Mariko Kondo

List of Publications by Year
in descending order

11 papers	1,064 citations	933447 10 h-index	1281871 11 g-index
12 all docs	12 docs citations	12 times ranked	1767 citing authors

#	Article	IF	Citations
1	Genome evolution in the allotetraploid frog Xenopus laevis. Nature, 2016, 538, 336-343.	27.8	849
2	A New Nomenclature of Xenopus laevis Chromosomes Based on the Phylogenetic Relationship to Silurana/Xenopus tropicalis. Cytogenetic and Genome Research, 2015, 145, 187-191.	1.1	50
3	Genomic insights of body plan transitions from bilateral to pentameral symmetry in Echinoderms. Communications Biology, 2020, 3, 371.	4.4	34
4	Conservatism and variability of gene expression profiles among homeologous transcription factors in Xenopus laevis. Developmental Biology, 2017, 426, 301-324.	2.0	24
5	Regeneration in crinoids. Development Growth and Differentiation, 2010, 52, 57-68.	1.5	21
6	Current Status of Echinoderm Genome Analysis - What do we Know?. Current Genomics, 2012, 13, 134-143.	1.6	18
7	Comprehensive analyses of hox gene expression in Xenopus laevis embryos and adult tissues. Development Growth and Differentiation, 2017, 59, 526-539.	1.5	16
8	Weighted gene co-expression network analysis reveals potential genes involved in early metamorphosis process in sea cucumber Apostichopus japonicus. Biochemical and Biophysical Research Communications, 2018, 495, 1395-1402.	2.1	15
9	Regeneration of the digestive tract of an anterior-eviscerating sea cucumber, Eupentacta quinquesemita, and the involvement of mesenchymalâ€“epithelial transition in digestive tube formation. Zoological Letters, 2019, 5, 21.	1.3	15
10	Pmar1/phb homeobox genes and the evolution of the double-negative gate for endomesoderm specification in echinoderms. Development (Cambridge), 2020, 147, .	2.5	14
11	de novotranscription of multiple Hox cluster genes takes place simultaneously in earlyXenopus tropicalisembryos. Biology Open, 2019, 8, .	1.2	8