

# Changmu Kim

## List of Publications by Year in descending order

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18  
papers

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| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Major Antioxidant Compound of <i>Polyporus Parvovarius</i> Culture Filtrate. Natural Product Communications, 2022, 17, 1934578X2110723.   | 0.5 | 3         |
| 2  | Two Unrecorded <i>Apiospora</i> Species Isolated from Marine Substrates in Korea with Eight New Combinations ( <i>A. piptatheri</i> and <i>A. rasikravindrae</i> ). Mycobiology, 2022, 50, 46-54.   | 1.7 | 6         |
| 3  | Antioxidant Activities and Mechanisms of Tomentosin in Human Keratinocytes. Antioxidants, 2022, 11, 990.  | 5.1 | 4         |
| 4  | The genus <i>Arthrinium</i> (Ascomycota, Sordariomycetes, Apiosporaceae) from marine habitats from Korea, with eight new species. IMA Fungus, 2021, 12, 13.   | 3.8 | 18        |
| 5  | First Report of Six Macrofungi from Daechongdo and Socheongdo Islands, Korea. Mycobiology, 2021, 49, 454-460.   | 1.7 | 2         |
| 6  | <i>Zygorulasporea cornina</i> sp. nov. and <i>Zygorulasporea smilacis</i> sp. nov., Two Novel Ascomycetous Yeast Species Isolated from Plant Flowers and Fruits. Mycobiology, 2021, 49, 1-6.  | 1.7 | 1         |
| 7  | Influence of Tree Vegetation on Soil Microbial Communities in Temperate Forests and Their Potential as a Proactive Indicator of Vegetation Shift Due to Climate Change. Sustainability, 2020, 12, 10591.  | 3.2 | 5         |
| 8  | Successional Variation in the Soil Microbial Community in Odaesan National Park, Korea. Sustainability, 2020, 12, 4795.   | 3.2 | 11        |
| 9  | Taxonomic revision of <i>Russula</i> subsection <i>Amoeninae</i> from South Korea. MycoKeys, 2020, 75, 1-29.  | 1.9 | 11        |
| 10 | Diversity and Ecology of Marine Algicolous <i>Arthrinium</i> Species as a Source of Bioactive Natural Products. Marine Drugs, 2018, 16, 508.  | 4.6 | 20        |
| 11 | Sequencing and de novo assembly of visceral mass transcriptome of the critically endangered land snail <i>Satsuma myomphala</i> : Annotation and SSR discovery. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2017, 21, 77-89.     | 1.0 | 10        |
| 12 | De novo Transcriptome Generation and Annotation for Two Korean Endemic Land Snails, <i>Aegista chejuensis</i> and <i>Aegista quelpartensis</i> , Using Illumina Paired-End Sequencing Technology. International Journal of Molecular Sciences, 2016, 17, 379. | 4.1 | 7         |
| 13 | MP-V1 from the Venom of Social Wasp <i>Vespula vulgaris</i> Is a de Novo Type of Mastoparan that Displays Superior Antimicrobial Activities. Molecules, 2016, 21, 512.  | 3.8 | 15        |
| 14 | Transcriptomic Analysis of the Endangered Neritid Species <i>Clithon retropictus</i> : De Novo Assembly, Functional Annotation, and Marker Discovery. Genes, 2016, 7, 35.   | 2.4 | 13        |
| 15 | Transcriptome sequencing and de novo characterization of Korean endemic land snail, <i>Koreanohadra kurodana</i> for functional transcripts and SSR markers. Molecular Genetics and Genomics, 2016, 291, 1999-2014.   | 2.1 | 14        |
| 16 | Sequencing, De Novo Assembly, and Annotation of the Transcriptome of the Endangered Freshwater Pearl Bivalve, <i>Cristaria plicata</i> , Provides Novel Insights into Functional Genes and Marker Discovery. PLoS ONE, 2016, 11, e0148622.                    | 2.5 | 61        |
| 17 | Transcriptome Characterization for Non-Model Endangered Lycaenids, <i>Protantigius superans</i> and <i>Spindasis takanosis</i> , Using Illumina HiSeq 2500 Sequencing. International Journal of Molecular Sciences, 2015, 16, 29948-29970.                    | 4.1 | 13        |
| 18 | Construction of PANM Database (Protostome DB) for rapid annotation of NGS data in Mollusks. Korean Journal of Malacology, 2015, 31, 243-247.  | 0.1 | 17        |