

1 **Whole Genome Sequence of an High Temperature Edible**
2 **Mushroom *Pleurotus giganteus* (Zhudugu)**

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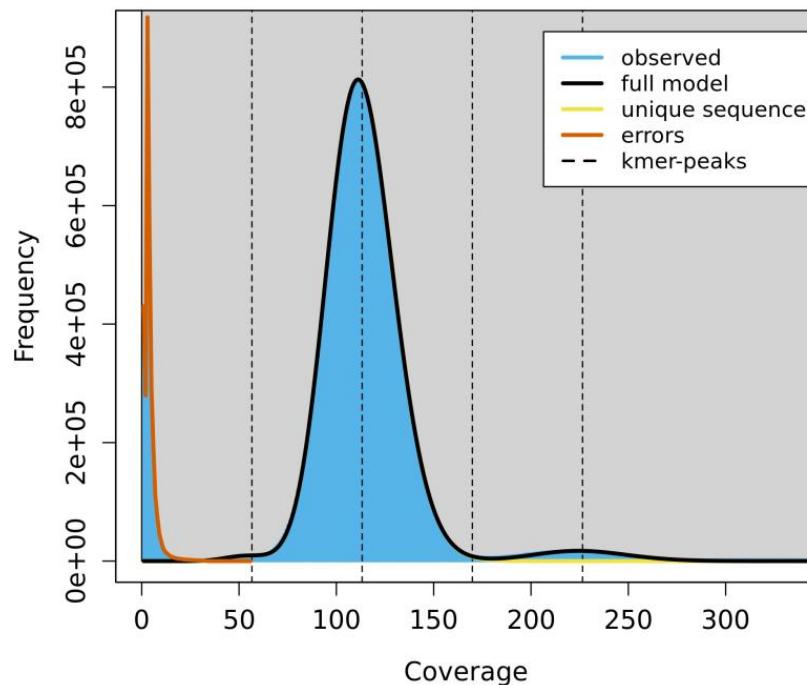
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20 **Keywords:** *Pleurotus giganteus*; genome; edible mushroom; white-rot fungi; mating locus, CAZymes

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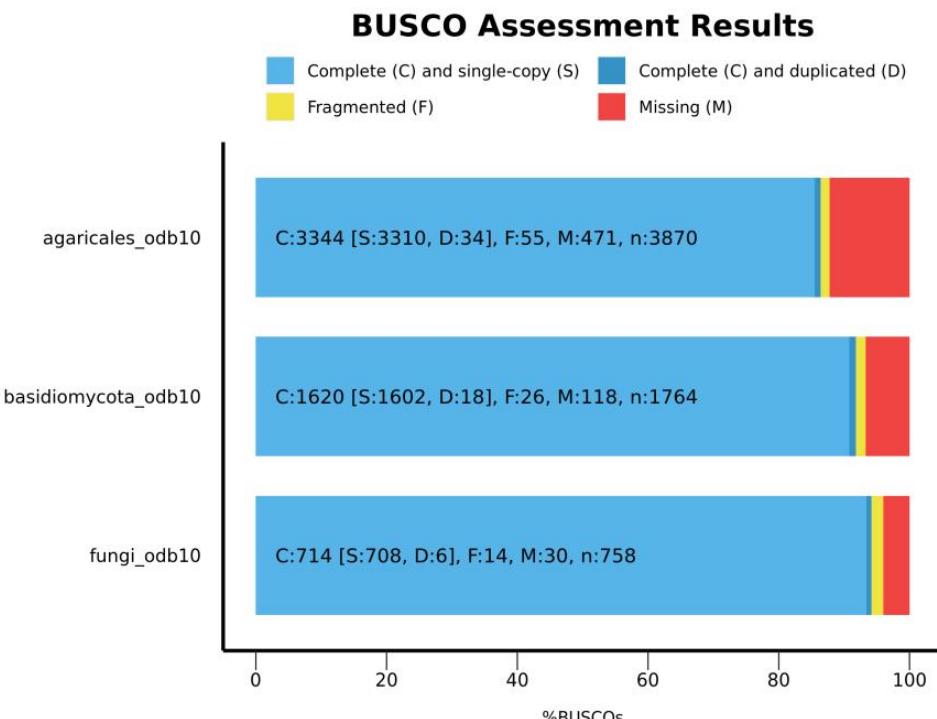
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24 **Figure S1**

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26 **Figure S1.** Histogram of the 19-mer depth distribution of the Illumina sequencing reads of *P. giganteus*
27 plotted in GenomeScope. Blue areas indicate the observed k-mer frequencies and the black line
28 indicates the fitted GenomeScope model ($p = 2$).

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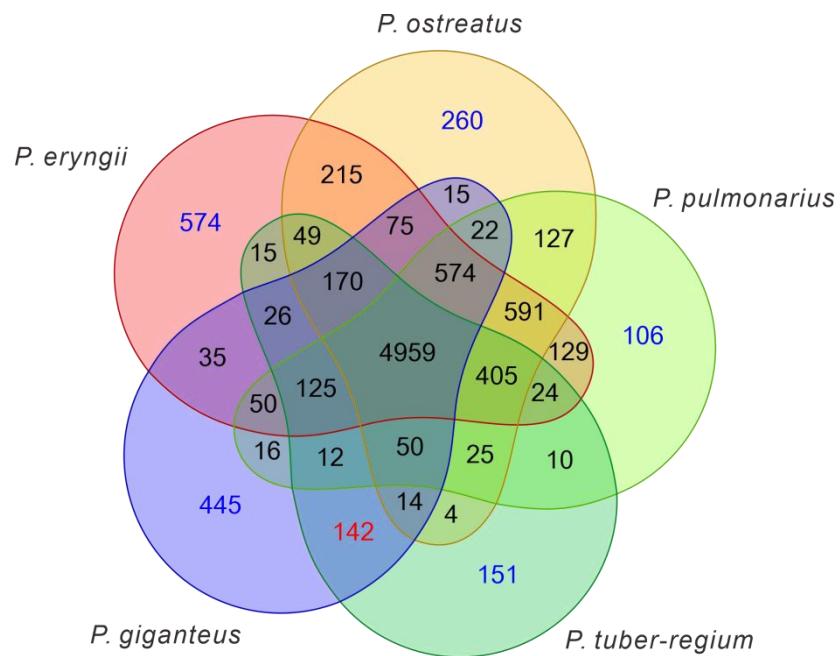
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32 **Figure S2.** BUSCO assessment of the protein annotation completeness in *P. giganteus* with different
33 BUSCO databases.

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36 **Figure S3**



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38 **Figure S3.** Venn diagram of the orthogroups of five *Pleurotus* species.

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41 **Table S1** Characteristics of the gene prediction of *P. giganteus*

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Content	Number/length
Gene number	12628
Average gene length	1761.11 bp
Average protein length	586.04 aa
Total exon number	90508
Total exon/gene length	22239338 bp
Average exon length	245.72 bp
Average exon number per gene	7.17 bp
Total intron number	77880
Total intron length	23734624 bp
Average intron length	304.76 bp
Average intron number per gene	6.17 bp

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