Theodore G Schurr

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2071312/publications.pdf

Version: 2024-02-01

66 papers

3,237 citations

201674 27 h-index 55 g-index

70 all docs

70 docs citations

times ranked

70

3044 citing authors

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Genetic Variation in the E6 and E7 Genes of Human Papillomavirus Type 16 in Northeastern Argentina. Journal of Medical Virology, 2022, 94, 745-751. | 5.0 | 3 |
| 2 | Contrasting maternal and paternal genetic histories among five ethnic groups from Khyber Pakhtunkhwa, Pakistan. Scientific Reports, 2022, 12, 1027. | 3.3 | O |
| 3 | Investigating variability in the frequency of fire use in the archaeological record of Late Pleistocene Europe. Archaeological and Anthropological Sciences, 2022, 14, 1. | 1.8 | 4 |
| 4 | Evolution and dispersal of mitochondrial DNA haplogroup U5 in Northern Europe: insights from an unsupervised learning approach to phylogeography. BMC Genomics, 2022, 23, 354. | 2.8 | 3 |
| 5 | Ancient DNA reveals five streams of migration into Micronesia and matrilocality in early Pacific seafarers. Science, 2022, 377, 72-79. | 12.6 | 13 |
| 6 | Ancestry, health, and lived experiences of enslaved Africans in 18th century Charleston: An osteobiographical analysis. American Journal of Physical Anthropology, 2021, 175, 3-24. | 2.1 | 15 |
| 7 | Y chromosome diversity in Aztlan descendants and its implications for the history of Central Mexico. IScience, 2021, 24, 102487. | 4.1 | 3 |
| 8 | Genetic landscape of Gullah African Americans. American Journal of Physical Anthropology, 2021, 175, 905-919. | 2.1 | 9 |
| 9 | Matrilineal diversity and population history of Norwegians. American Journal of Physical Anthropology, 2021, 176, 120-133. | 2.1 | 3 |
| 10 | Mitochondrial genetic variation in human bioenergetics, adaptation, and adult disease. American Journal of Human Biology, 2021, , e23629. | 1.6 | 1 |
| 11 | Host genetic factors and susceptibility to <scp>SARSâ€CoV</scp> â€2 infection. American Journal of Human Biology, 2020, 32, e23497. | 1.6 | 7 |
| 12 | Mitochondrial DNA diversity in the Khattak and Kheshgi of the Peshawar Valley, Pakistan. Genetica, 2020, 148, 195-206. | 1.1 | 3 |
| 13 | Ancient DNA and bioarchaeological perspectives on European and African diversity and relationships on the colonial Delaware frontier. American Journal of Physical Anthropology, 2019, 170, 232-245. | 2.1 | 6 |
| 14 | Genetic diversity of the JC polyomavirus (JCPyV) and mitochondrial DNA ancestry in Misiones, Argentina. Infection, Genetics and Evolution, 2019, 75, 104011. | 2.3 | 4 |
| 15 | Analysis of biogeographic ancestry reveals complex genetic histories for indigenous communities of St. Vincent and Trinidad. American Journal of Physical Anthropology, 2019, 169, 482-497. | 2.1 | 20 |
| 16 | Mitochondrial DNA ancestry, HPV infection and the risk of cervical cancer in a multiethnic population of northeastern Argentina. PLoS ONE, 2018, 13, e0190966. | 2.5 | 10 |
| 17 | Mitochondrial DNA diversity of present-day Aboriginal Australians and implications for human evolution in Oceania. Journal of Human Genetics, 2017, 62, 343-353. | 2.3 | 24 |
| 18 | Genetic diversity in <scp>S</scp> vaneti and its implications for the human settlement of the <scp>H</scp> ighland <scp>C</scp> aucasus. American Journal of Physical Anthropology, 2017, 164, 837-852. | 2.1 | 4 |

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| 19 | New native South American Y chromosome lineages. Journal of Human Genetics, 2016, 61, 593-603. | 2.3 | 28 |
| 20 | Genetic Diversity in the Lesser Antilles and Its Implications for the Settlement of the Caribbean Basin. PLoS ONE, 2015, 10, e0139192. | 2.5 | 22 |
| 21 | Oxytocin receptor gene sequences in owl monkeys and other primates show remarkable interspecific regulatory and protein coding variation. Molecular Phylogenetics and Evolution, 2015, 91, 160-177. | 2.7 | 11 |
| 22 | Genographic Project., 2015,, 22-31. | | 0 |
| 23 | Tracing Human Movements from Siberia to the Americas: Insights from Genetic Studies. , 2015, , 23-47. | | 27 |
| 24 | Genetic characterization and clinical implications of human papillomavirus type 16 (HPV16) variants from northeastern Argentina. Infection, Genetics and Evolution, 2015, 29, 103-109. | 2.3 | 6 |
| 25 | Mitochondrial DNA Variant in COX1 Subunit Significantly Alters Energy Metabolism of Geographically Divergent Wild Isolates in Caenorhabditis elegans. Journal of Molecular Biology, 2014, 426, 2199-2216. | 4.2 | 49 |
| 26 | Correlates of genetic monogamy in socially monogamous mammals: insights from Azara's owl monkeys. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20140195. | 2.6 | 73 |
| 27 | Prolactin Receptor Gene Diversity in Azara's Owl Monkeys (Aotus azarai) and Humans (Homo sapiens) Suggests a Non-Neutral Evolutionary History among Primates. International Journal of Primatology, 2014, 35, 129-155. | 1.9 | 5 |
| 28 | Genetic diversity in $\langle scp \rangle P \langle scp \rangle uerto \langle scp \rangle R \langle scp \rangle ico$ and its implications for the peopling of the $\langle scp \rangle I \langle scp \rangle sland$ and the $\langle scp \rangle W \langle scp \rangle est \langle scp \rangle I \langle scp \rangle ndies$. American Journal of Physical Anthropology, 2014, 155, 352-368. | 2.1 | 34 |
| 29 | Geographic population structure analysis of worldwide human populations infers their biogeographical origins. Nature Communications, 2014, 5, 3513. | 12.8 | 114 |
| 30 | Mitochondrial genetic background modulates bioenergetics and susceptibility to acute cardiac volume overload. Biochemical Journal, 2013, 455, 157-167. | 3.7 | 79 |
| 31 | The GenoChip: A New Tool for Genetic Anthropology. Genome Biology and Evolution, 2013, 5, 1021-1031. | 2.5 | 54 |
| 32 | Genetic Background and Climatic Droplet Keratopathy Incidence in a Mapuche Population from Argentina. PLoS ONE, 2013, 8, e74593. | 2.5 | 6 |
| 33 | Analysis of TNFα promoter SNPs and the risk of cervical cancer in urban populations of Posadas (Misiones, Argentina). Journal of Clinical Virology, 2012, 53, 54-59. | 3.1 | 27 |
| 34 | Y-chromosome analysis reveals genetic divergence and new founding native lineages in Athapaskan- and Eskimoan-speaking populations. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 8471-8476. | 7.1 | 54 |
| 35 | Clan, language, and migration history has shaped genetic diversity in Haida and Tlingit populations from Southeast Alaska. American Journal of Physical Anthropology, 2012, 148, 422-435. | 2.1 | 37 |
| 36 | Mitochondrial DNA and Y Chromosome Variation Provides Evidence for a Recent Common Ancestry between Native Americans and Indigenous Altaians. American Journal of Human Genetics, 2012, 90, 229-246. | 6.2 | 146 |

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|----|--|-----|-----------|
| 37 | An Optimized Microsatellite Genotyping Strategy for Assessing Genetic Identity and Kinship in Azara's Owl Monkeys (Aotus azarai). Folia Primatologica, 2011, 82, 107-117. | 0.7 | 12 |
| 38 | Y-Chromosome Variation in Altaian Kazakhs Reveals a Common Paternal Gene Pool for Kazakhs and the Influence of Mongolian Expansions. PLoS ONE, 2011, 6, e17548. | 2.5 | 58 |
| 39 | 7. The Prehistory of Mongolian Populations as Revealed by Studies of Osteological, Dental, and Genetic Variation., 2011,, 125-165. | | О |
| 40 | Biological Ancestries, Kinship Connections, and Projected Identities in Four Central Anatolian Settlements: Insights from Culturally Contextualized Genetic Anthropology. American Anthropologist, 2011, 113, 116-131. | 1.4 | 22 |
| 41 | Parallel Evolution of Genes and Languages in the Caucasus Region. Molecular Biology and Evolution, 2011, 28, 2905-2920. | 8.9 | 149 |
| 42 | mtDNA diversity in azara's owl monkeys (<i>Aotus azarai azarai</i>) of the Argentinean Chaco. American Journal of Physical Anthropology, 2011, 146, 209-224. | 2.1 | 31 |
| 43 | Genetic ancestry and indigenous heritage in a Native American Descendant Community in Bermuda. American Journal of Physical Anthropology, 2011, 146, 392-405. | 2.1 | 19 |
| 44 | Who Are the Anatolian Turks?. Anthropology and Archeology of Eurasia, 2011, 50, 6-42. | 0.0 | 10 |
| 45 | AVPR1A Sequence Variation in Monogamous Owl Monkeys (Aotus azarai) and Its Implications for the Evolution of Platyrrhine Social Behavior. Journal of Molecular Evolution, 2010, 71, 279-297. | 1.8 | 24 |
| 46 | Genetic heritage and native identity of the Seaconke Wampanoag tribe of massachusetts. American Journal of Physical Anthropology, 2010, 142, 579-589. | 2.1 | 16 |
| 47 | Evaluation of Group Genetic Ancestry of Populations from Philadelphia and Dakar in the Context of Sex-Biased Admixture in the Americas. PLoS ONE, 2009, 4, e7842. | 2.5 | 33 |
| 48 | Response to Decoding Implications of the Genographic Project. International Journal of Cultural Property, 2009, 16, 182-187. | 0.3 | 3 |
| 49 | Haplotypic Background of a Private Allele at High Frequency in the Americas. Molecular Biology and Evolution, 2009, 26, 995-1016. | 8.9 | 74 |
| 50 | Genetic variation in the enigmatic Altaian Kazakhs of Southâ€Central Russia: Insights into Turkic population history. American Journal of Physical Anthropology, 2008, 136, 278-293. | 2.1 | 46 |
| 51 | Russian Old Believers: Genetic Consequences of Their Persecution and Exile, as Shown by Mitochondrial DNA Evidence. Human Biology, 2008, 80, 203-237. | 0.2 | 10 |
| 52 | De novo COX2 mutation in a LHON family of Caucasian origin: implication for the role of mtDNA polymorphism in human pathology. Journal of Human Genetics, 2006, 51, 161-170. | 2.3 | 12 |
| 53 | Reconstructing the Origins and Migrations of Diasporic Populations: The Case of the European Gypsies. American Anthropologist, 2004, 106, 267-281. | 1.4 | 33 |
| 54 | Mitochondrial DNA and Y chromosome diversity and the peopling of the Americas: Evolutionary and demographic evidence. American Journal of Human Biology, 2004, 16, 420-439. | 1.6 | 162 |

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| 55 | The Peopling of the New World: Perspectives from Molecular Anthropology. Annual Review of Anthropology, 2004, 33, 551-583. | 1.5 | 168 |
| 56 | Mitochondrial DNA Variation and the Origins of the Aleuts. Human Biology, 2003, 75, 809-835. | 0.2 | 76 |
| 57 | Mitochondrial DNA Diversity in Southeast Asian Populations. Human Biology, 2002, 74, 431-452. | 0.2 | 42 |
| 58 | The Dual Origin and Siberian Affinities of Native American Y Chromosomes. American Journal of Human Genetics, 2002, 70, 192-206. | 6.2 | 169 |
| 59 | mtDNA Variation in the South African Kung and Khwe—and Their Genetic Relationships to Other African Populations. American Journal of Human Genetics, 2000, 66, 1362-1383. | 6.2 | 188 |
| 60 | Mitochondrial DNA and the Peopling of the New World. American Scientist, 2000, 88, 246. | 0.1 | 19 |
| 61 | Mitochondrial DNA variation in Koryaks and Itel'men: Population replacement in the Okhotsk Sea-Bering Sea region during the neolithic. American Journal of Physical Anthropology, 1999, 108, 1-39. | 2.1 | 186 |
| 62 | mtDNA Diversity in Chukchi and Siberian Eskimos: Implications for the Genetic History of Ancient Beringia and the Peopling of the New World. American Journal of Human Genetics, 1998, 63, 1473-1491. | 6.2 | 209 |
| 63 | mtDNA Haplogroup X: An Ancient Link between Europe/Western Asia and North America?. American Journal of Human Genetics, 1998, 63, 1852-1861. | 6.2 | 263 |
| 64 | Y chromosome polymorphisms in Native American and Siberian populations: identification of Native American Y chromosome haplotypes. Human Genetics, 1997, 100, 536-543. | 3.8 | 81 |
| 65 | The structure of human mitochondrial DNA variation. Journal of Molecular Evolution, 1991, 33, 543-555. | 1.8 | 213 |
| 66 | Mitochondrial DNA variation in Koryaks and Itel'men: Population replacement in the Okhotsk Seaâ \in Bering Sea region during the neolithic. , 0, . | | 1 |