. <i>Methods in Ecology and Evolution</i> , 2018, 9, 1453-1460. | 5.2 | 51 |
| 21 | Phylogeny, new generic-level classification, and historical biogeography of the <i>Eucera</i> complex (Hymenoptera: Apidae). <i>Molecular Phylogenetics and Evolution</i> , 2018, 119, 81-92. | 2.7 | 55 |
| 22 | Pollinivory and the diversification dynamics of bees. <i>Biology Letters</i> , 2018, 14, 20180530. | 2.3 | 21 |
| 23 | A global synthesis of the effects of diversified farming systems on arthropod diversity within fields and across agricultural landscapes. <i>Global Change Biology</i> , 2017, 23, 4946-4957. | 9.5 | 259 |
| 24 | Flower handling behavior and abundance determine the relative contribution of pollinators to seed set in apple orchards. <i>Agriculture, Ecosystems and Environment</i> , 2017, 246, 102-108. | 5.3 | 38 |
| 25 | Phylogenomic Insights into the Evolution of Stinging Wasps and the Origins of Ants and Bees. <i>Current Biology</i> , 2017, 27, 1019-1025. | 3.9 | 329 |
| 26 | The impact of GC bias on phylogenetic accuracy using targeted enrichment phylogenomic data. <i>Molecular Phylogenetics and Evolution</i> , 2017, 111, 149-157. | 2.7 | 50 |
| 27 | Long-legged bees make adaptive leaps: linking adaptation to coevolution in a plant-pollinator network. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20171707. | 2.6 | 22 |
| 28 | Playing with extremes: Origins and evolution of exaggerated female forelegs in South African Rediviva bees. <i>Molecular Phylogenetics and Evolution</i> , 2017, 115, 95-105. | 2.7 | 10 |
| 29 | Pollen preferences among the bee species visiting apple (<i>Malus pumila</i>) in New York. <i>Apidologie</i> , 2017, 48, 806-820. | 2.0 | 22 |
| 30 | Phylogenetic systematics and a revised generic classification of anthidiine bees (Hymenoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 30 | 2.7 | 34 |
| 31 | Crop domestication facilitated rapid geographical expansion of a specialist pollinator, the squash bee <i>Peponapis pruinosa</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20160443. | 2.6 | 94 |
| 32 | Per-visit pollinator performance and regional importance of wild <i>Bombus</i> and <i>Andrena</i> (<i>Melandrena</i>) compared to the managed honey bee in New York apple orchards. <i>Apidologie</i> , 2016, 47, 145-160. | 2.0 | 56 |
| 33 | Pollination services for apple are dependent on diverse wild bee communities. <i>Agriculture, Ecosystems and Environment</i> , 2016, 221, 1-7. | 5.3 | 121 |
| 34 | The challenge of accurately documenting bee species richness in agroecosystems: bee diversity in eastern apple orchards. <i>Ecology and Evolution</i> , 2015, 5, 3531-3540. | 1.9 | 58 |
| 35 | Nest Suitability, Fine-Scale Population Structure and Male-Mediated Dispersal of a Solitary Ground Nesting Bee in an Urban Landscape. <i>PLoS ONE</i> , 2015, 10, e0125719. | 2.5 | 44 |
| 36 | Introduction of Non-Native Pollinators Can Lead to Trans-Continental Movement of Bee-Associated Fungi. <i>PLoS ONE</i> , 2015, 10, e0130560. | 2.5 | 38 |

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|----|--|------|-----------|
| 37 | Negative effects of pesticides on wild bee communities can be buffered by landscape context. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015, 282, 20150299. | 2.6 | 144 |
| 38 | Delivery of crop pollination services is an insufficient argument for wild pollinator conservation. <i>Nature Communications</i> , 2015, 6, 7414. | 12.8 | 656 |
| 39 | Climate, physiological tolerance and sex-biased dispersal shape genetic structure of Neotropical orchid bees. <i>Molecular Ecology</i> , 2014, 23, 1874-1890. | 3.9 | 62 |
| 40 | Discovery and characterization of microsatellites for the solitary bee <i>Colletes inaequalis</i> using Sanger and 454 pyrosequencing. <i>Apidologie</i> , 2013, 44, 163-172. | 2.0 | 10 |
| 41 | The bee tree of life: a supermatrix approach to apoid phylogeny and biogeography. <i>BMC Evolutionary Biology</i> , 2013, 13, 138. | 3.2 | 134 |
| 42 | Biodiversity ensures plant-pollinator phenological synchrony against climate change. <i>Ecology Letters</i> , 2013, 16, 1331-1338. | 6.4 | 184 |
| 43 | Social Insects: Are Ants Just Wingless Bees?. <i>Current Biology</i> , 2013, 23, R1011-R1012. | 3.9 | 6 |
| 44 | The Impact of Molecular Data on Our Understanding of Bee Phylogeny and Evolution. <i>Annual Review of Entomology</i> , 2013, 58, 57-78. | 11.8 | 200 |
| 45 | A global quantitative synthesis of local and landscape effects on wild bee pollinators in agroecosystems. <i>Ecology Letters</i> , 2013, 16, 584-599. | 6.4 | 875 |
| 46 | ORIGINS, EVOLUTION, AND DIVERSIFICATION OF CLEPTOPARASITIC LINEAGES IN LONG-TONGUED BEES. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, n/a-n/a. | 2.3 | 34 |
| 47 | Historical changes in northeastern US bee pollinators related to shared ecological traits. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 4656-4660. | 7.1 | 432 |
| 48 | Bees diversified in the age of eudicots. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20122686. | 2.6 | 187 |
| 49 | Revision and reclassification of <i>Lasioglossum</i> (<i>Eulyaeus</i>), <i>L.</i> (<i>Hemihalictus</i>) and <i>L.</i> (<i>Sphecodogastra</i>) in eastern North America (Hymenoptera: Apoidea: Halictidae). <i>Zootaxa</i> , 2013, 3672, 1-117. | 0.5 | 114 |
| 50 | Phylogeny of halictine bees supports a shared origin of eusociality for <i>Halictus</i> and <i>Lasioglossum</i> (Apoidea: Anthophila: Halictidae). <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 926-939. | 2.7 | 137 |
| 51 | Taxonomic affinity of halictid bee fossils (Hymenoptera: Anthophila) based on geometric morphometrics analyses of wing shape. <i>Journal of Systematic Palaeontology</i> , 2012, 10, 755-764. | 1.5 | 33 |
| 52 | Biogeography and diversification of colletid bees (Hymenoptera: Colletidae): emerging patterns from the southern end of the world. <i>Journal of Biogeography</i> , 2012, 39, 526-544. | 3.0 | 95 |
| 53 | Phylogeny of the bee family Megachilidae (Hymenoptera: Apoidea) based on adult morphology. <i>Systematic Entomology</i> , 2012, 37, 261-286. | 3.9 | 59 |
| 54 | Identifying the sister group to the bees: a molecular phylogeny of Aculeata with an emphasis on the superfamily Apoidea. <i>Zoologica Scripta</i> , 2012, 41, 527-535. | 1.7 | 63 |

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|----|---|------|-----------|
| 55 | Morphology, Classification, and Antiquity of <i>Melittosphex burmensis</i> (Apoidea: Melittosphecidae) and Implications for Early Bee Evolution. <i>Journal of Paleontology</i> , 2011, 85, 882-891. | 0.8 | 31 |
| 56 | A simple and distinctive microbiota associated with honey bees and bumble bees. <i>Molecular Ecology</i> , 2011, 20, 619-628. | 3.9 | 462 |
| 57 | Rooting phylogenies using gene duplications: An empirical example from the bees (Apoidea). <i>Molecular Phylogenetics and Evolution</i> , 2011, 60, 295-304. | 2.7 | 19 |
| 58 | Isolation and cross-species characterization of polymorphic microsatellites for the orchid bee <i>Eulaema meriana</i> (Hymenoptera: Apidae: Euglossini). <i>Conservation Genetics Resources</i> , 2011, 3, 21-23. | 0.8 | 2 |
| 59 | Why do leafcutter bees cut leaves? New insights into the early evolution of bees. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2011, 278, 3593-3600. | 2.6 | 93 |
| 60 | Climate-associated phenological advances in bee pollinators and bee-pollinated plants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 20645-20649. | 7.1 | 402 |
| 61 | The Antiquity and Evolutionary History of Social Behavior in Bees. <i>PLoS ONE</i> , 2011, 6, e21086. | 2.5 | 161 |
| 62 | Unexpected Polylecty in the Bee Genus <i>Meganomia</i> (Hymenoptera: Apoidea: Melittidae). <i>Journal of the Kansas Entomological Society</i> , 2010, 83, 221-230. | 0.2 | 4 |
| 63 | Comprehensive phylogeny of apid bees reveals the evolutionary origins and antiquity of cleptoparasitism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 16207-16211. | 7.1 | 164 |
| 64 | Phylogeny of the bee family Melittidae (Hymenoptera: Anthophila) based on combined molecular and morphological data. <i>Systematic Entomology</i> , 2009, 34, 574-597. | 3.9 | 55 |
| 65 | Phylogeny of colletid bees (Hymenoptera: Colletidae) inferred from four nuclear genes. <i>Molecular Phylogenetics and Evolution</i> , 2009, 50, 290-309. | 2.7 | 70 |
| 66 | Episodes in insect evolution. <i>Integrative and Comparative Biology</i> , 2009, 49, 590-606. | 2.0 | 57 |
| 67 | Phylogeny and biogeography of bees of the tribe Osmiini (Hymenoptera: Megachilidae). <i>Molecular Phylogenetics and Evolution</i> , 2008, 49, 185-197. | 2.7 | 70 |
| 68 | Phylogeny of the Xeromelissinae (Hymenoptera: Colletidae) Based upon Morphology and Molecules. <i>Apidologie</i> , 2008, 39, 75-85. | 2.0 | 11 |
| 69 | Phylogeny of Halictidae with an emphasis on endemic African Halictinae. <i>Apidologie</i> , 2008, 39, 86-101. | 2.0 | 48 |
| 70 | Insights into Bee Evolution: A Tribute to Charles D. Michener. <i>Apidologie</i> , 2008, 39, 1-2. | 2.0 | 28 |
| 71 | Phylogenetic relationships and host-plant evolution within the basal clade of Halictidae (Hymenoptera, Apoidea). <i>Cladistics</i> , 2008, 24, 255-269. | 3.3 | 30 |
| 72 | Changing Paradigms in Insect Social Evolution: Insights from Halictine and Allodapine Bees. <i>Annual Review of Entomology</i> , 2007, 52, 127-150. | 11.8 | 198 |

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|----|---|------|-----------|
| 73 | Low nuclear DNA variation supports a recent origin of Hawaiian Hylaeus bees (Hymenoptera: Tj ETQq1 1 0.784314 rgBT /Overclock 10 T | 2.7 | 21 |
| 74 | Bees. <i>Current Biology</i> , 2007, 17, R156-R161. | 3.9 | 38 |
| 75 | Recent and simultaneous origins of eusociality in halictid bees. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2006, 273, 1643-1649. | 2.6 | 130 |
| 76 | Evolution and biogeography of native Hawaiian Hylaeus bees (Hymenoptera: Colletidae). <i>Cladistics</i> , 2006, 22, 393-411. | 3.3 | 76 |
| 77 | Analysis of family-level relationships in bees (Hymenoptera: Apiformes) using 28S and two previously unexplored nuclear genes: CAD and RNA polymerase II. <i>Molecular Phylogenetics and Evolution</i> , 2006, 39, 358-372. | 2.7 | 115 |
| 78 | A Fossil Bee from Early Cretaceous Burmese Amber. <i>Science</i> , 2006, 314, 614-614. | 12.6 | 94 |
| 79 | The history of early bee diversification based on five genes plus morphology. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 15118-15123. | 7.1 | 208 |
| 80 | How do insect nuclear ribosomal genes compare to protein-coding genes in phylogenetic utility and nucleotide substitution patterns?. <i>Systematic Entomology</i> , 2005, 30, 549-562. | 3.9 | 51 |
| 81 | Single-Copy Nuclear Genes Recover Cretaceous-Age Divergences in Bees. <i>Systematic Biology</i> , 2004, 53, 309-326. | 5.6 | 130 |
| 82 | Recent Intron Gain in Elongation Factor-1 α of Colletid Bees (Hymenoptera: Colletidae). <i>Molecular Biology and Evolution</i> , 2004, 21, 691-696. | 8.9 | 28 |
| 83 | Molecular Phylogenetics and Evolution of Maternal Care in Membracine Treehoppers. <i>Systematic Biology</i> , 2004, 53, 400-421. | 5.6 | 44 |
| 84 | How do insect nuclear and mitochondrial gene substitution patterns differ? Insights from Bayesian analyses of combined datasets. <i>Molecular Phylogenetics and Evolution</i> , 2004, 30, 686-702. | 2.7 | 314 |
| 85 | Phylogeny of Eusocial <i>Lasioglossum</i> Reveals Multiple Losses of Eusociality within a Primitively Eusocial Clade of Bees (Hymenoptera: Halictidae). <i>Systematic Biology</i> , 2003, 52, 23-36. | 5.6 | 123 |
| 86 | Evolution of sociality in a primitively eusocial lineage of bees. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 286-290. | 7.1 | 157 |
| 87 | PHYLOGEOGRAPHY OF THE SOCIALLY POLYMORPHIC SWEAT BEE HALICTUS RUBICUNDUS (HYMENOPTERA:) Tj ETQq1 1 0.784314 rgB | 2.3 | 9 |
| 88 | BEES IN THE NEW MILLENIUM. <i>BioScience</i> , 2002, 52, 848. | 4.9 | 0 |
| 89 | PHYLOGEOGRAPHY OF THE SOCIALLY POLYMORPHIC SWEAT BEE HALICTUS RUBICUNDUS (HYMENOPTERA:) Tj ETQq1 1 0.784314 rgB | 2.3 | 76 |
| 90 | Phylogenetic Utility of the Major Opsin in Bees (Hymenoptera: Apoidea): A Reassessment. <i>Molecular Phylogenetics and Evolution</i> , 2001, 19, 76-93. | 2.7 | 50 |

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|-----|--|-----|-----------|
| 91 | Australian <i>Lasioglossum</i> + <i>Homalictus</i> Form a Monophyletic Group: Resolving the "Australian Enigma". <i>Systematic Biology</i> , 2001, 50, 268-283. | 5.6 | 39 |
| 92 | Determining Parasitoid Species Composition in a Host Population: A Molecular Approach. <i>Annals of the Entomological Society of America</i> , 2000, 93, 640-647. | 2.5 | 52 |
| 93 | Male dimorphism in <i>Perdita portalis</i> (Hymenoptera, Andrenidae) has arisen from preexisting allometric patterns. <i>Insectes Sociaux</i> , 1999, 46, 18-28. | 1.2 | 33 |
| 94 | Phylogeny of the bee genus <i>Lasioglossum</i> (Hymenoptera: Halictidae) based on mitochondrial COI sequence data. <i>Systematic Entomology</i> , 1999, 24, 377-393. | 3.9 | 80 |
| 95 | Phylogeny of the Bee Genus <i>Halictus</i> (Hymenoptera: Halictidae) Based on Parsimony and Likelihood Analyses of Nuclear EF-1 α Sequence Data. <i>Molecular Phylogenetics and Evolution</i> , 1999, 13, 605-618. | 2.7 | 104 |
| 96 | Two New and Highly Apomorphic Species of the <i>Lasioglossum</i> Subgenus <i>Evylaeus</i> (Hymenoptera: Andrenidae). <i>Journal of Hymenoptera Research</i> , 2000, 10, 1-10. | 2.5 | 50 |
| 97 | Reply from W.T. Wcislo and B.N. Danforth. <i>Trends in Ecology and Evolution</i> , 1998, 13, 199. | 8.7 | 0 |
| 98 | Mitochondrial Dna Differentiation between Two Cryptic <i>Halictus</i> (Hymenoptera: Halictidae) Species. <i>Annals of the Entomological Society of America</i> , 1998, 91, 387-391. | 2.5 | 40 |
| 99 | Secondarily solitary: the evolutionary loss of social behavior. <i>Trends in Ecology and Evolution</i> , 1997, 12, 468-474. | 8.7 | 159 |
| 100 | NESTMATE RELATEDNESS IN A COMMUNAL BEE, <i>PERDITA TEXANA</i> (HYMENOPTERA: ANDRENIDAE), BASED ON DNA FINGERPRINTING. <i>Evolution; International Journal of Organic Evolution</i> , 1996, 50, 276-284. | 2.3 | 16 |
| 101 | Dynamics of a Host-Cleptoparasite Relationship: <i>Holcopasites ruthae</i> as a Parasite of <i>Calliopsis pugionis</i> (Hymenoptera: Anthophoridae, Andrenidae). <i>Annals of the Entomological Society of America</i> , 1993, 86, 833-840. | 2.5 | 16 |
| 102 | Biology of <i>Calliopsis pugionis</i> (Hymenoptera: Andrenidae): Nesting, Foraging, and Investment Sex Ratio. <i>Annals of the Entomological Society of America</i> , 1993, 86, 822-832. | 2.5 | 37 |
| 103 | Male Polymorphism and Polyethism in <i>Perdita texana</i> (Hymenoptera: Andrenidae). <i>Annals of the Entomological Society of America</i> , 1992, 85, 616-626. | 2.5 | 32 |
| 104 | A review of <i>Perdita</i> , subgenus <i>Macrotera</i> (Hymenoptera: Andrenidae). <i>Contributions in Science</i> , 1992, 436, 1-12. | 0.3 | 1 |
| 105 | The morphology and behavior of dimorphic males in <i>Perdita portalis</i> (Hymenoptera : Andrenidae). <i>Behavioral Ecology and Sociobiology</i> , 1991, 29, 235-247. | 1.4 | 99 |
| 106 | Provisioning behavior and the estimation of investment ratios in a solitary bee, <i>Calliopsis</i> (<i>Hypomacrotera</i>) <i>persimilis</i> (Cockerell) (Hymenoptera: Andrenidae). <i>Behavioral Ecology and Sociobiology</i> , 1990, 27, 159. | 1.4 | 70 |
| 107 | The evolution of hymenopteran wings: the importance of size. <i>Journal of Zoology</i> , 1989, 218, 247-276. | 1.7 | 50 |
| 108 | Wing Folding in the Hymenoptera. <i>Annals of the Entomological Society of America</i> , 1988, 81, 342-349. | 2.5 | 32 |

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|-----|---|-----|-----------|
| 109 | Mechanisms of spontaneous mutagenesis: An analysis of the spectrum of spontaneous mutation in the Escherichia coli lacI gene. <i>Journal of Molecular Biology</i> , 1986, 189, 273-284. | 4.2 | 330 |
| 110 | Rapid repeated cloning of mutant lac repressor genes. <i>Gene</i> , 1985, 39, 181-189. | 2.2 | 119 |
| 111 | The evolution of social behavior in the augochlorine sweat bees (Hymenoptera: Halictidae) based on a phylogenetic analysis of the genera. , 0, , 270-292. | | 54 |