

灰树花活性多糖构效关系研究进展

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摘 要: 灰树花是一种珍贵的食药菌, 具有降血糖、抗肿瘤、免疫调节和抗病毒等多种生物活性。灰树花多糖是其主要的活性成分, 多糖的生物活性与多糖的结构密切相关。本文综述了从20世纪80年代起国内外已报道的灰树花活性多糖的结构表征。部分研究认为多糖的降血糖活性可能与 β -1,6-葡聚糖主链化学结构相关, 而灰树花多糖结构为 β -1,6主链或 β -1,3主链葡聚糖时具有较好的抗肿瘤活性。然而多糖结构异常复杂, 精细结构的解析困难, 导致目前灰树花多糖结构表征一般止步于单糖组成、分子量、糖苷键类型、分支结构和粗略分子链构象, 但二维核磁(two dimensional nuclear magnetic resonance, 2D-NMR)和高分辨质谱联用等技术的发展将有助于解开灰树花多糖构效关系, 并为灰树花活性多糖的开发利用提供理论依据。

关键词: 灰树花; 多糖; 降血糖; 抗肿瘤; 结构表征; 构效关系

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Advances in structure-activity relationship of polysaccharides from *Grifola frondosa*

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Abstract: *Grifola frondosa* is a precious edible and medicinal fungus with anti-tumor, immunomodulatory, hypoglycemic, and antiviral activities. Polysaccharides are the main active component of *G. frondosa*, with the biological activity associated with the structure. We reviewed the structural characterization of active polysaccharides from *G. frondosa*. Some studies suggest that the blood sugar-lowering activity might be related to the chemical structure with β -1,6-glucan as the main chain, and the polysaccharides have better antitumor activity when the main chain is β -1,6-glucan or β -1,3-glucan. However, the complex structure makes it difficult to dissect the fine structure of *G. frondosa* polysaccharides. As a result, the structural characterization of *G. frondosa* polysaccharides is generally limited to monosaccharide composition, molecular weight, glycosidic bond type, branching structure, and rough molecular chain conformation. The advancing 2D-NMR and high-resolution mass spectrometry will help to disclose the structure-activity relationship and provide a theoretical basis for the development and utilization of *G. frondosa* polysaccharides.

Keywords: *Grifola frondosa*; polysaccharides; hypoglycemic; anti-tumor; structural characterization; structure-activity relationship

灰树花(*Grifola frondosa*)隶属于担子菌门层菌纲非褶菌目多孔菌科树花菌属^[1],是一种珍贵的食药兼用蕈菌^[2]。现代医学表明,灰树花多糖类物质(多糖、糖蛋白、蛋白多糖)具有显著的生理活性^[3],如抗肿瘤^[4]、调节机体免疫力^[5]、抗氧化^[6]、降血糖降脂^[7]、抗病毒^[8]和抗辐射^[9]等。

虽然多糖类活性物质的功能显著,但天然多糖本身分子量大且一级结构及空间结构复杂,尤其是天然多糖多和蛋白质以非共价键的形式结合在一起,使得多糖的结构解析异常困难,导致活性多糖的结构和活性关系不明晰。多糖的生物活性与其自身结构密切相关,多糖的结构在一定程度上决定了其生物活性。多糖的分子量、单糖组成^[10]、糖苷键连接方式^[4]、

分支度、支链类型及位置和构象^[11]等都会影响多糖的生物活性。活性多糖构效关系的不明晰严重制约其活性多糖的产业化进程。

从 20 世纪 80 年代开始国内外就开始了对灰树花多糖的探究。本文归纳了灰树花降血糖、抗肿瘤、免疫调节、抗病毒等活性多糖的一级结构及其作用机制,以期为进一步解开灰树花活性多糖构效关系提供参考。

1 灰树花降血糖活性多糖一级结构表征及其作用机制研究

降血糖灰树花多糖的分子质量、结构或组成见表 1。

表 1 降血糖灰树花多糖的分子质量、结构或组成

Table 1 Molecular weight, structure or composition of <i>G. frondosa</i> polysaccharides with hypoglycemic activities									
名称	提取部位	分子量	糖:蛋白	单糖组成	结构	实验动物模型	作用机制	血糖降低率	参考文献
Name	Extractive fraction	Mw (kDa)	Polysaccharides: proteins	Monosaccharide composition	Structure or construction	Experimental animal model	Mechanism of action	Hypoglycemic rate (%)	References
X-组分	子实体	200	65:35	不详	α -1,4 分支的	雌性自发高血	不详	50.0	[12-13]
X- fraction	Fruiting bodies			Uncertain	β -1-6-葡聚糖	压小鼠	Uncertain		
					β -1,6 main chain with	(KK-A ^y) female spontaneously			
SX-组分	子实体	20	10:90-25:75	不详	α -1,4 branches	hypertensive mice	改善糖耐量, 对外源性胰岛素的	11.9	[14-15]
SX- fraction	Fruiting bodies			Uncertain	不详	自发高血压	敏感性增强, 改善胰岛素敏感性		
					Uncertain	大鼠	Improvement of glucose tolerance and the sensitivity to exogenous insulin, and enhancement of insulin sensitivity		
MT- α -glucan	子实体	40-45	不详	葡萄糖	α -D-葡聚糖	雌性自发高血	增加胰岛素敏感性, 改善外周靶	22.0	[16-17]
	Fruiting bodies		Uncertain	Glc	α -D-glucan	压小鼠	组织的胰岛素抵抗		
						(KK-A ^y) female spontaneously and amelioration of insulin resistance in hypertensive mice peripheral tar-get tissues	Increase of insulin sensitivity		
GFP	子实体	45.3	不详	葡萄糖:鼠李糖:阿拉伯糖:木糖:甘露糖:半乳糖=31.29:24.74:5.1:3.42:6.89	β -1,4 和	体外/人 HepG2 细胞	通过 Akt/GSK-3 通路增加葡萄糖代谢, 刺激细胞内糖原合成	*	[18]
	Fruiting bodies		Uncertain		β -1-6-葡聚糖	细胞	代谢, 刺激细胞内糖原合成		
					β -1,6 main chain with	<i>In vitro</i> /Human liver cancer cells HepG2	Increase of glucose metabolism of and stimulation of intracellular glycogen synthesis through the Akt/GSK-3 pathway		
				Glc:Rha:Ara:Xyl:Man:Gal=31.29:24.74:5.1:3.42:6.89	α -1,4 branches				
GFP	子实体	18.18-15 850	不详	葡萄糖:甘露糖:半乳糖:鼠李糖:葡萄糖醛酸:半乳糖醛酸:岩藻糖=27.59:25.49:15.02:5.18:9.49:7.30:9.92	不详	STZ 和 HFD 联合诱导的糖尿	通过改变肠道菌群和调节肝脏糖脂代谢相关基因	*	[19]
	Fruiting bodies		Uncertain		Uncertain	病小鼠	脂代谢相关基因		
						Diabetic mice induced by STZ and HFD	Though modulation of gut microbiota and regulation of hepatic glycolipid metabolism related genes		
				Glc:Man:Gal:Rha:GlcUA:GalUA:Fuc=27.59:25.49:15.02:5.18:9.49:7.30:9.92					

(待续)

(待续)

GFWE	子实体 Fruiting bodies	不详	不详	半乳糖:鼠李糖:甘露糖:葡萄糖:岩藻糖=22.66:12.10:11.88:5.81:4.51 Gal:Rha:Man:Glc:Fuc=22.66:12.10:11.88:5.81:4.51	STZ 和 HFD 联合诱导的糖尿病 Wistar 大鼠 Diabetic Wistar rats induced by STZ and HFD	上调 AMPK- α 、PPAR- α 和 GK 的相对 mRNA 和蛋白表达水平, 下调 SREBP-1c 和 ACC, 提高盲肠总胆汁酸和短链脂肪酸水平 Up-regulation of the relative mRNA and protein expression levels of adenine monophosphate activated protein kinase- α , peroxisome proliferator-activated receptors- α , and glucokinase, while down-regulation of sterol regulatory element-binding transcription factor-1c and acetyl CoA carboxylase; enhancement of the levels of total bile acids and short-chain fatty acids in the cecum	* [20]	(续表 1)
GFP-N	子实体 Fruiting bodies	12 600	不详	葡萄糖:阿拉伯糖:甘露糖=49.70:3.79:1.00 Glc:Ara:Man=49.70:3.79:1.00	STZ 和 HFD 联合诱导的糖尿病 ICR 小鼠 Diabetic ICR mice induced by STZ and HFD	激活 IRS1、PI3K 和 GLUT4, 抑制 JNK, 改善口服糖耐量, 减轻胰岛素抵抗, 保护免受肾损伤, 减轻炎症反应并且调节 T2DM 小鼠肠道菌群 Activation of insulin receptor substrate 1, phosphatidylinositol-3-kinase, and glucose31 transporter 4 and inhibition of c-Jun N-terminal kinase 1/2 and improvement of oral glucose tolerance, alleviation of insulin resistance, and protection against liver and kidney injury with reduced inflammation and modulation of gut microbiota in diabetic mice	* [7]	(待续)
GFP-W	子实体 Fruiting bodies	66.1	不详	木糖:甘露糖:岩藻糖:葡萄糖:半乳糖:半乳糖=1.37:1.36:1.22:1.10:1.00 Xyl:Man:Fuc:Glc:Gal=1.37:1.36:1.22:1.10:1.00	人 HepG2 细胞 Human liver cancer cells HepG2	改善 IRS1、PI3K 的 mRNA 和蛋白表达, 上调 GLUT4, 下调 JNK1, 显著增加地塞米松诱导的胰岛素抵抗 HepG2 细胞对葡萄糖的摄取 Improvement of the uptake of glucose in dexamethasone induced insulin resistant HepG2 cells by improving the mRNA and protein expression of insulin receptor	* [21]	(待续)

(续表 1)

F2	子实体 Fruiting bodies	452	62.5:37.5	葡萄糖:甘露糖:半乳糖:木糖:阿拉伯糖:鼠李糖:核糖=26.74:22.79:16.76:16.02:14.29:2.05:1.35 Glc:Man:Gal:Xyl:Ara:Rha:Rib=26.74:22.79:16.76:16.02:14.29:2.05:1.35 核糖:阿拉伯糖:木糖=74.73:14.20:11.08 Rib:Ara:Xyl=74.73:14.20:11.08	Glcp-(1→,→2,6)-α-Galp-(1→,→2)-α-Manp-(1→, and→3)-α-L-Fucp-(1→含 β 糖苷键相连的吡喃环 Pyran rings linked by β glycosidic bonds STZ 和 HFD 联合诱导的糖尿病 SD 大鼠 STZ and HFD induced diabetes in SD rats 上调 IR (Try1361)蛋白磷酸化水平 * 和下调 IRS1 (Ser307)蛋白磷酸化水平, 改善胰岛素抵抗 Improvement of insulin sensitivity as a result of the increased protein levels of phospho-IR (Try 1361) and decreased levels of phospho-IRS-1 (Ser307)	[22-23]
F3			78.3:21.7	核糖:阿拉伯糖:木糖=74.73:14.20:11.08 Rib:Ara:Xyl=74.73:14.20:11.08		
GF5000	子实体 Fruiting bodies	5	38.2:37.8	葡萄糖:阿拉伯糖:甘露糖:核糖:鼠李糖:半乳糖:半乳糖=47.06:30.05:18.89:1.93:1.65:0.26:0.16 Glc:Ara:Man:Rib:Rha:Xyl:Gal=47.06:30.05:18.89:1.93:1.65:0.26:0.16	不详 Uncertain ALX 和 HFD 联合诱导的糖尿病雄性 SD 大鼠 Diabetic male SD rats induced by ALX and HFD 通过调节肠道菌群的组成来抑制 TLR4/MyD88/NF-κB 途径进而缓解炎症水平, 改善胰岛素抵抗 Improvement of insulin resistance is relation of modulating composition of gut microbiota, which may help to alleviate inflammation by inhibition of TLR4/MyD88/NF-κB pathway	[24]
GFP-2	菌丝体 Mycelia	45.3	不含蛋白、核酸 Contains no protein or nucleic acid	不详 Uncertain 含 β 型糖苷键和吡喃糖环 Contain β glycosidic bonds and pyranose ring 人 HepG2 细胞 Human liver cancer cells HepG2 提高糖原代谢能力, 活化胰岛素抵抗信号传导途径, 促进机体糖代谢能力 Improvement of glycogen metabolism, activation of insulin resistance signal transduction pathway, and promotion of glucose metabolism	[25]	

注: *: 文献中未标明具体数值, 但与糖尿病模型组相比具有统计学差异; STZ: 链脲佐菌素; HFD: 高脂饮食; ALX: 四氧嘧啶; JNK: c-Jun 氨基末端激酶; PI3K: 磷脂酰肌醇-3-激酶; GLUT4: 葡萄糖转运体 4; AMPK-α: 腺嘌呤核苷酸活化蛋白激酶-α; PPAR-α: 过氧化物酶体增殖物激活受体-α; GK: 葡萄糖激酶; SREBP-1c: 甾醇调控元素结合转录因子-1c; ACC: 乙酰辅酶 A 羧化酶

Note: *: Specific values are not indicated, but there are statistical differences compared with the diabetes model group; STZ: Streptozocin; HFD: High fat diet; ALX: Alloxan; JNK: c-Jun N-terminal kinase; PI3K: Phosphatidylinositol-3-kinase; GLUT4: Glucose transporters 4; AMPK-α: Adenine monophosphate activated protein kinase-α; PPAR-α: Peroxisome proliferator-activated receptors-α; GK: Glucokinase; SREBP-1c: Sterol regulatory element-binding transcription factor-1c; ACC: Acetyl CoA carboxylase.

1.1 灰树花降血糖活性多糖的单糖组成及其分子量

灰树花降血糖多糖既可以来源于子实体,也可以从发酵的菌丝体(孢内)或发酵液(孢外)中获取,但绝大多数是从灰树花子实体中分离提取获得。除了 MT- α -glucan^[16-17]外,灰树花降血糖活性多糖均为杂多糖,其单糖组成多为葡萄糖、甘露糖、葡萄糖、半乳糖、阿拉伯糖和木糖。几乎所有的降血糖活性多糖都含有葡萄糖,仅有本团队从灰树花子实体获得的活性多糖 F3 的单糖组成为核糖、阿拉伯糖和木糖^[22-23]。

灰树花活性多糖的降血糖活性与其分子量也有一定的关系。研究表明,多糖分子量较低则无法形成多糖产生活性的聚合结构,但分子质量太大则不利于多糖穿过多重细胞膜进入生物体内而发挥生物学活性^[26]。一般而言,把较高分子量的多糖降解为较低的分子量能显著提高其生物活性^[27]。然而从表 1 可以看出,灰树花降血糖活性多糖的分子量范围跨度较大,从 20 kDa 的灰树花 SX-组分^[14]到 12 600 kDa 的 GFP-N^[7],都表现出较好的降血糖活性。

1.2 灰树花降血糖活性多糖的糖苷键连接方式

多糖通常是由 10 个以上单糖通过糖苷键连接而成的生物大分子物质,糖苷键的连接方式对多糖的生物活性有重要的影响^[28]。1994 年, Kubo 等^[12]水提醇沉得到的灰树花子实体糖蛋白聚合物 X-组分(糖:蛋白=65:35),其结构为具有 α -1,4 支链的 β -1,6-葡聚糖主链,对自发性糖尿病 KK-A^y 小鼠口服给药,具有显著的降血糖活性,推测其高活性可能与 β -1,6-葡聚糖主链化学结构相关。2014 年 Ma 等^[18]提取得到的灰树花子实体多糖 GFP (分子量为 45.3 kDa)主链结构主要由 β -1,4 和 β -1,6-葡聚糖组成。

1.3 灰树花降血糖活性多糖的作用机制

1.3.1 调节胰岛素信号转导通路,增强胰岛素受体(insulin receptor, IR)和胰岛素受体底物 1 (insulin receptor substrate 1, IRS1)的表达

IR 酪氨酸磷酸化水平的降低和 IRS1 丝氨酸磷酸化水平的增加会导致胰岛素信号通路被阻断,引发胰岛素抵抗,进而造成 2 型糖尿病(diabetes mellitus type 2, T2DM)。研究者^[22-23]通过阴离子交换层析从灰树花子实体中分离纯化获得降血糖活性多糖 F2 和 F3,发现 F2 和 F3 均可上调 IR (Try1361)蛋白磷酸化水平并下调 IRS1 (Ser307)蛋白磷酸化水平,从而改善胰岛素抵抗,进而降低 T2DM 大鼠的空腹血糖水平。在体外高糖刺激培养的骨骼肌 L6 细胞中,灰树花 SX-组分(分子量为 20 kDa)也可通过增加 IR (γ)的磷酸化水平、降低 IRS1 (γ)的磷酸化、激活受损的胰岛素信号转导通路、增强胰岛素敏感性,从而增加葡萄糖摄取^[14,29]。

灰树花多糖 GFP-N 能通过激活 IRS1、phosphatidylinositol-3-kinase (PI3K)和 glucose transporters (GLUT4)等,调节 IRS1/PI3K 和 c-Jun N-terminal kinase (JNK)信号通路,改善肝脏胰岛素抵抗,发挥其降血糖作用^[7]。灰树花子实体多糖 GFP 能激活 HepG2 细胞膜 IRS1 信号,上调 Akt (蛋白激酶) (Ser473)的蛋白磷酸化表达水平并抑制糖原合成酶激酶(GSK-3)的表达水平,通过 Akt/GSK-3 途径促进葡萄糖代谢,刺激细胞内糖原合成^[18]。灰树花多糖 GFP-W 通过改善 IRS1、PI3K、GLUT4 的 mRNA 和蛋白表达上调,以及 JNK 的 mRNA 和蛋白表达下调来增加葡萄糖摄入,从而改善胰岛素抵抗和血糖升高^[21]。

1.3.2 抑制 α -葡萄糖苷酶活性

α -葡萄糖苷酶可分解二糖生成葡萄糖,从而升高血液的血糖值。通过抑制 α -葡萄糖苷酶

可减少葡萄糖生成量,从而抑制血糖升高。

MT- α -葡聚糖和 GF-H 具有明显的降血糖功能,其药效作用发挥与抑制 α -葡萄糖苷酶活性有关^[16-17,30-31]。

1.3.3 调节肠道菌群

对链脲佐菌素(streptozocin, STZ)、高脂饮食(high fat diet, HFD)联合诱导的糖尿病小鼠, HFD 诱导的 T2DM 小鼠灌胃给药不同的灰树花多糖 GFP,发现其均可通过调节肠道菌群、调节肝脏糖脂代谢相关基因的 mRNA 表达以及与胆固醇代谢相关基因的表达水平,增强肝内胆汁酸(bile acid, BA)合成和排泄,达到降血糖的效果^[19,32]。对 STZ、HFD 联合诱导 T2DM 小鼠灌胃给药灰树花多糖 GFP-N 发现,实验小鼠肠道菌群中 *Bacteroidetes* 丰度升高,而 *Firmicutes* 和 *Proteobacteria* 丰度降低,通过调节菌群丰度和细菌结构从而降低小鼠的血糖水平^[7]。

Xiao 等^[24]采用超滤法截留灰树花子实体水提取物中分子量大于 5 kDa 的大分子组分 GF 5000,通过测定灰树花 GF 5000 干预的 T2DM 大鼠肠道内容物的细菌 16S rRNA 基因的 V3-V4 区间序列并进行生物信息学分析,推测灰树花改善胰岛素抵抗的作用,与其通过改善肠道菌群的组分来抑制 toll 样受体 4/骨髓分化原发性反应 88/活化 B 细胞的核因子-轻链增强子(toll-like receptors, TLR4/myeloid differentiation primary response gene 88, MyD88/nuclearfactor κ B, NF- κ B)途径中关键因子的表达进而抑制靶器官的炎症水平密切相关。

2 灰树花抗肿瘤活性多糖一级结构表征及其作用机制研究

抗肿瘤灰树花多糖的分子质量、结构或组成见表 2。

2.1 灰树花抗肿瘤活性多糖的单糖组成及其分子量

与灰树花降血糖活性多糖一样,子实体也是灰树花抗肿瘤活性多糖的主要提取部位。抗肿瘤的灰树花活性多糖以杂多糖居多,其单糖组成主要为葡萄糖、半乳糖、甘露糖和木糖等(表 2)。除了灰树花多糖 GFPW (分子量为 15.7 kDa)的单糖组成不含葡萄糖外,其余的抗肿瘤活性多糖均含有葡萄糖^[48]。

多糖抗肿瘤活性多糖分子量分布范围尚无定论,灰树花抗肿瘤活性多糖的分子量范围广泛(1.79–12 600 kDa),而且大部分为较高分子量的多糖。有研究指出,相对分子质量太低的多糖生物活性较低甚至无活性,较高分子量的多糖才能够维持其空间构象,但陈向东等^[55]提取获得的灰树花菌丝体多糖 GFP 2 分子量仅为 2.6 kDa,荷瘤小鼠肿瘤生长抑制实验结果表明,其仍具有良好的抗肿瘤活性。

2.2 灰树花抗肿瘤活性多糖的糖苷键连接方式

灰树花多糖抗肿瘤活性如表 2 所示,灰树花多糖的结构为 β -1,6 主链或 β -1,3 主链葡聚糖时具有较好的抗肿瘤活性。灰树花菌丝体多糖 Grifolan (GRN)的结构为具有 β -1,6 分支的 β -1,3-葡聚糖,通过体外对 RAW 264.7 巨噬细胞的抑制实验表明,其能够激活巨噬细胞促进细胞因子的产生,增强免疫系统对肿瘤细胞的免疫应答^[54]。1987 年,日本的 Kodama 等从灰树花子实体里获得了一种不溶于酸、可溶于碱的蛋白多糖聚合物 D-组分(蛋白含量约 30%,分子量为 1 200–2 000 kDa),结构为具有 β -1,3 分支的 β -1,6-葡聚糖^[57],经腹腔注射和口服给药均能对荷瘤小鼠具有很强抗肿瘤作用^[38]。

另外,从表 2 还可以看出,多数活性多糖具有支链结构,因此可以推测多糖的支化度对

表 2 抗肿瘤灰树花多糖的分子质量、结构或组成

Table 2 Molecular weight, structure or composition of <i>G. frondosa</i> polysaccharides with anti-tumor activities									
Name	提取部位 Extractive fraction	分子量 Mw (kDa)	糖:蛋白 Polysaccharides: proteins	单糖组成 Monosaccharide composition	多糖分子结构 Structure or construction	动物/细胞类型 Animal/Cell type	作用机制 Mechanism of action	抑瘤率 Anti-tumor rate (%)	参考文献 References
Grifolan-7N	子实体 Fruiting bodies	1 200	不详 Uncertain	葡萄糖 Glc	β -1,3-葡萄糖 β -1,3-glucan	雄性 ICR 小鼠/180 肉瘤腹水细胞肿瘤 Male ICR mice/Sarcoma 180 tumor ascites cells	不详 Uncertain	97.00	[33]
AP-组分	子实体	不详	不详	不详	带分支的 β -1,3-葡 聚糖	雄性 ICR 小鼠/180 肉瘤腹水细胞肿瘤	不详 Uncertain	95.00	[34]
AS-组分	Fruiting bodies	Uncertain	Uncertain	Uncertain	β -1,3-glucan branched	Male ICR mice/Sarcoma 180 tumor ascites cells	Uncertain	59.00	
AS- component			糖:蛋白质:糖醛 酸=0.75:0.05:0.10 Sugar:protein: uronic acid= 0.75:0.05:0.10	不详 Uncertain					
CF-1	子实体 Fruiting bodies	不详 Uncertain	不详 Uncertain	葡萄糖、木糖 和甘露糖 Glc, Xyl and Man	β -1,6 分支的 β -1,3-葡萄糖 β -1,6-branched β -1,3-glucan	雄性 ICR 小鼠/180 肉瘤腹水细胞肿瘤 Male ICR mice/Sarcoma 180 tumor ascites cells	不详 Uncertain	98.80	[35]
GRN-LE	子实体 Fruiting bodies	5	91.0:0.3- 84.0:1.8	不详 Uncertain	β -1,6 分支的 β -1,3-葡萄糖 β -1,6-branched β -1,3-glucan	雄性 ICR 小鼠/180 肉瘤腹水细胞肿瘤 Male ICR mice/Sarcoma 180 tumor ascites cells	不详 Uncertain	99.00	[36-37]
D-组分	子实体	1 200-	70:30	半乳糖、甘露 糖、葡萄糖和 N-乙酰氨基葡 萄糖和岩藻糖 Gal, Man, Glc and N- acetylglucosamine and Fuc	β -1,3 分支的 β -1,6-葡萄糖 β -1,3 branched β -1,6-glucan	雄性 ICR 小鼠/180 肉瘤腹水细胞肿瘤 Male ICR mice/ Sarcoma 180 tumor ascites cells	通过激活细胞免疫抑制 小鼠异体肿瘤的生长 Immunosuppression of allogeneic tumor growth in mice by activating cells	>69.10	[38-41]
D- component	Fruiting bodies	2 000							

(待续)

(续表 2)									
MD-组分 MD- component L-MD-组分 L-MD- component	子实体 Fruiting bodies	1 000	80:20~99:1	葡萄糖 Glc	β -1,3 分支的 β -1,6-葡聚糖 β -1,3 branched β -1,6-glucan	雄性 C3H/HeJ 小鼠/ MM-46 肿瘤细胞 Male C3H/HeJ mice/ MM-46 carcinoma cells	增强免疫激活 Enhancement of immune activation	54.60	[41]
		250	不详 Uncertain					67.90	[42]
	子实体 Fruiting bodies	20	84:16	葡萄糖 Glc	β -1,6 和 β -1,3 分支 的 β -D-葡聚糖 β -1,6 and β -1,3 branches of β -D-glucan	雄性 C3H/HeJ 小鼠/小 鼠巨噬细胞细胞系 J774.1 Male C3H/HeJ mice/ Murine macrophage cell line J774.1	提高 TNF- α 和 IL-12 的 生产和小鼠巨噬细胞 J774.1 的抗原呈递 Increase of TNF- α and IL-12 production and antigen presentation of mouse macrophage J774.1	*	[43]
			500~250	不详 Uncertain	葡萄糖、木糖、 岩藻糖、甘露 糖、半乳糖 Glc, Xyl, Fuc, Man, and Gal	每隔 5 个葡萄糖以 β -1,6 连接的 β -1,3- 葡聚糖 β -1,3-glucan branched with β -1,6 linkage and average chain length of 5 residues	雌性 ICR/JCL 小鼠/ 180 肉瘤腹水细胞肿瘤 Female ICR/JCL mice/Sarcoma 180 tumor ascites cells	不详 Uncertain	50.00
MZF	子实体 Fruiting bodies	23	46:54	半乳糖:甘露糖: 岩藻糖:葡萄糖 =1.24:1.00:0.95: 0.88 Gal:Man:Fuc: Glc=1.24:1.00: 0.95:0.88	α -1,6-葡萄糖、 α -1,3- 岩藻糖、 α -1,6-甘露 糖、 β -1,3-葡萄糖、 β -1,3,6-葡萄糖组成	雌性 BALB/cA 小鼠/ Colon-26 癌细胞 Female BALB/cA mice/Colon-26 carcinoma cells	增加肿瘤部位 T 细胞的 积累, 诱导细胞免疫和 肿瘤消退 Increase of T cell accumulation in tumor site, inducement of cellular immunity and tumor regression	47.60	[45]
LMw-GFP	子实体 Fruiting bodies	1.79	糖:蛋白质:糖醛 酸=98.83:0.34: 0.21 Carbohydrates: proteins:uronic acid=98.83: 0.34:0.21	葡萄糖 Glc	α -D-葡萄糖、 α -1,4- 葡萄糖和 α -1,3,-葡 萄糖组成 α -D-glucose, α -1,4-glucose and α -1,3,-glucose composition	雌性 BALB/c 小鼠/ H22 肝癌细胞 Female BALB/c mice/H22 heptoma cells	保护免疫器官, 增强 NK 细胞、巨噬细胞和 淋巴细胞的活性, 通过 线粒体凋亡途径诱导 H22 实体瘤细胞凋亡 Protection of immune organs, and enhancement	40.10	[46]
(待续)									

(待续)

(续表 2)

GFAP	子实体 Fruiting bodies	644.9	糖含量=94.28% Carbohydrates content= 94.28%	葡萄糖:半乳糖:甘露糖: 甘露糖=2.18: 0.23:1.00 Glc:Gal:Man= 2.18:0.23:1.00	β -1,3-D-葡萄糖和 α -1,3-D-甘露糖组成 β -1,3-glucose and α -1,3-D-mannose	雌性 BALB/c 小鼠/ 小鼠 H22 肝癌细胞 Female BALB/c mice/Murine H22 hepatocarcinoma cells	of the killing activity of NK cells, the phagocytosis of peritoneal macrophages, inducing the tumor cells apoptosis <i>in vivo</i> via mitochondrial apoptotic pathway 显著提高 NK 细胞、巨 噬细胞、CD19 ⁺ B 细胞 和 CD4 ⁺ T 细胞的活性, 导致 H22 细胞的凋亡通 过 G0/G1 受阻 Improvement of the activities of NK cells, macrophages, CD19 ⁺ B cells and CD4 ⁺ T cells, leading to the apoptosis of H22 cells via G0/G1 phase arrested	36.72 [47]
GPII	子实体 Fruiting bodies	6.9-33.0	不详 Uncertain	葡萄糖:半乳糖:甘露糖: 甘露糖=5.04: 2.61:1.00 Glc:Gal:Man= 5.04:2.61:1.00 so on	1,6-D-甘露糖, 1,4-葡萄糖等 1,6-D-mannose, 1,4-glucose and so on	雌性 ICR 小鼠/Anti- mouse TLR2, TLR4/Heps 肿瘤细胞 Female ICR mice/Anti-mouse TLR2, TLR4/Heps tumor cells	增加胸腺和脾脏的相对 重量以及血清 TNF- α 和 IL-12 水平, 通过 TLR4 介导 NO 和 TNF- α 的上 调而改善免疫功能 Increase of the relative thymus and spleen weights as well as the serum levels of TNF- α and IL-2, and increase of NO and TNF- α secretion through the TLR4 signaling pathway and improvement of immune function	>15.75 [5]
GFPW	子实体 Fruiting bodies	15.7	不含蛋白 No protein	半乳糖:岩藻糖: 甘露糖=1.00: 0.44:0.41	O-2 位连接支链的 α -(1,6)-半乳糖 O-2- α -(1,6) galactose	体外/人脑微血管内皮 细胞 <i>In vitro</i> /Human	显著降低人脑微血管内皮 细胞的增殖(迁移和成管) 且呈剂量和时间依赖性	* [48]

(待续)

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LELFD 组分		菌丝体	Mycelia	不详	不详	不详	β -1,3-葡聚糖主链 β -1,3-glucan backbone	雌性 ICR/Slc 小鼠/ 180 肉瘤腹水细胞肿瘤 Female ICR/Slc mice/Sarcoma 180 tumor ascites cells	明显提高(肉瘤 180/小鼠) * 抗肿瘤活性, 且对巨噬细 胞具有补体 C3 活化作用 Increase of the antitumor activity of (sarcoma 180 mouse) and activation of macrophages with complement C3	[52]
GFPS1b		菌丝体	Mycelia	21	糖:蛋白质:糖醛 酸=81.32:16.60: 4.25	不详	葡萄糖:半乳糖: 阿拉伯糖=4.00: 2.00:1.00	体外/乳腺肿瘤 MCF-7 细胞 <i>In vitro</i> /mammary tumor MCF-7 cells	对乳腺癌 MCF-7 细胞 * 的增殖具有明显的抑制 作用 Inhibition of the proliferation of mammary tumor MCF-7 cells	[53]
GRN		菌丝体	Mycelia	500	Carbohydrates: proteins:uronic acid=81.32: 16.60:4.25	不详	Glc:Gal:Ara= 4.00:2.00:1.00	O-6 with glycosyl residues composed of α -L-arabinose-1,4- α - D-glucose (1 \rightarrow linked residues β -1,6 分支的 葡萄糖 Glc	体外/RAW264.7 细胞巨噬细胞 <i>In vitro</i> /RAW264.7 macrophages cells	[54]
GFP2		菌丝体	Mycelia	2.6	糖含量=98.4% Carbohydrates content=98.4%	不详	葡萄糖 Glc	ICR 小鼠/S180, Heps, EACICR mice/细胞 S180、Heps、EAC cells	激活巨噬细胞产生 IL-6、* IL-12 和 TNF- α , 促进细 胞因子的产生 Activation of macrophages to produce IL-6, IL-12 and TNF- α , and promotion of the production of cytokines	17.00-26.56 [55]
GFG-3a		菌丝体	Mycelia	88.01	糖含量=6.20% Carbohydrates content=6.20%	不详	甘露糖:葡萄糖: 果糖:阿拉伯糖= 4.510:2.460:1.331	体外/小鼠肉瘤细胞 S-180 和人肝癌细胞 <i>In vitro</i> /Bel-7402mouse sarcoma cell line S-180 and human hepatoma cell line Bel-7402	不详 Uncertain	[56]

注: *: 文献中未标明具体数值, 但多糖剂量组的肿瘤抑制结果较肿瘤模型组有显著性差异; TNF- α : 肿瘤坏死因子; IL: 白细胞介素; NK 细胞: 自然杀伤细胞; CTX: 环磷酰胺

Note: *: Specific values are not indicated, but the tumor inhibition results of polysaccharide dose group were significantly different from that of tumor model group; TNF- α : Tumor necrosis factor- α ; IL: Interleukin; NK cell: Natural killer cell; CTX: Cyclophosphamide.

于多糖的抗肿瘤活性很重要。Li 等^[58]对具有抗肿瘤活性的 GFP 多糖进行热处理降低其支化度后, 发现其体外抗肿瘤作用也随之降低。

2.3 灰树花抗肿瘤活性多糖的作用机制

2.3.1 对肿瘤细胞的直接抑制作用

灰树花多糖具有显著的抗肿瘤功效, 可直接抑制肿瘤细胞。灰树花多糖 GFP 对细胞体外增殖抑制实验[3-(4,5)-dimethylthiazoliummide, MTT]表明, 其可以抑制人类肝癌细胞 HepG2 的生长, 而且有明显的浓度依赖关系^[59]。灰树花菌丝体多糖 GFPS1b 对乳腺肿瘤 MCF-7 细胞的增殖具有明显的抑制作用^[53]。

2.3.2 诱导细胞凋亡

细胞凋亡调控发生异常是导致恶性肿瘤发生的关键原因之一。Konno 等^[60]通过观察体外培养的人类前列腺癌 PC-3 细胞加入灰树花 D 组分的实验, 发现 24 h 内有超过 95% 的细胞死亡, 推断很大可能是发生了细胞凋亡。

2.3.3 增强免疫功能

灰树花多糖的抗肿瘤效果显著, 其主要是通过增强机体免疫力, 包括提高对肿瘤细胞的杀伤能力。通过雌性 ICR 小鼠实验, Mao 等^[5]发现提取得到的灰树花子实体多糖 GP11 (分子量为 6.9–33.0 kDa) 能通过 TLR4 介导 NO 和 TNF- α (tumor necrosis factor, TNF) 上调, 改善免疫应答而间接参与抗肿瘤活性。此外, Matsui 等^[61]认为, D 组分除了激活细胞免疫能力细胞外还具有抗肿瘤作用。王莉蕊^[62]分离提取得到的 GFP-A 可以显著增强 RAW 264.7 细胞的吞噬活性, 并且在一定浓度范围内提高其 NO 的释放量, 上调细胞内 TNF- α 、IL-1 β 、IL-6、IL-12、IFN- γ 细胞因子及细胞中诱导型一氧化氮合酶的 mRNA 水平的表达, 实现免疫调节作用。本团队前期研究也发现灵芝孢子粉多糖能够增强

免疫功能, 显著降低脾脏及肿瘤微环境中某些淋巴细胞的表达, 能够恢复肿瘤微环境中肿瘤浸润淋巴细胞的特异性识别和杀伤功能^[63–64]。

2.3.4 抗肿瘤转移

肿瘤转移是指恶性肿瘤细胞离开原发肿瘤, 通过各种途径到达远处组织或器官并继续增殖生长而形成转移灶的过程。因此, 治愈恶性肿瘤的主要方法有抗肿瘤转移。Masuda 等^[65]对灰树花 MD 组分进行研究发现, 腹腔给药的时间会影响抗癌细胞转移的效果, 表明 MD 组分能够抑制肿瘤细胞转移。

3 灰树花免疫调节活性多糖及其一级结构表征

研究认为, 灰树花多糖免疫调节活性是其重要的生物活性, 能够增强免疫细胞的活性。灰树花子实体多糖 GFPBW1 的分子量为 300 kDa, 结构为 β -1,6-分支的 β -1,3-葡聚糖, 其能通过 Dectin-1/Syk/NF- κ B 信号通路激活巨噬细胞^[66]。灰树花子实体多糖 GFP-A 的分子量为 848 kDa, 其具有 TLR4 和丝裂原活化蛋白激酶介导的免疫活性^[67]。灰树花子实体多糖 GFPBW 2 的分子量为 26.2 kDa, 在 O-6 位有 β -1,3 分支的 β -1,3 和 β -1,4 葡萄糖, 雌性 ICR 小鼠实验表明, 其通过触发细胞因子分泌而激活巨噬细胞^[68]。Meng 等^[69]提取得到的灰树花子实体多糖 GFP, 其分子量为 155 kDa, 结构为 1,3 和 1,3,4-葡聚糖主链, 在体外 RAW 264.7 巨噬细胞实验中能够提高细胞增殖, 促进细胞因子或趋化因子的产生, 从而发挥免疫刺激活性。

4 灰树花其他生物活性多糖及其一级结构表征

灰树花多糖除了具有降血糖、抗肿瘤、免

疫调节活性等功能外, 还有抗病毒、抗氧化、抗辐射等生物活性。在多糖的单糖组成成分中, 葡萄糖仍然是最主要的成分; 多糖的结构分析表明, 最主要的糖苷键是 α -1,4、 β -1,3 和 β -1,6 连接键型(表 3)。

5 展望

综上所述, 灰树花多糖的生理功能活性可能与 β -1,6-葡聚糖主链化学结构相关(图 1), 但灰树花活性多糖构效尚未完全明晰, 首要原因是灰树花活性多糖的低得率。比如, 已被开发成食品补充剂的灰树花 D 组分和 X 组分, 其得率分别为 0.30%和 0.05%^[72-73]。活性多糖的低得率限制了活性研究和结构分析对原料的需要。食用菌子实体细胞壁组织含有丰富的几丁质结构, 致使多糖难以被高效提取^[3], 因此亟须采用新的辅助提取技术, 如微波、超声波、生物酶、超高压、亚临界和脉冲电场等创新技术应

用于多糖辅助提取, 实现多糖的高效制备^[74]。

大部分食用菌多糖以糖苷键相互交联, 作为细胞壁的主要成分, 也通过糖肽键与蛋白质结合^[75], 越来越多的研究表明, 蛋白质对多糖生理功能活性起着至关重要的作用, 当 β 葡聚糖与蛋白质以复合物形态存在时, 其生物效率得到提升^[76], 然而这些结合蛋白的存在更增加了多糖精细结构解析的困难。因此, 理清结构蛋白和活性多糖的相互作用, 对于进一步解开灰树花活性多糖的构效关系至关重要。

灰树花多糖经化学修饰如磷酸化^[77]、羧甲基化和硒化^[78]等能增加多糖的生物活性。同时, 灰树花多糖主要活性成分 β -1,3/1,6-葡聚糖呈高度分支并呈三股螺旋结构, 因此未来的研究需要进一步确定灰树花多糖的三维结构和功能之间的关系^[2]。灰树花功能多糖构效关系的探究将有助于设计出更多潜在的促进健康的药物和基于化学修饰的功能性食品。

表 3 其他生物活性灰树花多糖的分子质量、结构或组成

Table 3 Molecular weight, structure or composition of *G. frondosa* polysaccharides with other activities

名称	提取部位	分子量	糖:蛋白	单糖组成	结构	生理活性	参考文献
Name	Extractive fraction	Mw (kDa)	Polysaccharides: proteins	Monosaccharide/Composition	Structure/Construction	Physiological activity	References
GFP30-2-a	子实体 Fruiting bodies	2 040.0	不详 Uncertain	葡萄糖:半乳糖=1.000:0.098 Glc:Gal=1.000:0.098	α -D-1,4-葡萄糖 α -D-1,4-glucose	不详 Uncertain	[70]
Se-GFP-22	子实体 Fruiting bodies	4 130.0	糖含量=97% Carbohydrates content=97%	甘露糖:葡萄糖:半乳糖=13.30:23.30:1.00 Man:Glc:Gal=13.30:23.30:1.00	β -1,3,6-D-甘露糖、 α -1,4,6-D-半乳糖分支的 α -1,4-D-葡聚糖 α -1,4-D-glucan main chain with β -1,3,6-D-mannose and α -1,4,6-D-galactose branched	抗氧化 Antioxidative	[71]
GFP1	菌丝体 Mycelia	40.5	不详 Uncertain	葡萄糖:岩藻糖=2.3:0.5 Glc:Fuc=2.3:0.5	α -1,3-D-岩藻糖分支 β -1,6-D-葡聚糖 β -1,6-D-glucan main chain with α -1,3-D-fucose branched	抗病毒 Antiviral	[8]

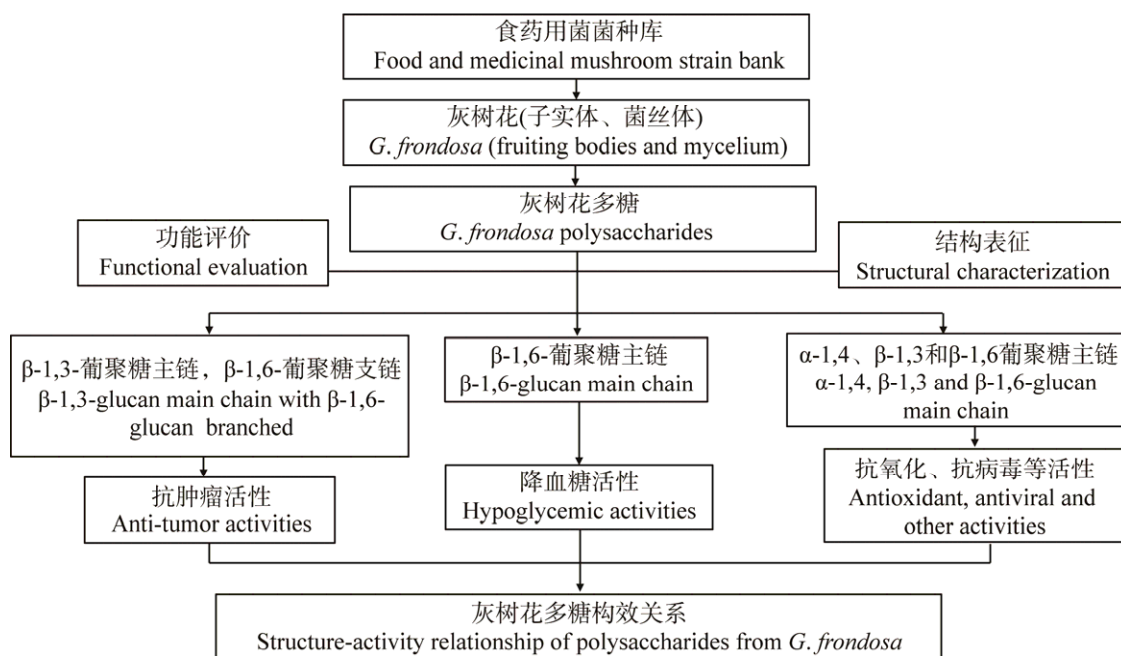


图1 灰树花多糖构效关系研究

Figure 1 Study on structure-activity relationship of *G. frondosa* polysaccharides.

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