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Optimising in vitro culture conditions for the truffle Tuber brumale

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ABSTRACT:

The vegetative propagation of ectomycorrhizal truffle fungi is limited by their slow mycelial growth. Many factors including media, isolate genotypes and environmental conditions can alter fungal mycelial growth rates. This study aimed to improve the *in vitro* growth rate of *Tuber brumale* by determining the optimal carbohydrate and nitrogen sources, temperature and pH. After 8 weeks, the highest level of growth and densest hyphal branching were recorded in the medium containing glucose as the main carbohydrate. For nitrogen, glutamine (200 mg N l⁻¹) provided the greatest hyphal growth and density compared to the other amino acid treatments. Regarding temperature, 16°C proved to be optimal for *T. brumale* growth and branching. Media of pH 6 and pH 7 were most favourable for the growth of *T. brumale*. The results from this research provide baseline data on the vegetative nutrition of *T. brumale* and have implications for the *in vitro* culture of winter truffle hyphae.

Keywords:

growth medium, hyphal branching, hyphal extension, mycelium, winter truffle

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INTRODUCTION

Hypogeous fungi belonging to the genus Tuber establish ectomycorrhizal symbiosis with diverse host plants. Species of Tuber are among the most valuable and esteemed fungal fruiting bodies (DONNINI et al. 2014; HILSZCZAŃSKA et al. 2016). The pungent and distinctive aroma released by truffles, especially when mature, attracts animals and entices them to consume the truffles. Unbeknownst to these animals, they act as vectors of truffle spore dispersal, thus playing a crucial role in the life cycle of these fungi (PACIONI et al. 2014; QIN & FENG 2022). Truffle cultivation is considered an arduous, expensive and long-term venture. However, the establishment of commercial farms is rapidly expanding as the truffle harvests from such farms have overtaken that of natural harvests (STOBBE et al. 2013; MOLINIER et al. 2016; MOSER et al. 2017).

The first mycorrhizal symbiosis under controlled conditions between Tuber and plants used spore inoculum and was carried out in Italy in the late 1960s; successes were followed by the commercial development of spore-inoculated seedlings by AGRITRUFFE in 1973 (MURAT 2015). IOTTI et al. (2016) demonstrated success in producing fruiting bodies of T. borchii via seedlings inoculated with pure mycelium. However, most commercially available truffle-inoculated (Tuber) seedlings are still established by inoculating seedlings with a suspension of truffle spores (NAKANO et al. 2020). Pure culture mycorrhizal synthesis of plants with *Tuber* may offer certain advantages over spore inoculations. First, this method can allow for the continuous production of ectomycorrhizal seedlings of consistent quality. Second, pure mycelium inoculation allows for a higher speed of seedling colonization compared to seedlings inoculated with ascospores (IOTTI et al. 2012). Finally, inoculation





Fig. 1. The morphological properties of the ascocarp, ascospores and hyphae of *Tuber brumale*: a) The ascocarp of *T. brumale*; b) The ascus with ascospores of *T. brumale* (1000X); c) The primary hyphal growth of *T. brumale* in the PDA medium.

with mycelium opens up the possibility for selecting improved strains in terms of sexual compatibility, efficiency of mycorrhizal infectivity, and adaptability to local soil and climate conditions (ZAMBONELLI *et al.* 2015; LEONARDI *et al.* 2017).

Tuber brumale is a sought after and commercial European truffle species (STROJNIK et al. 2020), which was recently reported from natural habitats in the Mazandaran Province of Iran (PULIGA et al. 2021). Previous studies showed that the growth of *Tuber* species in controlled conditions is very slow (IOTTI et al. 2002; PACIONI et al. 2007). Based on data released by IOTTI et al. (2002), the mean growth rate of various Tuber species in in vitro culture after 40 days ranged from 616 to 896 µm. In fact, T. borchii and T. maculatum showed the highest hyphal growth (896 and 842 µm, respectively), while T. brumale was the slowest growing truffle species of all of the species tested in the study (616 μ m). Given the commercial value of T. brumale, improving its mycelial growth in vivo has real-world applications, and strategies to improve its growth rate may be translatable to other truffle species. For these reasons, research was carried out to optimise the growth of T. brumale in pure culture by testing the carbohydrate, amino acid, pH and temperature preferences of this truffle species.

MATERIAL AND METHODS

Winter truffles (*Tuber brumale*) were harvested from the Yanesar region of Iran located at 36° 37′ 11.6″ latitude and 53° 43′ 8″ longitude, Behshahr, Iran in the autumn of 2019 (Fig. 1a, b). After collecting truffles from natural areas, all of the samples were transferred to the Sari Agricultural Sciences and Natural Resources University. Intact ascocarps were selected and washed under tap water and cleaned with a soft bristle brush. Their peridia were then disinfected with ethanol 70% (V/V). Tissue isolation was carried out with the aid of a stereomicroscope to choose ascocarps with intact peridia to minimise the growth of unwanted fungi including soil born and other saprotrophic fungi. For *in vitro* culturing, the truffle ascocarps were cut with a sterile razor and small fragments of less than 5 mm were separated from the inner part of the ascocarp tissue. These fragments were soaked in streptomycin (100 μ g/ml) for 7-10 min, then the sterile samples were dried on filter paper and transferred to Petri dishes containing potato dextrose agar (PDA) medium (Potato infusion: 250 g.l-1, Dextrose: 20 g.l-1, Agar: 20 g.l-1). A total of 19 truffle strains were isolated in this manner. Subcultures of the primary hyphae which emerged from the plated tissues were then transferred to a secondary plate after 20-30-days of growth and were used to carry four independent experiments. The size of the transferred plugs was 3-5 mm (Fig. 1c). To evaluate hyphal growth, the length of a single hypha was measured weekly with a graduated binocular. In order to assess the branching density, the average distances between the branches and the G index were also calculated. Five isolates were measured for each of the described assays below with 10 replicate plates per assay.

Calculating the G index. G is an indicator for evaluating the hyphal branching density and it shows the average length of the hyphae subtending from a primary hypha, according to the following equation:

$$G = \frac{Lt}{Nt}$$

Lt: total hyphal length; Nt: the total number of tips

The morphological and molecular identification of winter truffle ascocarps and pure hyphae. The morphological identification of the truffles was carried out by evaluating the width, length and shape of the asci and ascospores (Fig. 1b). These data and observations were compared with the descriptions provided by DIMITROVA & GYOSHEVA (2008) to assess the morphological characters and species designations. For molecular characterisation, DNA was isolated from the selected ascocarps and the mycelia were grown for 30 days on PDA following the protocols described by PACIONI *et al.* (2007). The internal transcribed spacer (*ITS*) nuclear rDNA was PCR amplified with the primer pair ITS5/ITS4 (WHITE *et al.* 1990) and was used for molecular identification. The generated sequences have been deposited in GenBank (Table 1). The

Isolate name	Sequences length (bp)	Accession number	Blast match	
			Species	Accession number
TB12	869	OL669325	Tuber brumale	MW829420
TB13	868	OL669326	Tuber brumale	MT495426
TB14	863	OL669327	Tuber brumale	MT495426
TB15	869	OL669322	Tuber brumale	MT495426
TB16	862	OL672491	Tuber brumale	MT495426
TB12ph	863	OL669331	Tuber brumale	MW829420
TB13ph	857	OL669329	Tuber brumale	MT495426
TB14ph	860	OL669330	Tuber brumale	MT495426
TB15ph	868	OL669323	Tuber brumale	MT495426
TB16ph	860	OL672490	Tuber brumale	MT495426
	Isolate name TB12 TB13 TB14 TB15 TB16 TB12ph TB13ph TB13ph TB14ph TB15ph TB16ph	Isolate name Sequences length (bp) TB12 869 TB13 868 TB14 863 TB15 869 TB16 862 TB13ph 857 TB14ph 860 TB15ph 868 TB15ph 868	Isolate name Sequences length (bp) Accession number TB12 869 OL669325 TB13 868 OL669326 TB14 863 OL669327 TB15 869 OL669322 TB16 862 OL672491 TB12ph 863 OL669331 TB13ph 857 OL669329 TB14ph 860 OL669323 TB15ph 868 OL669323	$\begin{array}{llllllllllllllllllllllllllllllllllll$

Table 1. Identification of the truffle ascocarps and pure culture isolates according to molecular methods.

Table 2. The media ingredients used for investigating the carbohydrate effect on the hyphal growth of *Tuber brumale*

Medium	Carbohydrates	Compounds
name	sources	
PA	-	Potato infusion + Agar
PDA	Dextrose (Glucose)	Glucose + Potato infusion + Agar
PFA	Fructose	Fructose + Potato infusion + Agar
PMA	Maltose	Maltose + Potato infusion + Agar
PSA	Sucrose	Sucrose + Potato infusion + Agar
PDexA	Dextrin	Dextrin + Potato infusion + Agar
PMtlA	Mannitol	Mannitol + Potato infusion + Agar

phylogenetic relationships were further evaluated based on the maximum likelihood method using MEGA7.

Carbohydrate assays. To determine the carbohydrate preferences of the truffle isolates, vegetative growth on a potato extract (250 g.l⁻¹) reference medium was compared to that of six media containing different carbohydrates (glucose, fructose, maltose, sucrose, dextrin and mannitol) at a concentration of (20 g.l⁻¹), in addition to a potato infusion (250 g.l⁻¹) reference medium (Table 2). A twenty-day old primary hypha (mother culture) was used to establish this experiment. Colony growth was measured weekly over an 8-week period.

Amino acids assays. A PDA medium was used as the basal medium. In this experiment, amino acids glutamine, L-asparagine, methionine, and phenyl alanine as well as a mixture of all these amino acids were compared. Assays were carried out at two concentrations, 100 and 200 mg of nitrogen per litre. The amino acids solutions were sterilised with a 0.22-micron filter and then added to the PDA medium which was autoclaved at 120°C for 20 min and cooled. A twenty-seven-day old primary hypha (mother culture) was used for this experiment. Colony growth was measured weekly over an 8-week period. **pH assays.** To determine the pH preference of *T. brumale*, PDA was chosen as the basal medium and was prepared at different pH levels. To stabilise the pH, two buffers comprising N-Tris(hydroxymethyl) methyl-3-amino-propanesulfonic acid and 2-morpholinoethanesulfonic acid were used. The media were pH adjusted with 1M NaOH and KOH to target pH levels of 5, 6, 7 and 8 following the protocols of NAKANO *et al.* (2020). Colony growth was measured weekly over an 8-week period.

Temperature assay. To determine the optimum temperature for the growth of *T. brumale* mycelium, a 27-day old hypha of *T. brumale* was placed on a PDA medium and the plates were incubated at three different temperatures including 8°C, 16°C and 24°C. Colony growth was measured weekly over an 8-week period.

Statistical analysis. Analysis of variance was determined using SAS ver. 9.1. Duncan's tests were used for mean comparisons.

RESULTS

The results from the morphological (Fig. 2a) and phylogenetic analyses demonstrate that the winter truffles collected for this study belong to *T. brumale* haplogroup II of clade A, based on MERÉNYI *et al.* (2014) (Fig. 2b). We further tested the carbohydrate, amino acid, pH and temperature preferences of the obtained isolates in order improve the rate of *in vitro* hyphal growth.

The effect of carbohydrate sources on *T. brumale* hyphal growth. After 8 weeks, the greatest hyphal length (1.1 mm) was observed in the potato dextrose agar (PDA) medium, which contained dextrose (glucose) as the main carbohydrate (Fig. 3a). All the other carbohydrate sources tested significantly improved the hyphal growth of *T. brumale* compared to the potato agar (PA) medium with the exception of mannitol, which inhibit-



Fig. 2. The distribution and phylogenetic relationship of *Tuber brumale* populations: a) The distribution and habitat of *T. brumale* according to the report made by MERÉNYI *et al.* (2014) in addition to our sampling location in Iran (blue dot); b) The phylogenetic relationship between Iranian *T. brumale* (blue asterisk) and populations in Eurasia.

ed hyphal growth (Fig. 3a). In fact, PDA supported the greatest growth of *T. brumale* across all weeks, except for during the eighth week (Supplementary Fig. S1). A comparison of eight weeks of experiments was conducted to shed light on how the mycelium lifespan can affect its growth rate. According to Fig. 4a and Supplementary Fig. S1, the highest and lowest mycelium growth rate was recorded in the first and eighth weeks after *in vitro* culture respectively.

Observations of the hyphal branching showed that the branching of hyphae only occurred in PDA, PFA, PSA and PMA media, as lateral branches (Fig. 5a; Supplementary Fig. S5a). In addition to hyphal length extension, glucose also supported the densest hyphal branching growth. The lowest average distances between the hyphal branches (0.373 mm) were measured in the PDA medium (Supplementary Fig. S5a). The lowest score in the hyphal branching *G* index (0.385) was related to the PSA medium. No



Fig. 3. The hyphal growth of *Tuber brumale* after 8 weeks: a) The hyphal growth of *T. brumale* as affected by different carbon sources after 8 weeks. The specific carbon source of different media: (PA: non carbohydrates), (PDA: Dextrose (Glucose)), (PFA: Fructose), (PMA: Maltose), (PSA: Sucrose), (PDexA: Dextrin), (PMtlA: Mannitol); b) The hyphal growth of *T. brumale* as affected by different amino acid treatments after 8 weeks (Basal medium: PDA). Abbreviations: Gln: glutamine, Asn: Asparagine, Met: Methionine, Phe: Phenylalanine. Suffixes1 and 2 are related to the concentration of amino acids (100 and 200 mg N l^{-1} , respectively); c) The hyphal growth of *T. brumale* as affected by different temperatures after 8 weeks (Basal medium: PDA), a), b, c and d - statistically significant at 1% level. The error bars depict the standard deviation. In each column, the means followed by the same letter are not significantly different according to Duncan's tests.

statistically significant differences in the G index were noted between the PDA, PMA and PSA media (Fig. 5a).

The effect of amino acids on *T. brumale* hyphal growth. Fifty-eight (58) days after *in vitro* culture, the highest hyphal length (2.05 mm) was observed in the medium containing 200 mg l⁻¹ nitrogen in the form of glutamine, which was significantly greater than for all the other treatments (Fig. 3b). The application of 200 mg l⁻¹ nitrogen in the complex amino acid treatment (contained: glutamine + L asparagine + methionine + phenyl alanine) ranked second for hyphal growth (1.9 mm) (Fig. 3b). The hyphal growth rates in all the evaluated weeks, with the exception of week 6, showed the highest quantitative values or exhibited no significant differences compared to the optimum treatment (Supplementary Fig. S2). Over the course of the experiment, mycelial growth (0.541 mm) was significantly greater during the first week, after which growth dropped until the fifth week. After week 5, the rate of hyphal growth did not change significantly (Fig. 4b).

There were no statistically significant differences in hyphal branch formation by either glutamine 200 mg l^{-1} or the complex amino acid 200 mg l^{-1} (Fig. 5b; Sup-



Fig. 4. The hyphal growth rate of *Tuber brumale* in the promising media of four carried out experiments: a) The hyphal growth rate of *T. brumale* in the PDA medium during 8 weeks (PDA: the medium which contained dextrose as the main source of carbohydrates); b) The hyphal growth rate of *T. brumale* in glutamine 200 mg N l⁻¹ treated medium during 8 weeks; c) The hyphal growth rate of *T. brumale* as affected by 16°C during 8 weeks; d) The hyphal growth rate of *T. brumale* as affected by pH 6 and 7 during 8 weeks. a, b, c and d - statistically significant at 1% level. The error bars depict the standard deviation. In each column, the means followed by the same letter are not significantly different according to Duncan's tests.

plementary Fig. S5b). The average distances between the new hyphal apexes of glutamine 200 mg l^{-1} and complex amino acid 200 mg l^{-1} were 0.275 and 0.285 mm, respectively (Supplementary Fig. S5b). Figure 5b shows that the lowest values for *G* were recorded in the media containing glutamine 200 mg l^{-1} (0.287) and complex amino acid 200 mg l^{-1} (0.285), which show denser hyphal branching than for the other treatments.

The effect of temperature on *T. brumale* hyphal growth. The highest growth of *T. brumale* hypha (1.03)

mm) after 8 weeks was recorded at 16° C (Fig. 3c). The optimum temperature for hyphal growth in the 1st, 3rd and 8th weeks was 16°C, however, in the other weeks of the experiment no significant differences were observed among the tested temperatures (Supplementary Fig. S3). At 16°C, the highest hyphal growth rate was measured as 0.316 mm in the first week, which was greater than all the other weeks and temperatures (Fig. 4c). The densest hyphal branching was observed at 16°C after 8 weeks (Fig. 5c; Supplementary Fig. S5c). Although there were no statistically significant differences in the *G* index of



Fig. 5. The *G* index of the hyphal branching of *Tuber brumale*: a) The *G* index of the hyphal branching of *Tuber brumale* in certain media with different carbon sources which lead to hyphal branching; after 8 weeks. The specific carbon source of different media: (PDA: Dextrose (Glucose)), (PFA: Fructose), (PMA: Maltose), (PSA: Sucrose); b) The *G* index of the hyphal branching of *T. brumale* as affected by different amino acid treatments after 8 weeks (Basal medium: PDA). Abbreviations: Gln: glutamine, Asn: Asparagine, Met: Methionine, Phe: Phenylalanine. Suffixes1 and 2 are related to the concentration of amino acids (100 and 200 mg N l⁻¹, respectively); c) The *G* index of the hyphal branching of *T. brumale* as affected by different temperatures after 8 weeks (Basal medium: PDA); d) The *G* index of the hyphal branching of *T. brumale* as affected by different pHs after 8 weeks (Basal medium: PDA). a, b, c and d - statistically significant at 1% level. The error bars depict the standard deviation. In each column, the means followed by the same letter are not significantly different according to Duncan's tests.

16°C (0.523) and 8°C (0.599) treatments, the average distance between the new hyphal apexes was 0.409 mm at 16°C, which was significantly lower than at the other temperatures (Fig. 5c; Supplementary Fig. S5c).

The effect of pH on *T. brumale* **hyphal growth.** The growth of *T. brumale* hypha was the greatest at pH 6 and 7, leading to hyphal lengths of 1.025 and 0.975 mm, respectively, after 8 weeks with no significant differ-

ences between the two pHs (Fig. 3d). The highest hyphal growth rate took place during the first week and the lowest growth rate was recorded in the seventh and eighth weeks (Fig. 4d; Supplementary Fig. S4). The observations of hyphal branching showed that the densest branches formed at pH 6 (Fig 5d; Supplementary Fig. S5d). At the densest hyphal branching junctions, the distance between the branches was 0.400 mm at pH 6. Figure 5d shows that there were no statistically signifi-

cant differences between the *G* index of pH 6 (0.555) and pH 5 (0.541). However, hyphal extension at pH 7 was significantly greater than at pH 5, while acidic conditions increased hyphal branching and branches did not form at pH 8 (Fig. 5d).

DISCUSSION

In this study we tested the carbohydrate, amino acid, pH and temperature preferences of *T. brumale* in order to improve the hyphal growth rate of this commercial truffle species *in vitro*. We found that *T. brumale* isolates have a nutritional preference for glucose as a carbohydrate source and glutamine as a nitrogen source. Mycelial growth was also improved by culturing on media at a pH of 6 and 7, and incubating at 16°C.

Optimising mycelial growth requires an understanding of fungal nutrition and physiology. The comparison of our results with previous studies indicates that truffle species may differ in their nutritional preferences. For example, SALTARELLI et al. (1998) reported that glucose and fructose are a superior source of carbon for T. borchii strain ATCC 96540. Yet, CECCAROLI et al. (2001) concluded that mannose was preferred as a carbohydrate for three strains of *T. borchii* (i.e. 1BO, 17BO and 10RA) in comparison with glucose and mannitol. Interestingly, the hyphal growth of the 17BO strain was curbed in the medium containing mannitol. ORIAIFO (2014) evaluated diverse carbohydrate sources on the growth of T. borchii and found that the highest growth rate in all the studied strains was observed with starch as the carbohydrate source in comparison with sucrose, fructose, xylose, glucose, mannose, glycerol sorbitol and mannitol. In contrast, research on T. maculatum showed that mycelial growth was highest when lactose was used as a carbon source, but the difference between it and other carbohydrate sources including sucrose, fructose, xylose, glucose, dextrose, mannose and maltose was not statistically significant (NADIM et al. 2016).

It is well documented that T. melanosporum and T. brumale have a close phylogenetic relationship (BONITO et al. 2013). Thus, our results for T. brumale may be generalisable to T. melanosporum, and vice versa. Although compounds such as cellulose, cellobiose and starch can be used as a carbon source for T. melanosporum (MAMOUN & OLIVIER 1991), genomic sequencing showed a limited gene repertoire in this species' coding for carbohydrate active enzymes (CAZymes), which are the enzymes responsible for biodegrading non-living organic matter. For example, just a few glycoside hydroxylase (GH) genes were identified in T. melanosporum in contrast to its saprotrophic relatives which contain many genes in the GH6 and GH7 cellulase gene families (MARTIN et al. 2010). These findings provide the genetic basis for the weak saprotrophic ability of *T. melanosporum* observed. However, T. melanosporum has an invertase enzyme,

enabling it to hydrolyse the sucrose provided by the host plant. Based on these data, it is expected that *T. brumale* grows in a similar way to *T. melanosporum* and prefers to assimilate simple sugars rather than polymer carbohydrates.

The proliferation and growth of mycelium is dependent on hyphal branching patterns. G is an efficient index for the assessment of branching conditions. The value of G depended on the total length of the mycelium and the number of tips. Thus, G is an indicator of growth density and when branching is rare, the G index will be higher (MOORE *et al.* 2020). We found that G was highest with fructose as a carbon source and asparagine as a nitrogen source. The G values were also greater at higher temperatures and the pH levels tested. Although many aspects of hyphal branching are not yet known, one interesting hypothesis is that heat shock proteins (polypeptides) are involved in the mechanisms of branching initiation. It is thought that conformational change in the cell wall proteins is required for branch initiation, and that heat shock proteins play a key role in the delivery of initiation polypeptides to the appropriate site (MOORE et al. 2020). The Spitzenkorper (SPK) is a multicomponent pleomorphic structure which plays a key role in maintaining hyphal extension (RIQUELME & SANCHEZ-LEON 2014). Lateral hyphal branching requires the emergence of a new SPK to support the growth of nascent hyphae (HARRIS 2019). Previous studies have revealed that the main components of the SPK include an accumulation of vesicles (containing the enzymes required for cell wall synthesis), ribosomes, actin microfilaments, chitosomes and an amorphous material of undefined nature (RI-QUELME & SANCHEZ-LEON 2014; HARRIS 2019). Amino acids are, in terms of actin (as a protein polymer) and enzyme structure, critical to the formation and activity of SPK (BEREPIKI et al. 2011).

Cell division is controlled by core cell cycle genes and transcription factors. Among the core cell cycle genes involved in the biosynthesis of cyclins and their catalytic partners, cyclin-dependent kinases (CDKs) play a key role in the regulation and progression of cell cycles (HYDBRING et al. 2016). Considering the structure of cyclins and CDKs (proteins), amino acids serve as their monomeric units, thus increasing the amino acid content of the cells would be effective for the progression of cell cycles. Furthermore, the amino acids and amino groups involved in histones and nucleotides play an important role in cell cycle division (MARIÑO-RAMÍREZ et al. 2005). Glutamine is an energetically favoured nitrogen source for fungi, and is involved in the biosynthesis pathways of many secondary metabolites and other amino compounds (TUDZYNSKI 2014). Glutamine serves as an efficient nitrogen source for T. brumale hyphal growth. Previous studies on nitrogen metabolism in ectomycorrhizal fungi demonstrated that some amino acids including glutamine, glutamate, arginine and asparagine are abundant in the fungal mycelium (JOHANSEN *et al.* 1996; BAGO *et al.* 1999). Ectomycorrhizal fungi are known to have the ability to uptake some forms of organic nitrogen, which they can transfer to the host plant in the form of glycine and glutamate (HAWKINS *et al.* 2000). Thus, it is interesting that *T. brumale* utilised and preferred glutamine over the other amino acids, raising the question as to whether *T. brumale* utilises and transfers organic N to host plants in natural systems.

The evaluation of the hyphal growth of *T. sinense* in media with different nitrogen sources showed that the highest growth occurred in yeast extract and peptone treatments, which were preferred as a nitrogen source compared to mineral forms of nitrogen such as ammonium nitrate and potassium nitrate (LIU et al. 2008). Alanine was also found to stimulate more hyphal growth of T. melanosporum in comparison with asparagine and inorganic sources (KAMAL 2011). The formation of D-type cyclins is critical for progression through the G1 phase of the cell cycle to drive DNA synthesis (S phase). The biosynthesis pattern of D-type cyclins changes in response to the intracellular conditions and environmental factors (Hydbring et al. 2016). In another study by ORIAIFO (2014), the effect of nitrogen sources was investigated on T. borchii. The results revealed that the effect of the nitrogen source was genotype-dependent, and the response of each strain to the nitrogen source was different.

According to ZAMBONELLI *et al.* (2016), the habitats of *T. brumale* have a 3.7-15.8°C annual mean temperature. We found that 16°C was the optimal incubation temperature for *T. brumale* growth among all the isolates tested. The optimum temperature for hyphal growth in *T. maculatum* was reported to be 15°C (NADIM *et al.* 2016). In another investigation, ORIAIFO (2014) pointed out that most strains of *T. borchii* preferred 20 and 17.5°C, but they noted that temperature preference was isolate-dependent.

Tuber species are known to prefer non-acidic environments, which is one of the reasons why truffle orchard soils are often limed or planted on naturally alkaline soils. Previous studies aimed at understanding the ecological parameters of T. brumale habitats elucidated that the average water pH in the Carpathian Basin is 6.9 (ZAMBONELLI et al. 2016). We found that T. brumale mycelium grew significantly better at a pH of 6 and 7 compared to the other pH levels tested. Hyphal growth rates and preferences differ between the different clades of *Tuber*. It was reported that the hyphal growth of *T*. japonicum in in vitro culture was observed in pH 5 and 6, while T. himalayense and T. longispinosum exhibited maximum growth in pH 7 (NAKANO et al. 2020). ORIAIFO (2014) tested a range of pH values (4-9) on T. borchii mycelial growth and found pH to be a significant factor for mycelial growth, however, the optimum pH varied by isolate. In another study, the optimum pH

hyphal growth of *T. maculatum* was reported to be in the range of pH 5.8-7.6 (NADIM *et al.* 2016). These results demonstrate that the response to the pH medium is both species and genotype dependent. Thus, each species of fungi, and possibly each strain, may need to be tested to determine the optimum pH for growth.

CONCLUSION

Tuber brumale is an important commercial truffle which grows across Europe and was recently reported from Iran. Through this research we were able to improve the culture medium and environmental conditions for the growth of T. brumale in pure culture. We showed that glucose and glutamine are the preferred carbon and nitrogen sources, respectively. For the growth of T. brumale mycelium, a PDA (potato extract + Dextrose (glucose) + agar) medium containing 20 g/L glucose is an appropriate option, and adding 200 mg l⁻¹ nitrogen from a glutamine source is recommended to promote faster growth. We found that the optimal temperature for hyphal growth is around 16°C and the optimal pH medium is 6-7. Importantly, the hyphal growth rates decreased significantly over time, indicating that fresh mycelium should be propagated and sub-cultured frequently to maintain active growth.

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REFERENCES

- BAGO B, PFEFFER PE, DOUDS JR DD, BROUILLETTE J, BECARD G & SHACHAR-HILL Y. 1999. Carbon metabolism in spores of the arbuscular mycorrhizal fungus *Glomus intraradices* as revealed by nuclear magnetic resonance spectroscopy. *Plant Physiology* **121**(1): 263-272.
- BEREPIKI A, LICHIUS A & READ ND. 2011. Actin organization and dynamics in filamentous fungi. *Nature Reviews Microbiology* **9**(12): 876-887.
- BONITO G, SMITH ME, NOWAK M, HEALY RA, GUEVARA G, CA-ZARES E, KINOSHITA A, NOUHRA ER, DOMINGUEZ LS, TEDERSOO L & MURAT C. 2013. Historical biogeography and diversification of truffles in the Tuberaceae and their newly identified southern hemisphere sister lineage. *PloS One* 8(1): e52765.
- CECCAROLI P, SALTARELLI R, CESARI P, ZAMBONELLI A & STOCCHI V. 2001. Effects of different carbohydrate sources on the growth of *Tuber borchii* Vittad. mycelium strains in pure culture. *Molecular and Cellular Biochemistry* **218**(1): 65-70.
- DIMITROVA E & GYOSHEVA M. 2008. Hypogeous ascomycetes in Bulgaria. *Phytologia Balcanica* 14(3): 309-314.

- DONNINI D, BENUCCI GM, BENCIYANGA M & FALINI LB. 2014. Quality assessment of truffle-inoculated seedlings in Italy: proposing revised parameters for certification. *Forest Systems* 23(2): 385-393.
- HARRIS SD. 2019. Hyphal branching in filamentous fungi. *Developmental Biology* **451**(1): 35-39.
- HAWKINS HJ, JOHANSEN A & GEORGE E. 2000. Uptake and transport of organic and inorganic nitrogen by arbuscular mycorrhizal fungi. *Plant and Soil* **226**(2): 275–285.
- HILSZCZAŃSKA D, SIEBYŁA M, HORAK J, KRÓL M, PODSADNI P, STECKIEWICZ P, BAMBUROWICZ-KLIMKOWSKA M, SZUTOWSKI, M & TURŁO J. 2016. Comparison of chemical composition in *Tuber aestivum* Vittad. of different geographical origin. *Chemistry & Biodiversity* 13(12): 1617-1629.
- HYDBRING P, MALUMBRES M & SICINSKI P. 2016. Non-canonical functions of cell cycle cyclins and cyclin-dependent kinases. *Nature Reviews Molecular Cell Biology* **17**(5): 280–292.
- IOTTI M, AMICUCCI A, STOCCHI V & ZAMBONELLI A. 2002. Morphological and molecular characterization of mycelia of some *Tuber* species in pure culture. *New Phytologist* **155**(3): 499-505.
- IOTTI M, PIATTONI F, LEONARDI P, HALL IR & ZAMBONELLI A. 2016. First evidence for truffle production from plants inoculated with mycelial pure cultures. *Mycorrhiza* 26(7): 793-798.
- IOTTI M, PIATTONI F & ZAMBONELLI A. 2012. Techniques for host plant inoculation with truffles and other edible ectomycorrhizal mushrooms. In: ZAMBONELLI A & BONITO G (eds.), *Edible ectomycorrhizal mushrooms*, pp. 145-161, Springer, Berlin, Heidelberg.
- JOHANSEN A, FINLAY RD & OLSSON PA. 1996. Nitrogen metabolism of external hyphae of the arbuscular mycorrhizal fungus *Glomus intraradices*. *New Phytologist* **133**(4): 705-712.
- KAMAL S. 2011. Effect of nutrient sources and plant hormones on mycelial morphology of the black perigord truffle *Tuber melanosporum*. Proceedings of the 7th International conference on mushroom biology and mushroom products (ICMBMP7), pp. 509-515, Arcachon, France.
- LEONARDI P, IOTTI M, ZEPPA SD, LANCELLOTTI E, AMICUCCI A & ZAMBONELLI A. 2017. Morphological and functional changes in mycelium and mycorrhizas of *Tuber borchii* due to heat stress. *Fungal Ecology* **29**: 20–29.
- LIU RS, LI DS, LI HM & TANG YJ. 2008. Response surface modeling the significance of nitrogen source on the cell growth and *Tuber* polysaccharides production by submerged cultivation of Chinese truffle *Tuber sinense*. *Process Biochemistry* **43**(8): 868-876.
- MAMOUN M & OLIVIER JM. 1991. Influence du substrat carboné et de la forme d'azote minéral sur la croissance de *Tuber melanosporum* (Vitt) en culture pure. Application à la production de biomasse mycélienne. *Agronomie* **11**(6): 521-527.
- MARIÑO-RAMÍREZ L, KANN MG, SHOEMAKER BA & LANDSMAN D. 2005. Histone structure and nucleosome stability. *Expert Review* of Proteomics **2**(5): 719-729.
- MARTIN F, KOHLER A, MURAT C, BALESTRINI R, COUTINHO PM, JAILLON O, MONTANINI B, MORIN E, NOEL B, PERCUDANI R & PORCEL B. 2010. Périgord black truffle genome uncovers evolutionary origins and mechanisms of symbiosis. *Nature* **464**(7291): 1033-1038.
- MERÉNYI Z, VARGA T, GEML J, ORCZÁN ÁK, CHEVALIER G & BRATEK Z. 2014. Phylogeny and phylogeography of the *Tuber* brumale aggr. *Mycorrhiza* **24**(1): 101-113.
- Molinier V, Murat C, Baltensweiler A, Büntgen U, Martin F, Meier B, Moser B, Sproll L, Stobbe U, Tegel W & Egli S.

2016. Fine-scale genetic structure of natural *Tuber aestivum* sites in southern Germany. *Mycorrhiza* **26**(8): 895–907.

- MOORE D, ROBSON GD & TRINCI AP. 2020. 21st century guidebook to fungi. Cambridge University Press.
- MOSER B, BÜNTGEN U, MOLINIER V, PETER M, SPROLL L, STOBBE U, TEGEL W & EGLI S. 2017. Ecological indicators of *Tuber aestivum* habitats in temperate European beech forests. *Fungal Ecology* **29**: 59-66.
- MURAT C. 2015. Forty years of inoculating seedlings with truffle fungi: past and future perspectives. *Mycorrhiza* **25**(1): 77-81.
- NADIM M, SAIDI N, HASANI IW, EL BANNA YY, SAMIR O, ASSAD MEH & SHAMEKH S. 2016. Effects of some environmental parameters on mycelia growth of Finnish truffle *Tuber Maculatum*. *International Journal of Applied Science and Engineering* 3: 2394–3661.
- NAKANO S, KINOSHITA A, OBASE K, NAKAMURA N, FURUSAWA H, NOGUCHI K & YAMANAKA T. 2020. Influence of pH on *in vitro* mycelial growth in three Japanese truffle species: *Tuber japonicum, T. himalayense,* and *T. longispinosum. Mycoscience* **61**(2): 58-61.
- ORIAIFO OF. 2014. In vitro mycelial growth and root infection of loblolly pine seedlings by bianchetto truffle (Tuber borchii). Doctoral dissertation, North Carolina Agricultural and Technical State University.
- PACIONI G, CERRETANI L, PROCIDA & G. CICHELLI A. 2014. Composition of commercial truffle flavored oils with GC–MS analysis and discrimination with an electronic nose. *Food Chemistry* **146**: 30–35.
- PACIONI G, LEONARDI M, AIMOLA P, RAGNELLI AM, RUBINI A & PAOLOCCI F. 2007. Isolation and characterization of some mycelia inhabiting *Tuber* ascomata. *Mycological Research* **111**(12): 1450-1460.
- PULIGA, F, ILLICE M, IOTTI, M, LEONARDI P, BAGHDADI A, MOZA-FARI AA & ZAMBONELLI A. 2021. True truffle diversity in Iran. *Italian Journal of Mycology* **50**: 52–62.
- QIN J & FENG B. 2022. Life cycle and phylogeography of true truffles. *Genes* **13**(1): 145.
- RIQUELME M & SÁNCHEZ-LEÓN E. 2014. The Spitzenkörper: a choreographer of fungal growth and morphogenesis. *Current Opinion in Microbiology* **20**: 27–33.
- SALTARELLI R, CECCAROLI P, VALLORANI L, ZAMBONELLI A, CIT-TERIO B, MALATESTA M & STOCCHI V. 1998. Biochemical and morphological modifications during the growth of *Tuber borchii* mycelium. *Mycological Research* **102**(4): 403-409.
- STOBBE U, EGLI S, TEGEL W, PETER M, SPROLL L & BÜNTGEN U. 2013. Potential and limitations of Burgundy truffle cultivation. *Applied Microbiology and Biotechnology* **97**(12): 5215-5224.
- STROJNIK L, GREBENC T & OGRINC N. 2020. Species and geographic variability in truffle aromas. *Food and Chemical Toxicology* **142**: 111434.
- TUDZYNSKI B. 2014. Nitrogen regulation of fungal secondary metabolism in fungi. *Frontiers in Microbiology* **5**: 656.
- WHITE TJ, BRUNS TD, LEE S & TAYLOR J. 1990. Analysis of phylogenetic relationship by amplification and direct sequencing of ribosomal RNA genes. In: INNIS MA, GELFAND DH, SNINSKY JJ & WHITE TJ (eds.), PCR Protocols: a guide to methods and applications, pp. 315–322, Academic Press, San Diego.
- ZAMBONELLI A, IOTTI M & HALL I. 2015. Current status of truffle cultivation: recent results and future perspectives. *Italian Journal of Mycology* **44**: 31-40.
- ZAMBONELLI A, IOTTI M & MURAT C. 2016. *True truffle (Tuber spp.) in the world*. Springer International Publishing, Cham.

Botanica

REZIME -

Optimizacija uslova za in vitro gajenje tartufa Tuber brumale

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Vegetativno razmnožavanje ektomikoriznih gljiva tartufa ograničeno je njihovim sporim rastom micelija. Mnogi faktori, uključujući medijume, izolovane genotipove i uslove životne sredine, mogu promeniti stopu rasta micelija gljiva. Ova studija je imala za cilj da poboljša *in vitro* brzinu rasta *Tuber brumale* određivanjem optimalnih izvora ugljenih hidrata i azota, temperature i pH. Posle 8 nedelja, najveći rast i najgušće grananje hifa zabeleženi su u medijumu koji je sadržao glukozu kao glavni ugljeni hidrat. Za azot, glutamin (200 mg N l⁻¹) je obezbedio najveći rast i gustinu hifa u poređenju sa drugim tretmanima amino kiselina. Što se tiče temperature, 16°C je bilo optimalno za rast i grananje *T. brumale*. Rezultati ovog istraživanja daju osnovne podatke o vegetativnoj ishrani *T. brumale* i imaju primenu u kulturi hifa zimskih tartufa *in vitro*.

Ključne reči: medijum za gajenje, grananje hifa, izduživanje hifa, micelijum, zimski tartuf