

# James Torrance

## List of Publications by Year in descending order

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Version: 2024-02-01

22  
papers

4,906  
citations

840776

11  
h-index

888059

17  
g-index

32  
all docs

32  
docs citations

32  
times ranked

4650  
citing authors

#	ARTICLE	IF	CITATIONS
1	The complete sequence of a human genome. <i>Science</i> , 2022, 376, 44-53.	12.6	1,222
2	Reference genome and demographic history of the most endangered marine mammal, the vaquita. <i>Molecular Ecology Resources</i> , 2021, 21, 1008-1020.	4.8	54
3	Significantly improving the quality of genome assemblies through curation. <i>GigaScience</i> , 2021, 10, .	6.4	739
4	A high-quality, chromosome-level genome assembly of the Black Soldier Fly ( <i>Hermetia illucens</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	1.8	37
5	Towards complete and error-free genome assemblies of all vertebrate species. <i>Nature</i> , 2021, 592, 737-746.	27.8	1,139
6	The genome sequence of the Norway rat, <i>Rattus norvegicus</i> Berkenhout 1769. <i>Wellcome Open Research</i> , 2021, 6, 118.	1.8	16
7	The genome sequence of the brown trout, <i>Salmo trutta</i> Linnaeus 1758. <i>Wellcome Open Research</i> , 2021, 6, 108.	1.8	15
8	The genome sequence of the European golden eagle, <i>Aquila chrysaetos chrysaetos</i> Linnaeus 1758. <i>Wellcome Open Research</i> , 2021, 6, 112.	1.8	3
9	A Chromosome-Level Genome Assembly of the Reed Warbler ( <i>Acrocephalus scirpaceus</i> ). <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	4
10	The complete genome sequence of <i>Eimeria tenella</i> (Tyzzer 1929), a common gut parasite of chickens. <i>Wellcome Open Research</i> , 2021, 6, 225.	1.8	14
11	A high-quality genome and comparison of short- versus long-read transcriptome of the palaeartic duck <i>Aythya fuligula</i> (tufted duck). <i>GigaScience</i> , 2021, 10, .	6.4	7
12	The genome sequence of the Eurasian red squirrel, <i>Sciurus vulgaris</i> Linnaeus 1758. <i>Wellcome Open Research</i> , 2020, 5, 18.	1.8	3
13	The genome sequence of the eastern grey squirrel, <i>Sciurus carolinensis</i> Gmelin, 1788. <i>Wellcome Open Research</i> , 2020, 5, 27.	1.8	4
14	The genome sequence of the channel bull blenny, <i>Cottoperca gobio</i> (GÃ¼nther, 1861). <i>Wellcome Open Research</i> , 2020, 5, 148.	1.8	18
15	Evaluation of GRCh38 and de novo haploid genome assemblies demonstrates the enduring quality of the reference assembly. <i>Genome Research</i> , 2017, 27, 849-864.	5.5	728
16	gEVALâ€”a web-based browser for evaluating genome assemblies. <i>Bioinformatics</i> , 2016, 32, 2508-2510.4.1		319
17	Modernizing Reference Genome Assemblies. <i>PLoS Biology</i> , 2011, 9, e1001091.	5.6	458
18	The genome sequence of the common pipistrelle, <i>Pipistrellus pipistrellus</i> Schreber 1774. <i>Wellcome Open Research</i> , 0, 6, 117.	1.8	2

#	ARTICLE	IF	CITATIONS
19	The genome sequence of the European water vole, <i>Arvicola amphibius</i> Linnaeus 1758. Wellcome Open Research, 0, 6, 162.	1.8	1
20	The genome sequence of the ringlet, <i>Aphantopus hyperantus</i> Linnaeus 1758. Wellcome Open Research, 0, 6, 165.	1.8	4
21	The genome sequence of the European robin, <i>Erithacus rubecula</i> Linnaeus 1758. Wellcome Open Research, 0, 6, 172.	1.8	2
22	The genome sequence of the European turtle dove, <i>Streptopelia turtur</i> Linnaeus 1758. Wellcome Open Research, 0, 6, 191.	1.8	4