

Charley G P Mccarthy

List of Publications by Year in descending order

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12
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840776

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694
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Pangloss: A Tool for Pan-Genome Analysis of Microbial Eukaryotes. <i>Genes</i> , 2019, 10, 521. | 2.4 | 14 |
| 2 | Proteome and allergenome of the European house dust mite <i>Dermatophagoides pteronyssinus</i> . <i>PLoS ONE</i> , 2019, 14, e0216171. | 2.5 | 26 |
| 3 | Whole Genome Sequence of the Commercially Relevant Mushroom Strain <i>Agaricus bisporus</i> var. <i>bisporus</i> ARP23. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 3057-3066. | 1.8 | 13 |
| 4 | Pan-genome analyses of model fungal species. <i>Microbial Genomics</i> , 2019, 5, . | 2.0 | 59 |
| 5 | Antibiotic resistomes of healthy pig faecal metagenomes. <i>Microbial Genomics</i> , 2019, 5, . | 2.0 | 23 |
| 6 | Investigating the pangenomes of microbial eukaryotes. <i>Access Microbiology</i> , 2019, 1, . | 0.5 | 0 |
| 7 | Evolutionary, structural and functional analysis of the caleosin/peroxygenase gene family in the Fungi. <i>BMC Genomics</i> , 2018, 19, 976. | 2.8 | 11 |
| 8 | Genome analysis of the yeast <i>Diutina catenulata</i> , a member of the Debaryomycetaceae/Metschnikowiaceae (CTG-Ser) clade. <i>PLoS ONE</i> , 2018, 13, e0198957. | 2.5 | 13 |
| 9 | Phylogenomic Reconstruction of the Oomycete Phylogeny Derived from 37 Genomes. <i>MSphere</i> , 2017, 2, . | 2.9 | 84 |
| 10 | Draft Genome Sequence of <i>Dermatophagoides pteronyssinus</i> , the European House Dust Mite. <i>Genome Announcements</i> , 2017, 5, . | 0.8 | 18 |
| 11 | Multiple Approaches to Phylogenomic Reconstruction of the Fungal Kingdom. <i>Advances in Genetics</i> , 2017, 100, 211-266. | 1.8 | 19 |
| 12 | Systematic Search for Evidence of Interdomain Horizontal Gene Transfer from Prokaryotes to Oomycete Lineages. <i>MSphere</i> , 2016, 1, . | 2.9 | 22 |