

# Liz Milla

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5333934/publications.pdf>

Version: 2024-02-01

12  
papers

603  
citations

1039406

9  
h-index

1199166

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

1531  
citing authors

#	ARTICLE	IF	CITATIONS
1	Monitoring of honey bee floral resources with pollen DNA metabarcoding as a complementary tool to vegetation surveys. <i>Ecological Solutions and Evidence</i> , 2022, 3, .	0.8	13
2	A molecular phylogeny and revision of the genus <i>Pyropteron</i> Newman, 1832 (Lepidoptera, Sesiidae) reveals unexpected diversity and frequent hostplant switch as a driver of speciation. <i>Zootaxa</i> , 2021, 4972, 175.	0.2	1
3	Pollen DNA metabarcoding identifies regional provenance and high plant diversity in Australian honey. <i>Ecology and Evolution</i> , 2021, 11, 8683-8698.	0.8	22
4	Phylotranscriptomics resolves phylogeny of the Heliozelidae (Adeloidea: Lepidoptera) and suggests a Late Cretaceous origin in Australia. <i>Systematic Entomology</i> , 2020, 45, 128-143.	1.7	8
5	Menstrual fluid factors facilitate tissue repair: identification and functional action in endometrial and skin repair. <i>FASEB Journal</i> , 2019, 33, 584-605.	0.2	22
6	CARD11 is dispensable for homeostatic responses and suppressive activity of peripherally induced FOXP3 + regulatory T cells. <i>Immunology and Cell Biology</i> , 2019, 97, 740-752.	1.0	10
7	A preliminary molecular phylogeny of shield-bearer moths (Lepidoptera: Adeloidea: Heliozelidae) highlights rich undescribed diversity. <i>Molecular Phylogenetics and Evolution</i> , 2018, 120, 129-143.	1.2	13
8	DNA repair processes are critical mediators of p53-dependent tumor suppression. <i>Nature Medicine</i> , 2018, 24, 947-953.	15.2	122
9	A new extant family of primitive moths from Kangaroo Island, Australia, and its significance for understanding early Lepidoptera evolution. <i>Systematic Entomology</i> , 2015, 40, 5-16.	1.7	32
10	An Inducible Lentiviral Guide RNA Platform Enables the Identification of Tumor-Essential Genes and Tumor-Promoting Mutations In Vivo. <i>Cell Reports</i> , 2015, 10, 1422-1432.	2.9	337
11	Association of candidate single nucleotide polymorphisms with somatic mutation of the epidermal growth factor receptor pathway. <i>BMC Medical Genomics</i> , 2013, 6, 43.	0.7	8
12	Cre transgene results in global attenuation of the cAMP/PKA pathway. <i>Cell Death and Disease</i> , 2012, 3, e365-e365.	2.7	15