

专论与综述

# 低温刺激对食用菌子实体形成的研究进展

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**摘要:** 探究低温刺激子实体形成过程的组学数据能为食用菌的工厂化低碳高效栽培及广温型品种的选育提供参考。本文首先通过主成分分析对食用菌低温刺激子实体形成的类型进行低温型( $11\text{--}19\text{ }^{\circ}\text{C}$ )和冷压型的划分( $\leq 10\text{ }^{\circ}\text{C}$ )。在此基础上, 综述了食用菌两种类型的低温刺激子实体形成过程的研究进展。食用菌低温刺激子实体形成过程中都涉及信号转导、胁迫响应、基础代谢、细胞分化、细胞结构变化等代谢过程。随着低温型到冷压型的刺激温度下降, 糖代谢可能转向脂代谢为子实体形成提供能量。

**关键词:** 食用菌; 低温刺激; 子实体形成; 功能基因组学; 功能基因

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## Advances in low temperature stimulation for the fruiting body formation of edible fungi

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**Abstract:** Exploring the process of low temperature stimulating fruiting body formation (LSFF) can provide a reference for the industrialized low-carbon efficient cultivation of edible fungi and the breeding of varieties with wide temperature ranges. This study first performed principal component analysis to classify LSFF of edible fungi into the low-temperature type (11–19 °C) and the cold stress type ( $\leq 10$  °C). On this basis, the research progress in the two types of LSFF in edible fungi was reviewed. The LSFF in edible fungi involves the metabolic processes such as signal transduction, stress response, basic metabolism, cell differentiation, and cell structure changes. With the decrease in the stimulation temperature from the low-temperature type to cold stress type, the sugar metabolism may shift to lipid metabolism to provide energy for fruiting body formation.

**Keywords:** edible fungi; low temperature stimulation; fruiting body formation; functional genomics; functional gene

我国的食用菌资源丰富，将近千种，约一百多种食用菌被广泛栽培食用，市面常见的食用菌约 20 种<sup>[1]</sup>。香菇(*Lentinus edodes*)、金针菇(*Flammulina velutipes*)、黑木耳(*Auricularia auricula*)、杏鲍菇(*Pleurotus eryngii*)等品种年产量可达  $2 \times 10^6$ – $1 \times 10^7$  t<sup>[2]</sup>。食用菌从孢子萌发为营养菌丝到成熟的子实体再产生孢子，这一生长周期受多种环境因子的影响，诸如温度、湿度、光照、空气和酸碱度等。环境因子的变化可以直接影响子实体的形成。例如：高温会使香菇菌丝抗逆性减弱，超过阈值后造成菌丝死亡<sup>[3]</sup>；强光照射抑制菌丝生长，适当强度的光照对子实体分化有利<sup>[4]</sup>；空气中氧气浓度高有

助于菌丝生长及其子实体分化，高浓度二氧化碳则对食用菌生长有抑制作用<sup>[5]</sup>；多数食用菌的最适生长 pH 值为 5.0–8.0，在 pH 8.0 以上的碱性环境会停止生长<sup>[6]</sup>。工厂化栽培是食用菌生产发展的重要方向，而工厂化栽培重要的特点就是需要环境因子模块化，选择最适宜的环境因子，对工厂化低碳高效栽培具有重要的理论和实践价值。

温度作为影响子实体发育的关键因子<sup>[7]</sup>，影响菌丝的生长速度和子实体的质量<sup>[8]</sup>。这与食用菌的生命活动依赖于一系列功能蛋白有关，而蛋白活性与温度的变化密切相关<sup>[6]</sup>。如草菇(*Volvariella volvacea*)纤维二糖水解酶 Vvcbh1-1

在 4 °C 时活性升高引发草菇菌丝的低温自溶<sup>[9]</sup>; 张介弛等<sup>[10]</sup>发现黑木耳在 20 °C 发菌时菌丝生长较好, 耳质好、产量高, 相较 25 °C 发菌产量提高 37.9%; 田景花等<sup>[11]</sup>发现杏鲍菇子实体对温度较敏感, 错过原基形成温度(15 °C)将不再出菇。食用菌子实体分化需要较低的温度, 最佳分化温度往往低于菌丝生长温度<sup>[12]</sup>, 食用菌原基形成和子实体分化需要低温诱导或者温差的刺激<sup>[13]</sup>。

低温诱导是促进许多食用菌子实体结实的主要手段<sup>[14]</sup>。Weiser 等<sup>[15]</sup>首先提出了植物抗寒过程中需要特异基因的激活和新蛋白诱导合成的观点, 低温刺激食用菌结实的分子机理核心也可以认为是这一概念的延伸, 即冷诱导基因的表达与调控。

目前, 转录组学、蛋白质组学、代谢组学等功能基因组学能系统地研究冷诱导基因的表达与调控<sup>[16]</sup>。例如, 蛋白质组学技术可以研究蛋白质的组成、表达、结构、功能和相互作用, 揭示特定生命过程的内在机制, 在更深层次上探索生命活动的规律<sup>[17]</sup>。考虑到低温刺激子实体结实的温度区间范围差异幅度较大(大约 20 °C), 我们首先在已有食用菌温型分类的基础上, 对食用菌低温刺激温型进行了进一步的划分。鉴于组学技术已经在食用菌子实体形成的研究中大量应用<sup>[18]</sup>, 重点对不同低温型的组学研究进行了总结。围绕低温刺激子实体形成的相关通路和基因的功能进行了讨论, 以期为食用菌工厂化低碳高效栽培及广温性品种选育提供理论参考。

## 1 食用菌温型划分

温度是食用菌生长的重要环境因素, 在其菌丝生长、子实体形成等阶段都有相应的最适温度。一些食用菌能在环境温度不变即恒温条件下完成子实体的分化, 称为恒温结实。一些

食用菌需要温差刺激才能分化出子实体, 称为变温结实, 如香菇、金针菇等<sup>[6]</sup>。

根据原基分化所需的温度不同可将食用菌分为低温型、中温型和高温型<sup>[19]</sup>。庞杰等<sup>[14]</sup>根据菌丝和子实体生长的温度需求详细分析了 29 种食用菌, 将其细分为四大类: 第一类为低温结实型食用菌, 子实体分化温度范围 13.8–19.4 °C, 低于菌丝生长温度范围(21.1–24.4 °C); 第二类为中温型食用菌, 菌丝和子实体发育所需温度相近(17.9–27 °C); 第三类为高温结实型, 菌丝生长温度(24–29 °C)低于子实体的分化温度(27.6–31.7 °C); 第四类为高温型, 菌丝和子实体发育所需温度均较高(27–37 °C)。本文为探究低温刺激对子实体形成的影响, 将子实体形成温度低于 20 °C 的低温结实型进一步分为 2 个类型(图 1, 表 1): (1) 低温型(11–19 °C), 包括香菇、真姬菇(*Hypholoma marmoreus*)、双孢蘑菇(*Agaricus bisporus*)和秀珍菇(*Pleurotus geesteranus*)等。(2) 冷压型(≤10 °C), 包括金针菇、白灵菇(*Pleurotus nebrodensis*)和滑菇(*Pholiota nameko*)等。

## 2 低温刺激对低温型食用菌子实体形成的影响

食用菌在长期的进化中产生许多适应温度逆境的能力。基因组、转录组等组学分析技术的发展拓宽了人们对食用菌子实体形成的认知, 成为研究细胞功能和表型的热点<sup>[29]</sup>。当外界温度下降时, 一系列特异基因的表达集中于能量代谢、氧化还原酶活性和靶蛋白降解等过程<sup>[30]</sup>。诱导蛋白在抵御低温渗透、稳定酶的结构、胞外酶代谢中发挥作用, 协调细胞内各项代谢水平从而使细胞适应新的环境<sup>[31]</sup>。

### 2.1 低温型食用菌的信号感知与转导

G 蛋白偶联受体(G protein-coupled receptor, GPCR)在 cAMP 信号通路中起着关键作用。碳

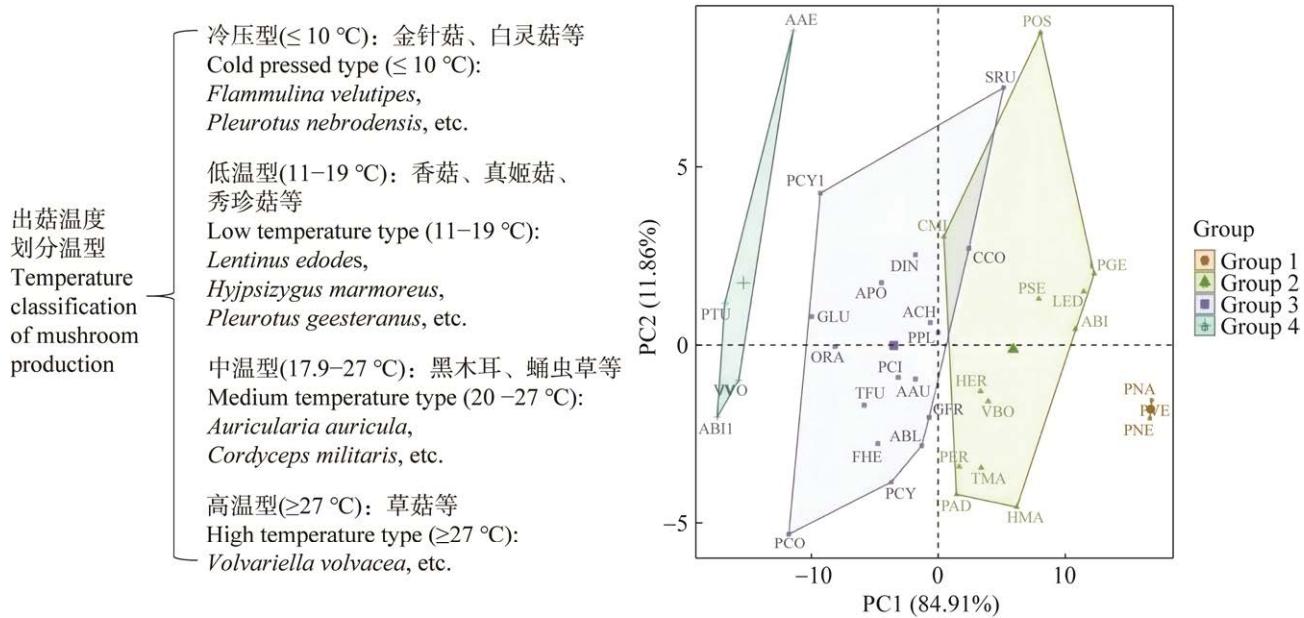


图 1 食用菌子实体分化温度的主成分分析 以子实体分化的最适温度区间的均值作为划分标准: 低于 11 °C 为冷压型(group 1), 11–19 °C 为低温型(group 2), 17.9–27 °C 为中温型(group 3), 高于 27 °C 的作为高温型(group 4); FVE 为 *Flammulina velutipes* 的第一个单词的第一位和后一个单词的前两位的缩写, 以此类推; 子实体分化温度数据引用庞杰等报道的数据<sup>[14]</sup>

Figure 1 Principal component analysis of fruiting body differentiation temperature. The average value of the optimal temperature range for the differentiation of the fruiting body is taken as the division standard, the temperature below 11 °C is cold pressing type (group 1), 11–19 °C is low temperature type (group 2), 17.9–27 °C is medium temperature type (group 3), and the temperature above 27 °C is high temperature type (group 4); FVE is the abbreviation of the first digit of the first word and the first two digits of the last word of *Flammulina velutipes*, and so on; The fruiting body differentiation temperature data refer to the date reported by Pang<sup>[14]</sup>.

源会被 G 蛋白偶联受体 1 (G protein-coupled receptor 1, GPR1) 感知, 激活 G 蛋白  $\alpha$  亚单位 2 (G protein alpha subunit 2, GPA2), GPA2 激活腺苷酸环化酶(adenylyl cyclase, AC)提高胞内 cAMP 浓度<sup>[32]</sup>, 过表达 G 蛋白  $\alpha$  亚基基因 *GANI* 也能提高内源 cAMP 水平, 产生的 cAMP 经过 G 蛋白介导的信号转导诱导结实<sup>[33-34]</sup>。灰盖鬼伞 (*Coprinopsis cinerea*) 在菌丝扭结时加入 AC 抑制剂后无法形成原基, 外源添加 cAMP 能诱导子实体的形成<sup>[35]</sup>; 杏鲍菇(12 °C刺激)原基形成差异基因富集于 cAMP 信号转导、碳水化合物代谢和细胞壁降解<sup>[36]</sup>; 高水平的 cAMP 能提高淀粉酶和漆酶活性<sup>[34]</sup>, 表明子实体形成需要大量能量,

cAMP/PKA 信号转导途径是子实体形成的关键。

## 2.2 低温型食用菌子实体发育的调控基因表达

锌指蛋白、DNA 结合蛋白均可作为转录因子调控子实体的发育<sup>[37-38]</sup>。C2H2 (Cys<sub>2</sub>His<sub>2</sub>)型锌指蛋白在双孢蘑菇子实体整个发育阶段明显上调<sup>[39]</sup>, 可能参与菌丝聚集到原基的转换<sup>[40]</sup>。C2H2 型锌指蛋白在干旱、低温等胁迫下有增强作物低温耐受力的作用<sup>[41]</sup>, 能调控大部分胁迫基因的表达<sup>[42]</sup>, 在丝状真菌中广泛参与碳代谢调控, 参与纤维素降解相关酶的表达<sup>[43]</sup>, 因此 C2H2 型锌指蛋白上调表明双孢蘑菇子实体形成期间对碳源的需求增加。DNA 结合蛋白基因

表 1 低温结实型食用菌类型划分

Table 1 Classification of low temperature fruiting edible fungi

食用菌种类 Edible fungi species	菌丝生长适宜温度 Suitable temperature for mycelium growth (°C)	子实体分化适宜温度 Suitable temperature for fruiting body differentiation (°C)	参考文献 References
滑菇 <i>Pholiota nameko</i> **	20–25	5–15	[14]
白灵菇 <i>Pleurotus nebrodensis</i> **	22–25	–2–5	[20]
金针菇 <i>Flammulina velutipes</i> **	22–24	6–14	[14]
双孢蘑菇 <i>Agaricus bisporus</i> *	22–25	13–16	[14]
香菇 <i>Lentinus edodes</i> *	23–25	12–17	[14]
姬菇 <i>Pleurotus cornucopiae</i> *	20–25	12–15	[21]
大球盖菇 <i>Stropharia rugoso-annulata</i> *	24–28	15–19	[14]
羊肚菌 <i>Morchella esculenta</i> *	20–24	15–18	[22]
杏鲍菇 <i>Pleurotus eryngii</i> *	20–22	14–15	[23]
秀珍菇 <i>Pleurotus geesteranus</i> *	20–24	15–18	[24]
榛蘑 <i>Armillariella mellea</i> *	23–25	14–20	[25]
元蘑 <i>Hohenbuehelia serotina</i> *	22	14	[26]
真姬菇 <i>Hypholoma marmoreum</i> *	20–25	12–16	[27]
大肥蘑菇 <i>Agaricus bitorquis</i> *	18–22	15–18	[28]

\*: 低温型; \*\*: 冷压型

\*: Low temperature type; \*\*: Cold pressed type.

*Le.cdc5* 在香菇原基和幼子实体中高表达, 延伸因子基因 *pri10022* 在香菇原基和成熟子实体中显著表达<sup>[44]</sup>, 香菇冷刺激后基因表达和蛋白翻译高度活跃。新研究发现一个香菇 DNA 结合蛋白基因 *Le-Dd10*, 敲除后不形成子实体<sup>[45]</sup>, *Le-Dd10* 蛋白有 56、27 和 14 kDa 三个分子量, 在子实体形成较快的香菇品种中 14 kDa 蛋白表达较强; 56–14 kDa 的变化可能对刺激子实体形成极其重要, 然而 56、27 和 14 kDa 蛋白在原基和子实体之间的关系尚不清楚<sup>[45]</sup>。*large fruiting body cap 1 (LFC1)*基因编码的转录因子 LFC1 负调节金针菇(15 °C刺激)子实体形成, 敲除该基因促进子实体发育, 出菇时间缩短 2 d, LFC1 同源物广泛存在于食用菌中, 可用于蘑菇育种<sup>[46]</sup>。

泛素化修饰在细胞生长中有重大作用。通过泛素激活酶 E1、泛素结合酶 E2、泛素连接酶 E3 共同作用使蛋白质降解, 参与信号转导、

DNA 修复、异常蛋白代谢和应激反应等<sup>[47–48]</sup>。担子菌泛素蛋白复合物随着发育进程而变化<sup>[49]</sup>, 鸡腿菇(*Coprinus comatus*)原基形成过程中, 泛素与泛素蛋白结合物同时增加, 在子实体发育成熟过程中菌柄下部比上部高<sup>[50]</sup>, 真姬菇原基和幼子实体中也存在类似现象<sup>[51]</sup>。降解异常蛋白维持细胞稳态, 得到的短肽和氨基酸可作为原料用于抵抗胁迫, 这可能有利于原基形成。F-box 蛋白参与真核生物蛋白泛素化等多种生物学过程<sup>[52]</sup>, F-box 编码基因也被发现在金针菇、蜜环菌(*Armillaria mellea*)、裂褶菌(*Schizophyllum commune* Fr.)子实体阶段特异性表达<sup>[53]</sup>, 表明 F-box 基因在伞菌子实体发育中具有重要作用。

真核翻译起始因子 2 (eukaryotic initiation factor 2, eIF2)几乎是所有 mRNA 翻译起始必需的, general control non-derepressible 2 (GCN2) 是一种广泛分布在真核生物且参与氨基酸缺

乏、冷刺激等各种胁迫响应的蛋白激酶, 通过磷酸化 eIF2 的  $\alpha$  亚基抑制 eIF2-GDP 向 eIF2-GTP 转化, 阻止翻译起始。然而在真姬菇 15 °C刺激出菇时, GCN2 介导的翻译通路促进蛋白合成<sup>[54]</sup>。GCN2 介导的 eIF2 $\alpha$  磷酸化的应激反应机制在所有真核生物中并不相同。

### 2.3 低温型食用菌子实体中功能基因表达

细胞色素 P450 属于氧化酶系统, 在真菌中参与解毒反应、次生代谢和生长发育调节<sup>[55]</sup>。已发现一个细胞色素 P450 基因在双孢蘑菇子实体中特异表达<sup>[56]</sup>。灰盖鬼伞 P450 编码基因突变能导致菌柄变短<sup>[57]</sup>, P450 基因在子实体形成中可能是保守的。双孢蘑菇子实体形成中(15 °C 低温刺激)的差异基因主要参与氨基酸代谢。大量氨基酸合成酶基因高表达, 积累具有生理活性的氨基酸如精氨酸、脯氨酸、酪氨酸等<sup>[39]</sup>。氨基酸可作为合成其他物质的前体, 如精氨酸可分解为鸟氨酸和尿素<sup>[58]</sup>。鸟氨酸可合成多胺提供氮源。脯氨酸在植物和食用菌非生物胁迫下维持代谢, 也能作为分子伴侣清除活性氧, 在低温胁迫下调节渗透压维持细胞稳态<sup>[59]</sup>。

疏水蛋白可以在潮湿环境中促进表面菌丝聚集<sup>[60]</sup>, 当疏水蛋白分泌至细胞壁中时, 菌丝越过水汽进入空气中形成气生菌丝, 更多的气生菌丝有助于原基形成<sup>[61]</sup>。对外阻止膜对水的吸收, 防止子实体被水浸泡<sup>[62]</sup>。香菇中疏水蛋白基因 *SCI*、*SC3* 在子实体形成时高表达<sup>[63]</sup>。水通路蛋白负责水和溶质运输, 在金针菇菌柄中特异表达, 可能加强水分和营养物质运输, 促进菌柄和孢子发育。疏水蛋白和水通路蛋白特异表达时间可能与金针菇菌柄生长有关。水运输也是腐生真菌发育中的一个重要过程<sup>[64]</sup>, 在白灵菇等多种食用菌子实体发育中受调控<sup>[53]</sup>。

漆酶是一类多酚氧化酶, 在木质素降解、子实体发育中起重要作用<sup>[65]</sup>。漆酶基因在真姬

菇发育过程中的原基期后表达逐渐上升, 子实体形成后表达下降<sup>[66]</sup>; 金针菇 15 °C刺激出菇时胞外漆酶活性较高<sup>[65]</sup>, 也能进一步证明漆酶与食用菌子实体形成密切相关。

碳水化合物活性酶(carbohydrate-active enzymes, CAZymes)调控基因在真菌中参与细胞壁构建, 细胞壁自溶和重建是担子菌类食用菌子实体发育的基本过程<sup>[67]</sup>。丝状真菌细胞壁主要由几丁质和  $\beta$ -1,3-葡聚糖组成<sup>[68]</sup>, 细胞壁合成、裂解, 多糖组分的变化在担子菌子实体菌柄发育中必不可少<sup>[69-70]</sup>。几丁质酶、甘露糖苷酶基因在香菇和金针菇子实体发育中上调表达<sup>[67,71]</sup>, 赋予菌丝相互黏附扭结的可塑性<sup>[53]</sup>。有学者认为真菌细胞在增大过程中要依靠水解酶类的降解, 使细胞壁松弛解除束缚满足生长需求<sup>[67]</sup>。灰盖鬼伞(高温型)子实体成熟过程中几丁质酶 *ChiB1* 表达量明显增加<sup>[72]</sup>。正常褐变的香菇糖苷水解酶基因在子实体形成时高表达<sup>[63]</sup>, 表明细胞壁降解重塑可能是伞菌子实体发育的重要过程。真菌中还存在一类扩张蛋白, 通过无水解过程破坏葡聚糖链或几丁质链之间的非共价键, 促进细胞壁聚合物滑移, 疏松重塑细胞壁结构<sup>[73]</sup>。扩张蛋白基因 *Fvexpl1* 在金针菇(15 °C刺激)菌柄快速伸长中起作用, 可作为提高金针菇产量的候选基因<sup>[74]</sup>。

## 3 低温刺激对冷压型食用菌子实体发育的影响

冷压型食用菌往往需要 10 °C以下的低温刺激才可以出菇。Sakamoto 等<sup>[75]</sup>将金针菇置于 8 °C下 14 d 后形成子实体, 认为温度是诱导金针菇子实体的关键因素<sup>[76]</sup>。-1~2 °C冷压刺激可以促进白灵菇子实体形成, 刺激的时间越长, 菇蕾形成时间越短<sup>[20]</sup>。下面列举冷压型食用

菌的子实体形成的研究,重点对金针菇在低温和冷压两种刺激下的应答机理进行比较总结。

### 3.1 冷压型食用菌信号感知与转导

细胞膜流动性的改变是食用菌感知冷刺激的首要变化,膜脂物理变化可直接调节膜上感应蛋白活性,以此传递低温信号到下级通路<sup>[77]</sup>。杏鲍菇-3 °C刺激后磷脂酶和脂肪酸去饱和酶基因表达上调<sup>[78]</sup>,改变细胞膜组成以适应温度变化<sup>[79]</sup>。真菌中麦角甾醇参与膜功能域的形成、膜流动性和胞内转运,以及影响膜蛋白的定位等<sup>[80]</sup>。金针菇和肺形侧耳(*Pleurotus pulmonarius*)X57 菌株子实体形成时麦角甾醇合成通路活力上调<sup>[81-82]</sup>,推测麦角甾醇合成通路与原基形成有关。脂膜中磷脂酰肌醇 4,5-二磷酸[PI(4,5)P<sub>2</sub>]含量随温度降低而降低,被磷脂酶水解产生第二信使 IP3,在内质网与配体门钙通道结合,开启钙通道,使胞内 Ca<sup>2+</sup>浓度升高,激活各种钙依赖性蛋白<sup>[83]</sup>。金针菇菌丝和原基中钙依赖性蛋白激酶表达明显不同也证实了 Ca<sup>2+</sup>作为第二信使参与冷信号转导<sup>[84-85]</sup>。Ca<sup>2+</sup>信号和渗透调节基因在白灵菇 4 °C刺激 3 d 后表达降低<sup>[86]</sup>,这与冷胁迫能诱导 Ca<sup>2+</sup>浓度瞬时升高的结论<sup>[87]</sup>相一致。拟南芥(*Arabidopsis thaliana*)冷刺激时 OST1/SNRK2.6 激酶与渗透转运蛋白 ATANN1 相互作用磷酸化增强 Ca<sup>2+</sup>转运活性<sup>[87]</sup>, Ca<sup>2+</sup>信号与活性氧、MAPKS 信号通路交互作用提高细胞抗冻性<sup>[88]</sup>,真菌中可能也存在类似信号转导模式。

HOG-MAPK 传递低温等多种胁迫信号并促进甘油、阿拉伯糖醇、海藻糖合成。在高渗透环境下控制信号传导和基因表达<sup>[89]</sup>,在酿酒酵母(*Saccharomyces cerevisiae*)、蓝藻(*Cyanobacteria*)、金针菇子实体发育中也被证实能响应低温胁迫<sup>[85]</sup>。低温胁迫响应通过 SLN1 支路实现<sup>[90-91]</sup>,外界低温刺激使得 SLN1 支路磷酸化被抑制,去磷酸化的 SSK1 结合 SSK2

和 SSK22,从而激活 PBS2 和 HOG1,激活的 HOG1 进入细胞核发挥功能<sup>[92-93]</sup>,开启转录调控。肺形侧耳 4 °C低温处理后 HOG 通路中大部分基因都持续显著上调<sup>[94]</sup>。金针菇 10 °C、白灵菇 4 °C刺激 3 d 后膜系统、HOG1 型 MAPK 通路相关基因上调表达明显<sup>[51,85,95]</sup>。在拟南芥等模式生物中的 HOG 同源途径都能响应低温胁迫<sup>[96]</sup>。这表明 HOG-MAPK 通路对低温胁迫响应具有重要研究意义。

G 蛋白偶联受体(GPCRs)是最大的跨膜受体家族,介导运输功能蛋白到细胞内特定位置,调节膜运输以及连接到脂膜受体对不同信号分子做出反应<sup>[32,97]</sup>,参与真菌抵抗胁迫<sup>[98]</sup>。Yamada 等<sup>[99]</sup>将金针菇菌丝置于 8 °C诱导子实体形成,发现 58i cDNA 编码的氨基酸序列与 G 蛋白 α 亚基类似,Northern blotting 结果表明不同种类的 G 蛋白在子实体形成时起作用。信息素受体基因 PRLGs 属于 GPCRs 家族,对外界刺激产生反应,在金针菇子实体期高表达<sup>[100]</sup>。GPCRs 家族在刺激机制中起着重要作用,深入研究 PRLGs 的多种功能将更有利于阐明食用菌子实体发育机理。

### 3.2 冷压型食用菌子实体发育的调控基因表达

转录因子基因 *PDD1* 的过表达能促进金针菇子实体形成,产量提高 33%,缩短出菇时间 4 d<sup>[101]</sup>。余颖豪等<sup>[13]</sup>利用转录组测序技术分析了金针菇菌丝和原基冷压(10 °C)刺激后的 7 935 个差异表达基因,差异基因功能注释显示糖酵解过程中异构酶、磷酸化酶基因表达下调,柠檬酸合酶和脂代谢相关基因表达上调,提示冷压作用下主要由三羧酸循环(tricarboxylic acid cycle, TCA cycle)和脂质代谢产能;几丁质合酶基因表达下调而 β-葡聚糖合成相关基因表达上调,细胞壁成分发生变化,可能与原基形成相关。与

此对比, 15 °C刺激下金针菇原基中糖酵解相关酶基因表达较高, 在原基形成时参与能量代谢<sup>[102]</sup>。金针菇在长期 12 °C冷胁迫后葡萄糖-6-磷酸异构酶、β-葡萄糖苷酶、海藻糖酶活性显著提高, 海藻糖酶能催化海藻糖转化为葡萄糖<sup>[103]</sup>。以上研究提示, 冷压作用下金针菇启动脂代谢的供应能量的方式, 而低温刺激则采用糖代谢的供应能量的方式。这与食用菌糖代谢相关酶受到冷压胁迫调控的报道<sup>[104]</sup>相吻合。

### 3.3 冷压型食用菌子实体中的功能基因表达

冷压型刺激温度较低, 细胞膜首先对低温做出反应, 感受到冷信号后金针菇转变供能模式, 可能由糖代谢转向脂代谢, 秀珍菇也调节了营养物供应。前述冷压型食用菌冷压后差异基因富集于碳水化合物代谢过程、能量代谢、翻译调节、氧化还原酶活性和细胞壁重塑等, 保护细胞免受损伤的同时准备形成子实体, 这些研究都表明 10 °C以下低温对于冷压型食用菌的子实体形成是必需的。

阿魏菇(*Pleurotus ferulae*)在 10 °C冷压处理后漆酶活力明显提高, 漆酶酶谱也发生变化, 出现一条 33 kDa 的新酶带<sup>[105]</sup>。秀珍菇成熟菌丝胞外羧甲基纤维素酶、滤纸纤维素酶、半纤维素酶和淀粉酶活性与对照相比都随温度下降而增强, 胞外漆酶活性则随温度下降而降低<sup>[106]</sup>。此结果表明冷压处理后秀珍菇酶谱和酶的活性都发生了变化, 从利用木质素转为利用纤维素的酶谱转变。 $\beta$ -1,6-葡聚糖合成酶在细胞壁交联中具有重要作用, 过表达  $\beta$ -1,6-葡聚糖合成酶基因 *FFGS6* 能增加金针菇菌丝细胞宽度, 提高搔菌后机械损伤恢复能力, 但原基和子实体形成(8 °C刺激)受到抑制,  $\beta$ -1,6-葡聚糖合成酶基因在大型真菌子实体发育和逆境反应中的作用尚不清楚<sup>[107]</sup>。

在金针菇原基阶段疏水蛋白基因显著上调<sup>[64]</sup>,

并且基因有阶段表达的特性<sup>[68]</sup>。Northern blotting结果显示金针菇疏水蛋白基因 *Fv-hydI* 在菌丝中未表达, 在原基和子实体(8 °C刺激)中表达较高<sup>[108]</sup>。沉默疏水蛋白基因 *Hyd9* 能导致金针菇(10 °C刺激)原基和子实体减少, 高表达能增加气生菌丝有助于原基形成, 这表明 *Hdy9* 在金针菇菌丝扭结时具有重要作用<sup>[61]</sup>。疏水蛋白在菌丝聚集时上调表达<sup>[109]</sup>, 对疏水蛋白家族基因进行验证将有助于更深入地理解疏水蛋白的功能以及原基形成的机制。

本研究总结了子实体形成过程中差异基因的主要功能和富集通路, 以及两种低温刺激类型的不同之处(图 2)。该模型突出 GPCRs、HOG-MAPK、cAMP/PKA 等信号转导在子实体形成过程中的调控作用, 还提示了随着温度的下降功能模式从糖代谢到脂代谢的转换。

## 4 讨论与结论

工厂化栽培低温结实菌类能源消耗巨大, 从分子层面研究不同类型的食用菌低温应答机理有助于培育广温型食用菌, 对食用菌低碳高效栽培具有重大意义。温度是影响食用菌子实体分化发育的关键因素, 大部分食用菌菌丝营养生长阶段的最适温度相近, 而子实体生殖生长阶段的温度则有较大差距。因此, 本文将低温结实类型进一步分为低温型(11–19 °C)和冷压型( $\leq 10$  °C), 结合子实体发育保守基因综述了它们受低温胁迫后的应答机制。

信号转导在两种温型的食用菌中暂未发现差异, G 蛋白、Ca<sup>2+</sup>、cAMP/PKA、HOG-MAPK 等信号途径均有积极调控作用。冷压型食用菌会降低膜的流动性从而限制细胞生长。真菌通过增加膜脂中不饱和脂肪链的比例来保持低温下的膜流动性<sup>[110]</sup>, 温度下降时细胞周期蛋白依赖性激酶连接了不同脂质合成途径, 平衡各种

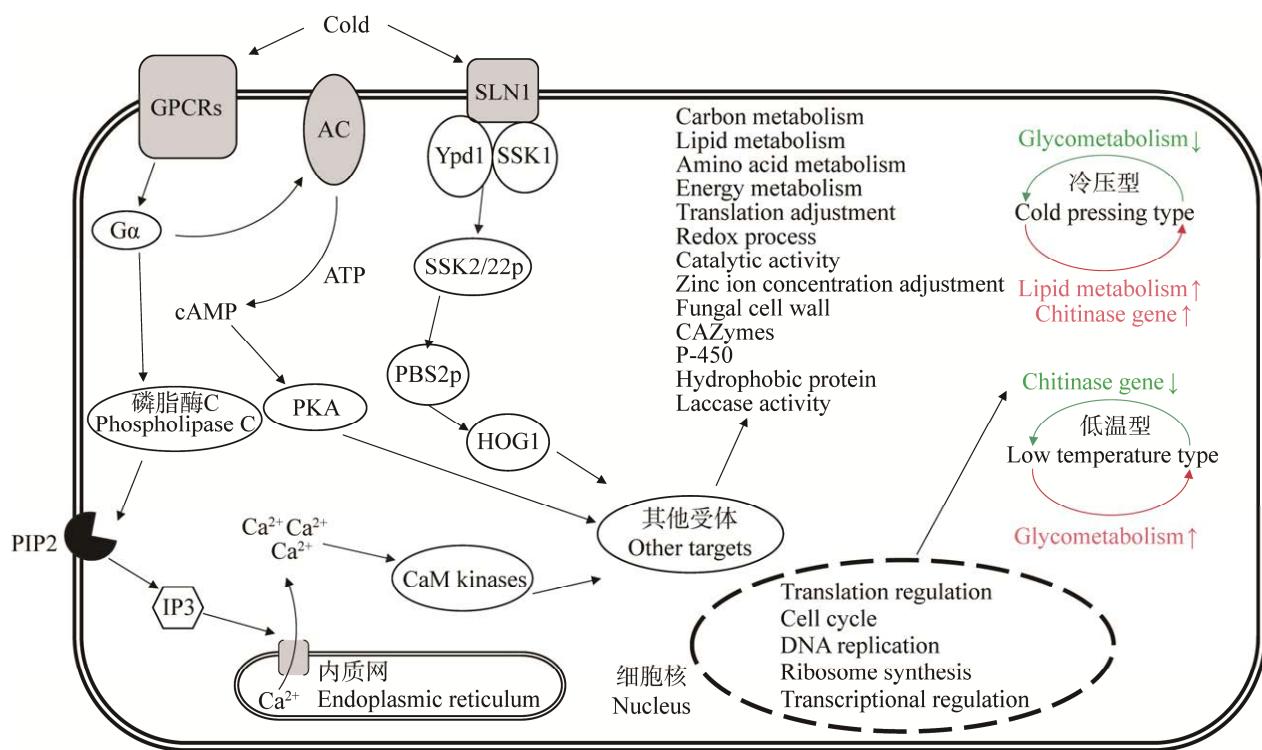


图 2 子实体形成过程中差异基因的主要功能和富集通路

Figure 2 Main functions and enrichment pathways of differential genes during fruiting body formation.

脂质丰度,是维持脂质稳定的核心调节因子<sup>[83]</sup>,这暗示了冷压刺激更多地影响了细胞膜的脂质结构,但细胞如何感知膜流动性改变,以及不同脂质丰度还需进一步研究<sup>[77]</sup>。

子实体发育调节中不同温型刺激下的金针菇能量利用存在明显的差异。金针菇 10 °C冷压刺激时大部分糖酵解相关基因和几丁质酶基因下调,供能方式转向脂代谢;而在 15 °C低温刺激下,糖酵解相关酶基因上调参与能量代谢<sup>[13,102]</sup>,这种由于冷压刺激带来的糖代谢转向脂代谢表达的模式是否是食用菌一般的调控模式,以及其在子实体形成中的作用仍需进一步探究。漆酶在两种温型食用菌子实体发育时均有重要作用(如金针菇、真姬菇、阿魏菇等)<sup>[111]</sup>,而秀珍菇漆酶活性随温度下降,各种纤维素酶活性增强,低温和冷压刺激下秀珍菇发生了酶谱转变,

摄取的营养物变化与其子实体形成的关系有待进一步探究。

有研究显示锌离子浓度和酵母胁迫耐性有关,胞内有机锌可促进酵母菌体生长<sup>[112]</sup>,外源添加锌离子有利于细胞能量代谢和膜稳定,提高抗应激能力<sup>[113-114]</sup>。低温型香菇和冷压型金针菇 *pria* 基因在原基和幼子实体中高表达,此基因可显著降低锌离子积累,参与调控子实体发育<sup>[115]</sup>,而在其他中高温型食用菌如蜜环菌、虎皮香菇(*Lentinus tigrinus*)、灰盖鬼伞、裂褶菌中表达较低<sup>[64]</sup>。这表明 *pria* 基因调节锌离子浓度与不同温型的食用菌子实体形成变化趋势关联性不大,这种调节模式机理在子实体早期发育中仍然未知。

生物体是一个基因、蛋白、信号传递的综合体,由一条调控途径可以发掘出更多的调控网

络。探究食用菌在冷刺激后的胁迫响应有助于更好地理解食用菌子实体形成的分子机制。未来可根据多组学数据对冷胁迫相关基因进行功能验证, 进而进行培育广温型食用菌的遗传工程改造, 助力于食用菌高效栽培和低碳经济的发展。

## REFERENCES

- [1] 汤昕明, 冯云利, 郭相, 马明, 陈正启, 吴素蕊. 我国食药用菌栽培现状分析及展望[J]. 北方园艺, 2019(8): 148-153.  
TANG XM, FENG YL, GUO X, MA M, CHEN ZQ, WU SR. Current situation and development trend of edible medicinal fungi in China[J]. Northern Horticulture, 2019(8): 148-153 (in Chinese).
- [2] 中国食用菌协会. 2021 年度全国食用菌统计调查结果分析[J]. 中国食用菌, 2023, 42(1): 118-127.  
China Edible Fungi Association. Analysis on the results of national edible fungi statistical survey in 2021[J]. Edible Fungi of China, 2023, 42(1): 118-127 (in Chinese).
- [3] 梁宇庭, 周骏辉, 袁媛, 赵玉洋, 黄璐琦. 环境胁迫影响药用真菌生长及生理生化机制的研究进展[J]. 中国现代中药, 2017, 19(12): 1780-1784.  
LIANG YT, ZHOU JH, YUAN Y, ZHAO YY, HUANG LQ. Research progress on growth and metabolism changes under environmental stress of medicinal fungi[J]. Modern Chinese Medicine, 2017, 19(12): 1780-1784 (in Chinese).
- [4] 李玉, 于海龙, 周峰, 王瑞娟, 李正鹏, 郭倩. 光照对食用菌生长发育影响的研究进展[J]. 食用菌, 2011, 33(2): 3-4.  
LI Y, YU HL, ZHOU F, WANG RJ, LI ZP, GUO Q. Advances in researching for the effect of illumination on the growth and development of edible fungi[J]. Edible Fungi, 2011, 33(2): 3-4 (in Chinese).
- [5] YANG HP, LEI MG, HUANG L, WANG Y, SUN N, BAN L, WANG XF, ZHANG HY. Study on the effects of environmental factors on enzyme activities during growth of *Hypsizygus marmoreus*[J]. PLoS One, 2022, 17(8): e0268107.
- [6] 周砾红. 秀珍菇变温结实相关 *Ppcsl-1* 基因的克隆及表达分析[D]. 金华: 浙江师范大学硕士学位论文, 2016.  
ZHOU SH. Cloning and expression analysis of *Ppcsl-1* gene related to variable temperature fruiting of *Pleurotus geesteranus*[D]. Jinhua: Master's Thesis of Zhejiang Normal University, 2016 (in Chinese).
- [7] 王海燕, 孙国琴, 庞杰, 解亚杰, 李亚娇. 环境胁迫对食用菌生长发育影响的研究进展[J]. 食药用菌, 2017, 25(2): 110-112.  
WANG HY, SUN GQ, PANG J, XIE YJ, LI YJ. Research progress on the influence of environmental stress on the growth and development of edible fungi[J]. Edible and Medicinal Mushrooms, 2017, 25(2): 110-112 (in Chinese).
- [8] 王再明, 石堃, 潘玲玲, 马玉心, 崔大练. 温度对8种食用菌菌丝生长的影响[J]. 林业科技, 2015, 40(3): 20-23.  
WANG ZM, SHI K, PAN LL, MA YX, CUI DL. The influence of temperature on mycelial growth of eight kinds of food[J]. Forestry Science & Technology, 2015, 40(3): 20-23 (in Chinese).
- [9] GONG M, WANG Y, BAO DP, JIANG S, CHEN HY, SHANG JJ, WANG XJ, HNIN YU H, ZOU G. Improving cold-adaptability of mesophilic cellulase complex with a novel mushroom cellobiohydrolase for efficient low-temperature ensiling[J]. Bioresource Technology, 2023, 376: 128888.
- [10] 张介驰, 韩增华, 张丕奇, 马银鹏, 王玉文, 陈鹤. 发菌温度对黑木耳菌丝和子实体生长的影响[J]. 食用菌学报, 2014, 21(2): 36-40.  
ZHANG JC, HAN ZH, ZHANG PQ, MA YP, WANG YW, CHEN H. Effects of cultivation temperature on *auricularia auricula-judae* mycelial growth and fruit body development[J]. Acta Edulis Fungi, 2014, 21(2): 36-40 (in Chinese).
- [11] 田景花, 赵润鹏, 李明, 李守勉. 温度和湿度对杏鲍菇子实体原基分化的影响[J]. 湖北农业科学, 2013, 52(20): 4975-4976, 4990.  
TIAN JH, ZHAO RP, LI M, LI SM. Effects of temperature and humidity on fruitbody primordium differentiation of *Pleurotus eryngii*[J]. Hubei Agricultural Sciences, 2013, 52(20): 4975-4976, 4990 (in Chinese).
- [12] 刘雁英, 李林辉, 谢富刚, 邱慧兰. 食用菌子实体分化发育研究进展[J]. 菌物研究, 2010, 8(1): 57-62.  
LIU YY, LI LH, XIE FG, QIU HL. Advances of research on fruiting body formation and development of edible fungi[J]. Journal of Fungal Research, 2010, 8(1): 57-62 (in Chinese).
- [13] 余颖豪, 伍土恒, 叶志伟, 陈柏雄, 郭丽琼, 林俊芳. 基于转录组分析的金针菇冷诱导原基形成的调控网络[J]. 菌物学报, 2020, 39(6): 1065-1076.

- YU YH, WU TH, YE ZW, CHEN BX, GUO LQ, LIN JF. Regulation network of cold-induced primordium formation of *Flammulina velutipes* based on transcriptome analysis[J]. *Mycosistema*, 2020, 39(6): 1065-1076 (in Chinese).
- [14] 庞杰, 孙国琴, 于静, 孟虎, 王海燕, 刘燕, 杨志刚. 35种食用菌菌丝生长和子实体分化温度的聚类分析[J]. 中国食用菌, 2015, 34(6): 34-37, 41.
- PANG J, SUN GQ, YU J, MENG H, WANG HY, LIU Y, YANG ZG. Cluster analysis based on the growth temperature of 35 edible fungi[J]. *Edible Fungi of China*, 2015, 34(6): 34-37, 41 (in Chinese).
- [15] WEISER CJ. Cold Resistance and Injury in Woody Plants: knowledge of hardy plant adaptations to freezing stress may help us to reduce winter damage[J]. *Science*, 1970, 169(3952): 1269-1278.
- [16] 龚明, 汪滢, 尚俊军, 唐利华, 尚晓冬, 张劲松, 谭琦, 鲍大鹏. 食药用菌系统演化及多组学研究进展[J]. 食用菌学报, 2018, 25(4): 85-96.
- GONG M, WANG Y, SHANG JJ, TANG LH, SHANG XD, ZHANG JS, TAN Q, BAO DP. Advances in phylogeny and multi-omics of edible and medicinal fungi[J]. *Acta Edulis Fungi*, 2018, 25(4): 85-96 (in Chinese).
- [17] 赵凤霞, 高相彬, 王正平, 李海峰, 宋学立, 冀敏. 蛋白质组学技术在烟草研究中的应用进展[J]. 中国烟草学报, 2014, 20(1): 103-110.
- ZHAO FX, GAO XB, WANG ZP, LI HF, SONG XL, JI M. Research and application of proteomics in tobacco[J]. *Acta Tabacaria Sinica*, 2014, 20(1): 103-110 (in Chinese).
- [18] 余金凤, 周汐, 赵春燕, 陈正启, 吴素蕊. 食用菌基因组学研究进展[J]. 蔬菜, 2017(3): 33-38.
- YU JF, ZHOU X, ZHAO CY, CHEN ZQ, WU SR. Research progress of edible fungi genomics[J]. *Vegetables*, 2017(3): 33-38 (in Chinese).
- [19] 刘红, 黄建军, 贺月林. 食用菌的温型及周年生产技术[J]. 现代农业科技, 2010(17): 147-148.
- LIU H, HUANG JJ, HE YL. Warm type and annual production technology of edible fungi[J]. *Xiandai Nongye Keji*, 2010(17): 147-148 (in Chinese).
- [20] 孔维威, 康源春, 胡素娟, 宋志波, 袁瑞奇, 韩玉娥, 张玉亭. 冷刺激温度和时间对瓶栽白灵菇子实体形成的影响[J]. 食药用菌, 2021, 29(2): 141-143.
- KONG WW, KANG YC, HU SJ, SONG ZB, YUAN RQ, HAN Y, ZHANG YT. Effect of temperature and time of cold stimulation on fruiting bodies formation of *Pleurotus uoliensis* under industrial bottle cultivation[J]. *Edible and Medicinal Mushrooms*, 2021, 29(2): 141-143 (in Chinese).
- [21] 姬婧, 蒋学杰. 姬菇栽培技术[J]. 特种经济动植物, 2020, 23(8): 39.
- JI J, JIANG XJ. Cultivation techniques of *Pleurotus cornucopiae*[J]. *Special Economic Animal and Plant*, 2020, 23(8): 39 (in Chinese).
- [22] 马建昌, 马小剑. 羊肚菌温室大棚种植技术要点[J]. 农业技术与装备, 2022(4): 136-138, 141.
- MA JC, MA XJ. Cultivation techniques of *Morchella* in greenhouse[J]. *Agricultural Technology & Equipment*, 2022(4): 136-138, 141 (in Chinese).
- [23] 赵书光, 李正鹏, 李玉, 张亚丽, 张翠娥, 徐相如. 杏鲍菇智能化高效生产关键技术研究[J]. 上海蔬菜, 2020(2): 65-67.
- ZHAO SG, LI ZP, LI Y, ZHANG YL, ZHANG CE, XU XR. Study on key technologies of intelligent and efficient production of *Pleurotus eryngii*[J]. *Shanghai Vegetables*, 2020(2): 65-67 (in Chinese).
- [24] 刘朋虎, 曾维源, 林占熲. 巨菌草工厂化栽培秀珍菇技术初探[J]. 食用菌, 2021, 43(2): 54-55.
- LIU PH, ZENG WY, LIN ZX. Preliminary study on industrial cultivation technology of *Pleurotus geesteranus* by giant weed[J]. *Edible Fungi*, 2021, 43(2): 54-55 (in Chinese).
- [25] 邬俊财, 张忠伟, 薛光艳, 邬昕彤, 于洋, 殷竹芬, 乔黎明. 蜜环菌(榛蘑)林地栽培技术[J]. 辽宁林业科技, 2009(3): 61-62.
- WU JC, ZHANG ZW, XUE GY, WU XT, YU Y, YIN ZF, QIAO LM. Planting techniques of *Armillaria mellea* on forest lands[J]. *Liaoning Forestry Science and Technology*, 2009(3): 61-62 (in Chinese).
- [26] 许泽成, 余悦, 尹龙川, 任美瑶. 北方元蘑冬季出菇栽培技术[J]. 食用菌, 2021, 43(2): 46-48.
- XU ZC, YU Y, YIN LC, REN MY. Cultivation techniques of northern yuanmo in winter[J]. *Edible Fungi*, 2021, 43(2): 46-48 (in Chinese).
- [27] 赖淑芳. 真姬菇工厂化栽培技术分析[J]. 农业与技术, 2021, 41(7): 98-100.
- LAI SF. Analysis of industrial cultivation techniques in *Hypsizygus marmoreus*[J]. *Agriculture & Technology*, 2021, 41(7): 98-100 (in Chinese).
- [28] 马宇生, 张利鹏. 大肥蘑菇低温型菌株的生物特性及其驯化栽培[J]. 食用菌, 2010, 32(6): 25-26.
- MA YS, ZHANG LP. Biological characteristics and domestication cultivation of low temperature strain of big fat mushroom[J]. *Edible Fungi*, 2010, 32(6): 25-26 (in Chinese).

- [29] de OLIVEIRA JMPF, de GRAAFF LH. Proteomics of industrial fungi: trends and insights for biotechnology[J]. *Applied Microbiology and Biotechnology*, 2011, 89(2): 225-237.
- [30] HAJELA RK, HORVATH DP, GILMOUR SJ, THOMASHOW MF. Molecular cloning and expression of cor (cold-regulated) genes in *Arabidopsis thaliana*[J]. *Plant Physiology*, 1990, 93(3): 1246-1252.
- [31] 黄桂英. 真姬菇低温胁迫下菌丝体酶活变化及差异蛋白质组学研究[D]. 福州: 福建农林大学硕士学位论文, 2008.  
HUANG GY. Changes and differences of enzyme activities of mycelium under low temperature stress in *Hypsizygus marmoreus*: a protein omics study[D]. Fuzhou: Master's Thesis of Fujian Agriculture and Forestry University, 2008 (in Chinese).
- [32] BAHN YS, XUE CY, IDNURM A, RUTHERFORD JC, HEITMAN J, CARDENAS ME. Sensing the environment: lessons from fungi[J]. *Nature Reviews Microbiology*, 2007, 5(1): 57-69.
- [33] YAMAGISHI K, KIMURA T, SUZUKI M, SHINMOTO H. Suppression of fruit-body formation by constitutively active G-protein  $\alpha$ -subunits ScGP-A and ScGP-C in the homobasidiomycete *Schizophyllum commune* the GenBank accession numbers for the sequences reported in this paper are AB066503, AB051903 and AB051904[J]. *Microbiology*, 2002, 148(9): 2797-2809.
- [34] CAO JX, SUN MJ, YU MM, XU YF, XIE JC, ZHANG HR, CHEN JY, XU T, QIAN X, SUN SJ. Transcriptome analysis reveals the function of a G-protein  $\alpha$  subunit gene in the growth and development of *Pleurotus eryngii*[J]. *Journal of Fungi*, 2023, 9(1): 69.
- [35] 卢园萍. 腺苷酸环化酶对金针菇菌丝生长和子实体形成的调控[D]. 福州: 福建农林大学博士学位论文, 2016.  
LU YP. Regulation of adenylate cyclase on mycelium growth and fruiting body formation of *Flammulina velutipes*[D]. Fuzhou: Doctoral Dissertation of Fujian Agriculture and Forestry University, 2016 (in Chinese).
- [36] YE D, DU F, ZOU YJ, HU QX. Transcriptomics analysis of primordium formation in *Pleurotus eryngii*[J]. *Genes*, 2021, 12(12): 1863.
- [37] 易勇. 水稻锌指蛋白 OsC3H54 互作蛋白鉴定[D]. 荆州: 长江大学硕士学位论文, 2021.  
YI Y. Identification of rice zinc finger protein OsC3H54 interacting protein[D]. Jingzhou: Master's Thesis of Yangtze University, 2021 (in Chinese).
- [38] LIU YH, KHAN AR, GAN YB. C<sub>2</sub>H<sub>2</sub> zinc finger proteins response to abiotic stress in plants[J]. *International Journal of Molecular Sciences*, 2022, 23(5): 2730.
- [39] 吴小梅, 张昕, 李南羿. 双孢蘑菇子实体不同发育时期的转录组分析[J]. *菌物学报*, 2017, 36(2): 193-203.  
WU XM, ZHANG X, LI NY. Transcriptome analysis of *Agaricus bisporus* fruiting at different stages[J]. *Mycosistema*, 2017, 36(2): 193-203 (in Chinese).
- [40] PELKMANS JF, VOS AM, SCHOLTMEIJER K, HENDRIX E, BAARS JJP, GEHRMANN T, REINDERS MJT, LUGONES LG, WÖSTEN HAB. The transcriptional regulator *c2h2* accelerates mushroom formation in *Agaricus bisporus*[J]. *Applied Microbiology and Biotechnology*, 2016, 100(16): 7151-7159.
- [41] ZHANG X, GUO XP, LEI CL, CHENG ZJ, LIN QB, WANG JL, WU FQ, WANG J, WAN JM. Overexpression of *SlCZFP1*, a novel TFIIIA-type zinc finger protein from tomato, confers enhanced cold tolerance in transgenic *Arabidopsis* and rice[J]. *Plant Molecular Biology Reporter*, 2011, 29(1): 185-196.
- [42] HAN GL, LU CX, GUO JR, QIAO ZQ, SUI N, QIU NW, WANG BS. C2H2 zinc finger proteins: master regulators of abiotic stress responses in plants[J]. *Frontiers in Plant Science*, 2020, 11: 115.
- [43] ZNAMEROSKI EA, CORADETTI ST, ROCHE CM, TSAI JC, IAVARONE AT, CATE JHD, GLASS NL. Induction of lignocellulose-degrading enzymes in *Neurospora crassa* by cellobextrins[J]. *Proceedings of the National Academy of Sciences of the United States of America*, 2012, 109(16): 6012-6017.
- [44] NAKAZAWA T, MIYAZAKI Y, KANEKO S, SHISHIDO K. Stimulative effects of light and a temperature downshift on transcriptional expressions of developmentally regulated genes in the initial stages of fruiting-body formation of the basidiomycetous mushroom *Lentinula edodes*[J]. *FEMS Microbiology Letters*, 2008, 289(1): 67-71.
- [45] KISHIKAWA A, HAMADA S, KAMEI I, FUJIMOTO Y, MIYAZAKI K, YOSHIDA M. A novel gene, Le-Dd10, is involved in fruiting body formation of *Lentinula edodes*[J]. *Archives of Microbiology*, 2022, 204(10): 1-13.
- [46] WU TJ, HU CC, XIE BG, WEI SL, ZHANG L, ZHU ZX, ZHANG ZY, LI SJ. A putative transcription factor LFC1 negatively regulates development and yield of

- winter mushroom[J]. *Applied Microbiology and Biotechnology*, 2020, 104(13): 5827-5844.
- [47] 乌凤章, 王贺新. 蛋白质泛素化介导的植物低温胁迫反应[J]. 生物技术通报, 2021, 37(6): 225-235.
- WU FZ, WANG HX. Low temperature stress response mediated by protein ubiquitination in plant[J]. *Biotechnology Bulletin*, 2021, 37(6): 225-235 (in Chinese).
- [48] GONG M, LI ZP, WAN JN, CHEN MJ, WANG H, SHANG JJ, ZHOU SC, TAN Q, WANG Y, BAO DP. Chilling stress reduced protein translation by the ubiquitination of ribosomal proteins in *Volvariella volvacea*[J]. *Journal of Proteomics*, 2020, 215: 103668.
- [49] HENSOLD JO, SWERDLOW PS, HOUSMAN DE. A transient increase in histone H2A ubiquitination is coincident with the onset of erythroleukemic cell differentiation[J]. *Blood*, 1988, 71(4): 1153-1156.
- [50] KANDA T, TANAKA N, TAKEMARU T. Ubiquitin immunoreactivity shows several proteins varying with development and sporulation in the basidiomycete *Coprinus cinereus*[J]. *Biochemistry and Cell Biology*, 1990, 68(7/8): 1019-1025.
- [51] YANG XQ, LIN RM, XU K, GUO LZ, YU H. Comparative proteomic analysis within the developmental stages of the mushroom white *Hypsizygus marmoreus*[J]. *Journal of Fungi*, 2021, 7(12): 1064.
- [52] CHAE E, TAN QKG, HILL TA, IRISH VF. An *Arabidopsis* F-box protein acts as a transcriptional co-factor to regulate floral development[J]. *Development*, 2008, 135(7): 1235-1245.
- [53] KRIZSÁN K, ALMÁSI É, MERÉNYI Z, SAHU N, VIRÁGH M, KÓSZÓ T, MONDO S, KISS B, BÁLINT B, KÜES U, BARRY K, CSEKLYE J, HEGEDÜS B, HENRISSAT B, JOHNSON J, LIPZEN A, OHM RA, NAGY I, PANGILINAN J, YAN JY, et al. Transcriptomic atlas of mushroom development reveals conserved genes behind complex multicellularity in fungi[J]. *Proceedings of the National Academy of Sciences of the United States of America*, 2019, 116(15): 7409-7418.
- [54] GONG M, HUANG TY, LI Y, LI JX, TANG LH, SU EZ, ZOU G, BAO DP. Multi-omics analysis of low-temperature fruiting highlights the promising cultivation application of the nutrients accumulation in *Hypsizygus marmoreus*[J]. *Journal of Fungi*, 2022, 8(7): 695.
- [55] WANG ST, LI WL, LIU L, QI H, YOU H. Biodegradation of decabromodiphenyl ethane (DBDPE) by white-rot fungus *Pleurotus ostreatus*: characteristics, mechanisms, and toxicological response[J]. *Journal of Hazardous Materials*, 2022, 424: 127716.
- [56] de GROOT PWJ, SCHAAP PJ, van GRIENSVEN LJLD, VISSER J. Isolation of developmentally regulated genes from the edible mushroom *Agaricus bisporus*[J]. *Microbiology*, 1997, 143(6): 1993-2001.
- [57] MURAGUCHI H, KAMADA T. A mutation in the *eln2* gene encoding a cytochrome P450 of *Coprinus cinereus* affects mushroom morphogenesis[J]. *Fungal Genetics and Biology*, 2000, 29(1): 49-59.
- [58] SHAH P, SWIATLO E. A multifaceted role for polyamines in bacterial pathogens[J]. *Molecular Microbiology*, 2008, 68(1): 4-16.
- [59] GHOSH UK, ISLAM MN, SIDDIQUI MN, CAO X, KHAN MR. Proline, a multifaceted signalling molecule in plant responses to abiotic stress: understanding the physiological mechanisms[J]. *Plant Biology*, 2022, 24(2): 227-239.
- [60] PELKMANS JF, LUGONES LG, WÖSTEN HAB. 15 fruiting body formation in basidiomycetes[M]//Growth, Differentiation and Sexuality. Cham: Springer International Publishing, 2016: 387-405.
- [61] TAO YX, CHEN RL, YAN JJ, LONG Y, TONG ZJ, SONG HB, XIE BG. A hydrophobin gene, *Hyd9*, plays an important role in the formation of aerial hyphae and primordia in *Flammulina filiformis*[J]. *Gene*, 2019, 706: 84-90.
- [62] HAMLETT CAE, SHIRTCLIFFE NJ, NEWTON MI, McHALE G, KOCH K. Passive water control at the surface of a superhydrophobic lichen[J]. *Planta*, 2011, 234(6): 1267-1274.
- [63] YOO SI, LEE HY, MARKKANDAN K, MOON S, AHN YJ, JI SM, KO J, KIM SJ, RYU H, HONG CP. Comparative transcriptome analysis identified candidate genes involved in mycelium browning in *Lentinula edodes*[J]. *BMC Genomics*, 2019, 20(1): 1-13.
- [64] LIU XB, XIA EH, LI M, CUI YY, WANG PM, ZHANG JX, XIE BG, XU JP, YAN JJ, LI J, NAGY LG, YANG ZL. Transcriptome data reveal conserved patterns of fruiting body development and response to heat stress in the mushroom-forming fungus *Flammulina filiformis*[J]. *PLoS One*, 2020, 15(10): e0239890.
- [65] CESUR A, YAMAMOTO R, ASADA Y, WATANABE A. Relationship between fruiting body development and

- extracellular laccase production in the edible mushroom *Flammulina velutipes*[J]. Biochemistry and Biophysics Reports, 2022, 29: 101204.
- [66] 陈辉, 郝海波, 赵静, 王倩, 隽加香, 陈明杰, 冯志勇, 张津京. 漆酶同工酶基因在斑玉蕈生长发育过程中的表达分析[J]. 菌物学报, 2020, 39(6): 1038-1048. CHEN H, HAO HB, ZHAO J, WANG Q, JUAN JX, CHEN MJ, FENG ZY, ZHANG JJ. Expression analysis of laccase isozyme gene during *Hypsizygus marmoreus* growth and development[J]. Mycosistema, 2020, 39(6): 1038-1048 (in Chinese).
- [67] FUKUDA K, HIRAGA M, ASAKUMA S, ARAI I, SEKIKAWA M, URASHIMA T. Purification and characterization of a novel exo- $\beta$ -1,3-1,6-glucanase from the fruiting body of the edible mushroom enoki (*Flammulina velutipes*)[J]. Bioscience Biotechnology and Biochemistry, 2008, 72(12): 3107-3113.
- [68] PARK YJ, BAEK JH, LEE S, KIM C, RHEE H, KIM H, SEO JS, PARK HR, YOON DE, NAM JY, KIM HI, KIM JG, YOON H, KANG HW, CHO JY, SONG ES, SUNG GH, YOO YB, LEE CS, LEE BM, KONG WS. Whole genome and global gene expression analyses of the model mushroom *Flammulina velutipes* reveal a high capacity for lignocellulose degradation[J]. PLoS One, 2014, 9(4): e93560.
- [69] KAMADA T, HAMADA Y, TAKEMARU T. Autolysis *in vitro* of the stipe cell wall in *Coprinus macrorhizus*[J]. Microbiology, 1982, 128(5): 1041-1046.
- [70] WESSELS JGH. Fruiting in the higher fungi[M]//Advances in Microbial Physiology. Amsterdam: Elsevier, 1993: 147-202.
- [71] KONNO N, SAKAMOTO Y. An endo- $\beta$ -1,6-glucanase involved in *Lentinula edodes* fruiting body autolysis[J]. Applied Microbiology and Biotechnology, 2011, 91(5): 1365-1373.
- [72] ZHOU YJ, KANG LQ, NIU X, WANG J, LIU ZH, YUAN S. Purification, characterization and physiological significance of a chitinase from the pilei of *Coprinopsis cinerea* fruiting bodies[J]. FEMS Microbiology Letters, 2016, 363(12): fnw120.
- [73] FANG HJ, ZHANG WM, NIU X, LIU ZH, LU CM, WEI H, YUAN S. Stipe wall extension of *Flammulina velutipes* could be induced by an expansin-like protein from *Helix aspersa*[J]. Fungal Biology, 2014, 118(1): 1-11.
- [74] HUANG QH, HAN X, MUKHTAR I, GAO LL, HUANG RM, FU LP, YAN JJ, TAO YX, CHEN BZ, XIE BG. Identification and expression patterns of fvexpl1, an expansin-like protein-encoding gene, suggest an auxiliary role in the stipe morphogenesis of *Flammulina velutipes*[J]. Journal of Microbiology and Biotechnology, 2018, 28(4): 622-629.
- [75] SAKAMOTO Y, ANDO A, TAMAI Y, MIURA K, YAJIMA T. Protein expressions during fruit body induction of *Flammulina velutipes* under reduced temperature[J]. Mycological Research, 2002, 106(2): 222-227.
- [76] SAKAMOTO Y. Protein expression during *Flammulina velutipes* fruiting body formation[J]. Mycoscience, 2010, 51(3): 163-169.
- [77] LOS DA, MURATA N. Membrane fluidity and its roles in the perception of environmental signals[J]. Biochimica et Biophysica Acta (BBA)-Biomembranes, 2004, 1666(1/2): 142-157.
- [78] FU YP, LIANG Y, DAI YT, YANG CT, DUAN MZ, ZHANG Z, HU SN, ZHANG ZW, LI Y. *De novo* sequencing and transcriptome analysis of *Pleurotus eryngii* subsp. *tuoliensis* (bailinggu) mycelia in response to cold stimulation[J]. Molecules, 2016, 21(5): 560.
- [79] ERNST R, EJSING CS, ANTONNY B. Homeoviscous adaptation and the regulation of membrane lipids[J]. Journal of Molecular Biology, 2016, 428(24): 4776-4791.
- [80] SEZGIN E, LEVENTAL I, MAYOR S, EGELING C. The mystery of membrane organization: composition, regulation and roles of lipid rafts[J]. Nature Reviews Molecular Cell Biology, 2017, 18(6): 361-374.
- [81] 解凡, 宋燕娇, 程冰, 孟国良, 郝金斌, 叶丽云, 吴小平. 低温胁迫诱导肺形侧耳合成麦角甾醇的通路分析[J]. 菌物学报, 2019, 38(12): 2221-2231. XIE F, SONG YJ, CHENG B, MENG GL, HAO JB, YE LY, WU XP. Cold stress induction of ergosterol biosynthesis in *Pleurotus pulmonarius*[J]. Mycosistema, 2019, 38(12): 2221-2231 (in Chinese).
- [82] WANG RH, MA PD, LI C, XIAO LG, LIANG ZS, DONG JE. Combining transcriptomics and metabolomics to reveal the underlying molecular mechanism of ergosterol biosynthesis during the fruiting process of *Flammulina velutipes*[J]. BMC Genomics, 2019, 20(1): 1-12.
- [83] PRIETO JA, ESTRUCH F, CÓRCOLES-SÁEZ I, del POETA M, RIEGER R, STENZEL I, RANDÉZ-GIL F. Pho85 and PI(4,5)P<sub>2</sub> regulate different lipid metabolic pathways in response to cold[J]. Biochimica et

- Biophysica Acta (BBA)-Molecular and Cell Biology of Lipids, 2020, 1865(2): 158557.
- [84] KNIGHT H, TREWAVAS AJ, KNIGHT MR. Cold calcium signaling in *Arabidopsis* involves two cellular pools and a change in calcium signature after acclimation[J]. *The Plant Cell*, 1996, 8(3): 489-503.
- [85] WU TH, YE ZW, GUO LQ, YANG XQ, LIN JF. *De novo* transcriptome sequencing of *Flammulina velutipes* uncover candidate genes associated with cold-induced fruiting[J]. *Journal of Basic Microbiology*, 2018, 58(8): 698-703.
- [86] 华霜. 低温胁迫白灵菇原基形成过程中 DNA 甲基化变异及调控基因研究[D]. 长春: 吉林农业大学博士学位论文, 2017.
- HUA S. Study on DNA methylation variation and regulatory genes during the formation of *Pleurotus nebrodensis* primordium under low temperature stress[D]. Changchun: Doctoral Dissertation of Jilin Agricultural University, 2017 (in Chinese).
- [87] LIU QB, DING YL, SHI YT, MA L, WANG Y, SONG CP, WILKINS KA, DAVIES JM, KNIGHT H, KNIGHT MR, GONG ZZ, GUO Y, YANG SH. The calcium transporter ANNEXIN<sub>1</sub> mediates cold-induced calcium signaling and freezing tolerance in plants[J]. *The EMBO Journal*, 2021, 40(2): e104559.
- [88] YUAN PG, YANG TB, POOVAIAH BW. Calcium signaling-mediated plant response to cold stress[J]. *International Journal of Molecular Sciences*, 2018, 19(12): 3896.
- [89] 吴雪昌, 胡森杰, 钱凯先. 酵母 HOG-MAPK 途径[J]. *细胞生物学杂志*, 2005(3): 247-252.
- WU XC, HU SJ, QIAN KX. HOG-MAPK pathway in yeast[J]. *Chinese Journal of Coll Biology*, 2005(3): 247-252 (in Chinese).
- [90] 金华燕. 蝇虫草组氨酸激酶基因(CmHK)和 Kexin 基因(CmKexin)的研究[D]. 南昌: 江西师范大学硕士学位论文, 2013.
- JIN HY. The Study of Histidine Kinase Gene (CmHK) and Kexin Gene (CmKexin) in *Cordyceps militaris*[D]. Nanchang: Master's Thesis of Jiangxi Normal University, 2013 (in Chinese).
- [91] SUZUKI I, LOS DA, MURATA N. Perception and transduction of low-temperature signals to induce desaturation of fatty acids[J]. *Biochemical Society Transactions*, 2000, 28(6): 628-630.
- [92] 李琼洁. 冷诱导金针菇原基形成的相关基因研究[D]. 广州: 华南农业大学硕士学位论文, 2016.
- LI QJ. Study on genes related to cold-induced primordium formation of *Flammulina velutipes*[D]. Guangzhou: Master's Thesis of South China Agricultural University, 2016 (in Chinese).
- [93] JIA HH, HAO LL, GUO XL, LIU SC, YAN Y, GUO XQ. A Raf-like MAPKKK gene, *GhRaf19*, negatively regulates tolerance to drought and salt and positively regulates resistance to cold stress by modulating reactive oxygen species in cotton[J]. *Plant Science*, 2016, 252: 267-281.
- [94] 解凡, 赵丽丽, 叶丽云, 吴小平. 肺形侧耳低温胁迫时期的转录组分析[J]. *菌物学报*, 2018, 37(12): 1598-1607.
- XIE F, ZHAO LL, YE LY, WU XP. Transcriptome analyses of *Pleurotus pulmonarius* after cold stress challenge[J]. *Mycosistema*, 2018, 37(12): 1598-1607 (in Chinese).
- [95] MATTEI AL, BAILLY N, MEISSNER A. DNA methylation: a historical perspective[J]. *Trends in Genetics*, 2022, 38(7): 676-707.
- [96] SOTO T, BELTRÁN FF, PAREDES V, MADRID M, MILLAR JBA, VICENTE-SOLER J, CANSADO J, GACTO M. Cold induces stress-activated protein kinase-mediated response in the fission yeast *Schizosaccharomyces pombe*[J]. *European Journal of Biochemistry*, 2002, 269(20): 5056-5065.
- [97] ITO E, UEMURA T. RAB GTPases and SNAREs at the trans-Golgi network in plants[J]. *Journal of Plant Research*, 2022, 135(3): 389-403.
- [98] YAN JJ, XIE B, ZHANG L, LI SJ, van PEER AF, WU TJ, CHEN BZ, XIE BG. Small GTPases and stress responses of vvrn1 in the straw mushroom *Volvariella volvacea*[J]. *International Journal of Molecular Sciences*, 2016, 17(9): 1527.
- [99] YAMADA M, SAKURABA S, SHIBATA K, TAGUCHI G, INATOMI S, OKAZAKI M, SHIMOSAKA M. Isolation and analysis of genes specifically expressed during fruiting body development in the basidiomycete *Flammulina velutipes* by fluorescence differential display[J]. *FEMS Microbiology Letters*, 2006, 254(1): 165-172.
- [100] MENG L, CHOU TS, JIANG SY, WANG L, ZHU MJ, MUKHTAR I, XIE BG, WANG W. Characterization and expression pattern analysis of pheromone receptor-like genes in winter mushroom *Flammulina filiformis*[J]. *Archives of Microbiology*, 2020, 202(10): 2671-2678.
- [101] WU TJ, HU CC, XIE BG, ZHANG L, YAN SJ, WANG W, TAO YX, LI SJ. A single transcription

- factor (PDD1) determines development and yield of winter mushroom (*Flammulina velutipes*)[J]. Applied and Environmental Microbiology, 2019, 85(24): e01735-19.
- [102] 刘芳, 王威, 谢宝贵. 金针菇菌丝与原基差异表达基因分析[J]. 食用菌学报, 2014, 21(1): 1-7.
- LIU F, WANG W, XIE BG. Comparison of gene expression patterns in the Mycelium and primordia of *Flammulina velutipes*, strain 1123[J]. Acta Edulis Fungi, 2014, 21(1): 1-7 (in Chinese).
- [103] REIS FS, BARROS L, MARTINS A, FERREIRA ICFR. Chemical composition and nutritional value of the most widely appreciated cultivated mushrooms: an inter-species comparative study[J]. Food and Chemical Toxicology, 2012, 50(2): 191-197.
- [104] GONG M, WANG Y, SU EZ, ZHANG JG, TANG LH, LI ZP, ZHANG LL, ZOU G, WAN JN, BAO DP. The promising application of a  $\beta$ -glucosidase inhibitor in the postharvest management of *Volvariella volvacea*[J]. Postharvest Biology and Technology, 2022, 185: 111784.
- [105] 张国良. 低温处理对阿魏茹菌丝体生理生化及营养积累的影响[D]. 乌鲁木齐: 新疆农业大学硕士学位论文, 2013.
- ZHANG GL. Effects of low temperature treatment on physiology, biochemistry and nutrient accumulation of *Pleurotus ferulae* mycelium[D]. Urumqi: Master's Thesis of Xinjiang Agricultural University, 2013 (in Chinese).
- [106] 吕静, 王伟科, 袁卫东, 陆娜, 宋吉玲, 周祖法. 温度对秀珍菇生长发育及胞外酶活性的影响[J]. 浙江大学学报(农业与生命科学版), 2020, 46(2): 161-167.
- YAN J, WANG WK, YUAN WD, LU N, SONG JL, ZHOU ZF. Effects of temperature on the growth, development and extracellular enzyme activities of *Pleurotus pulmonarius*[J]. Journal of Zhejiang University (Agriculture & Life Sciences Edition), 2020, 46(2): 161-167 (in Chinese).
- [107] LIU YY, MA XB, LONG Y, YAO S, WEI CZ, HAN X, GAN BC, YAN JJ, XIE BG. Effects of  $\beta$ -1,6-glucan synthase gene (*FfGS6*) overexpression on stress response and fruit body development in *Flammulina filiformis*[J]. Genes, 2022, 13(10): 1753.
- [108] YAMADA M, SAKURABA S, SHIBATA K, INATOMI S, OKAZAKI M, SHIMOSAKA M. Cloning and characterization of a gene coding for a hydrophobin, *Fv-hyd1*, specifically expressed during fruiting body development in the basidiomycete *Flammulina velutipes*[J]. Applied Microbiology and Biotechnology, 2005, 67(2): 240-246.
- [109] OHM RA, de JONG JF, LUGONES LG, AERTS A, KOTHE E, STAJICH JE, de VRIES RP, RECORD E, LEVASSEUR A, BAKER SE, BARTHOLOMEW KA, COUTINHO PM, ERDMANN S, FOWLER TJ, GATHMAN AC, LOMBARD V, HENRISSAT B, KNABE N, KÜES U, LILLY WW, et al. Genome sequence of the model mushroom *Schizophyllum commune*[J]. Nature Biotechnology, 2010, 28(9): 957-963.
- [110] HOLTHUIS JCM, MENON AK. Lipid landscapes and pipelines in membrane homeostasis[J]. Nature, 2014, 510(7503): 48-57.
- [111] GIANFREDA L, XU F, BOLLAG JM. Laccases: a useful group of oxidoreductive enzymes[J]. Bioremediation Journal, 1999, 3(1): 1-26.
- [112] 徐桂红, 赵心清, 李宁, 白凤武. 锌离子提高絮凝酶母乙酸胁迫耐受性[J]. 化工学报, 2012, 63(6): 1823-1829.
- XU GH, ZHAO XQ, LI N, BAI FW. Improvement of acetic acid tolerance of self-flocculating yeast by zinc supplementation[J]. CIESC Journal, 2012, 63(6): 1823-1829 (in Chinese).
- [113] 吴又多. 锌离子对丙酮丁醇梭菌发酵的调控作用及机制[D]. 大连: 大连理工大学博士学位论文, 2017.
- WU YD. Regulation and mechanism of zinc ion on *Clostridium acetobutylicum* fermentation[D]. Dalian: Doctoral Dissertation of Dalian University of Technology, 2017 (in Chinese).
- [114] ZHAO XQ, BAI FW. Zinc and yeast stress tolerance: micronutrient plays a big role[J]. Journal of Biotechnology, 2012, 158(4): 176-183.
- [115] ISHIZAKI T, SHISHIDO K. Decreased zinc ion accumulation by the basidiomycete *Lentinus edodes* over-expressing *L. edodes priA* gene[J]. FEMS Microbiology Letters, 2000, 193(1): 111-115.