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Natural decomposition of hornbeam wood decayed by the white rot fungus *Trametes versicolor*

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ABSTRACT

The impacts of white-rot fungi on altering wood chemistry have been studied mostly *in vitro*. However, *in vivo* approaches may enable better assessment of the nature of interactions between saprotrophic fungi and host tree in nature. Hence, decayed and sound wood samples were collected from a naturally infected tree (*Carpinus betulus* L.). Fruiting bodies of the white rot fungus *Trametes versicolor* grown on the same tree were identified using rDNA ITS sequencing. Chemical compositions (cellulose and lignin) of both sound and infected wood were studied. FT-IR spectroscopy was used to collect spectra of decayed and un-decayed wood samples. The results of chemical compositions indicated that *T. versicolor* reduced cellulose and lignin in similar quantities. Fungal activities in decayed wood causes serious decline in pH content. The amount of alcohol-benzene soluble extractives was severely decreased, while a remarkable increase was found in 1% sodium hydroxide soluble and hot water extractive contents in the decayed wood samples, respectively. FT-IR analyses demonstrated that *T. versicolor* causes simultaneous white rot in the hornbeam tree *in vivo* which is in line with *in vitro* experiments.

Key words: Hornbeam wood, *Trametes versicolor*, chemical compositions, FT-IR analyses.

INTRODUCTION

Major decay process caused by fungi initially starts through wounds and cracks in the stem branches and roots of living trees. Later, when tree dies, other fungi gain access to the woody material and

immediately speed up the decomposition process (Stokland et al. 2012, Schwarze et al. 2004, Schmidt 2006). Several microorganisms may be involved in the degradation. The most efficient wood destroyers are the white-rot and brown-rot fungi among the Basidiomycetes (Eriksson et al. 1990, Eaton and Hale 1993, Schmidt 2006). In addition, some Ascomycetes and Deuteromycetes cause a

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different type of wood rot called soft rot. However, some white rot fungi are able to decompose wood in a similar way to soft rot decay (Schwarze et al. 2004, Bari et al. 2017).

The majority of the white-rot fungi use cellulose and hemicelluloses at nearly the same rate; however, lignin is usually utilized at a somewhat faster rate. A few white-rot fungi, on the other hand, remove lignin and hemicelluloses preferentially, but eventually degrade all wood cell wall components. These fungi invade tissues of hardwoods and they rarely have been seen on softwoods (Schmidt 2006, Kubicek 2013). White rot has been classified by macroscopic characteristics into simultaneous and selective white rot and (Schmidt 2006, Kubicek 2013). Many white-rot fungi seem to produce just one of these behaviors, but some species can cause alternative forms under different conditions (Martínez et al. 2005, Bari et al. 2015c). Many white-rot fungi occur as dormant spores within the live sapwood and immediately after falling moisture content in dead cells, the mycelium grows and the decomposition starts (Boddy 1994).

The impacts of white-rot fungi on wood chemistry have been studied by various methods (e.g. Martínez et al. 2001, Mohebbi 2005, Bari et al. 2015a, Karim et al. 2016) But the most of the above mentioned methods have been carried out *in vitro*. However, there are inherent difficulties in emulating micro-environmental conditions occurring in the field (Rayner and Webber 1984) and *in vivo* approaches may enable better assessment of the nature of interactions between saprotrophic fungi and the wood (Woodward and Boddy 2008). In this regards, the white-rot fungus *Trametes versicolor* is a unique microorganisms that has a widespread host (Eaton and Hale 1993, Schmidt 2006) and act as great carbon recycler in the natural forest ecosystems. Hornbeam (*Carpinus betulus*) is an important tree species in Iran, economically and ecologically which accounts for 30.5% of the standing volume and 30% of the stem number

in the Hyrcanian forests of Iran. This species in combination with beech compose 54 and 60% of the stem number and standing volume of the Iranian Northern Forests, respectively (Sagheb Talebi et al. 2014), and therefore, it has been studied from different aspects (Taghiyari and Sarvari Samadi 2010).

More than 90% of wood decaying fungi in standing and felled trees in the north of Iran and America are caused by white-rotting fungi (Gilbertson 1980, Bari et al. 2015a). Thus, the white rot fungi called as predator and terminator for natural forest ecosystems (Bari et al. 2016) and many studies were carried out on them (Taghiyari et al. 2015). Former studies (Bari et al. 2015a, b, c, d, 2016) demonstrated that the white-rot fungus *Trametes versicolor* produces simultaneous decay in beech wood in laboratory condition (*in vitro*) (Bari et al. 2016). In this respect, a question arises that whether the degradation patterns of hornbeam wood decayed by this fungus is the same as in beech or not? In addition, will the fungus produce simultaneous or selective decay? The naturally decomposition of wood gives us new and significant knowledge about the behavior of fungi in nature compared to controlled conditions. Moreover, investigations of decomposition processes in forest ecosystems will be useful from the natural recycling of organic substance point of views (Stokland et al. 2012). Since there is no information about the behavior of this fungus in natural conditions, this study is aimed to assess the chemical composition of wood naturally decayed by *T. versicolor*. FT-IR spectroscopic analysis was accompanied with conventional chemical analysis to study the chemical changes in hornbeam wood attacked by the white-rot fungus *T. versicolor*. Comparable studies had been already done with the same fungus and *Fagus orientalis* wood (Bari et al. 2015a).

MATERIALS AND METHODS

FUNGI IDENTIFICATION

The fruiting bodies of white rot fungus *Trametes versicolor* (L.: Fr.) Pilát were collected from a fallen hornbeam tree (*Carpinus betulus* L.) at the Guilan forest, northwest Iran. Then morphology identification was done according to Ryvarden and Gilbertson (1993, 1994) and molecular identification was carried out according to Schmidt et al. (2012) and Bari (2014) with ten replicates. In brief, the DNA of the infected wood and fruiting bodies of fungi were extracted; PCR was amplified with primers ITS1 and ITS4, electrophoresed, purified, and sequenced. Thereafter, the identification was accomplished by the sequence comparison with sequences deposited in the DNA databases using the BLAST in the NCBI. The same tree was used for further chemical analyses (see below).

CHEMICAL ANALYSIS

Un-decayed and parts of the naturally decayed wood specimen were cut, dried at 103 ± 2 °C for 24 hours, and then ground to pass a 40-mesh (420- μ m) screen. The preparation of the wood powder for chemical composition analysis was done according to TAPPI standard (T 264 om-88, 1988). Lignin and cellulose content of both kinds of wood samples were measured using T 222 om-98 (1998), and T 17 wd-97 (1997a) procedures, respectively. The chemical percent of wood extractives was evaluated in accordance with TAPPI T 212 om-93 (1993a), TAPPI T 207 om-93 (1993b), and TAPPI T 204 cm-97 (1997b) standard methods, respectively. The pH values of the samples were calculated using a pH meter device. For this purpose, about 1 grams of meshed wood powder was added to 25ml distilled water and they were kept in room condition overnight. Then, the pH meter was calibrated with distilled water and pH values of the samples were

then determined. All analyses were done with five replicates.

FT-IR SPECTROSCOPY

Dried samples were milled and passed through a mesh 40 sieve mesh. FT-IR spectra were obtained directly from wood powder. Using a Shimadzu 8400s FT-IR Spectrometer equipped with DLATGS detector, all samples were examined at a spectral resolution of 4 cm^{-1} : Spectra from 30 scans of sample and background scans were measured. The background spectra were collected using an empty collector. A rubber band method was used for baseline correction. The band for CO_2 was removed to make a suitable baseline correction (Mohebbi 2005).

STATISTICAL ANALYSIS

To compare chemical losses, a Student t-test was performed (95% confidence level) between decayed and un-decayed samples. Statistical analysis was performed using the SPSS software program, version 17 (2010).

RESULTS AND DISCUSSION

FUNGI IDENTIFICATION

The results indicated that three fungi i.e. *Trametes versicolor*, *Trichoderma harzianum* (Rifai.) and *Mortierella elongata* (Linnem.) were present in the samples, while *T. versicolor* was the only found white-rot fungus. On the other hand, *T. harzianum* and *M. elongata* as members of the Pezizomycetes and Mucoromycetes might have participated in the decomposition of wooden disc. *T. harzianum* is known as a mycoparasite and a cellulolytic fungus, while *M. elongata* is a terrestrial saprophytic fungus inhabiting soil that may use sugar material from wood or other plants (Eaton and Hale 1993).

CHEMICAL ANALYSIS

The average chemical components in un-decayed and naturally-decayed hornbeam samples by *T. versicolor* are shown in Table I. As depicted in this figure, mean cellulose loss was a bit more pronounced compared to lignin degradation. On the other hand, at the examined stage of decay, cellulose and lignin were degraded approximately at the same rate. *T. versicolor* contains the full lingocellulosic enzyme system for degrading all wood cell walls layers. Entire decomposition of wood cell walls by fungi will take place only when the fungi have all necessary enzymes for degradation. In this regard, only some wood decaying fungi such as *Phanerochaete chrysosporium*, *P. carnosa* and *T. versicolor* have the full genome for degradation (Canam et al. 2013). However, fungi which are specialized may have different behavior in nature than under controlled conditions. There are also evidences that certain cell wall components may induce the production of the necessary enzymes. Thus cellulose induces the formation of carbohydrases in white-rotting fungi (Fengel and Wegener 1989). Nonetheless, since *T. versicolor* is a member of the white-rot fungi, these behaviors were expected. Several literatures (e.g. Eriksson et al. 1990) as well as our former studies (Bari et al. 2015a, c) in controlled situation, demonstrated the simultaneous decay behavior of this fungi. The average losses of cellulose and lignin were 32.52% and 26.10%, respectively which were in accordance with former studies and our investigations on *Fagus orientalis* wood (Bari et al. 2015a). A significant decrease was seen in pH value in naturally decayed wood. According to Humar et al. (2001), the fungi are able to severely reduce the pH in wood after advancement of decay. Nonetheless, it has been known that white-rot fungi do not cause such massive decline in pH of wood comparing to brown-rot fungi (Shortle 1990). On the other hand, considerable changes were recorded in extractives

constituent values of decayed wood. Reduction of alcohol-benzene soluble extractives was observed in decayed samples. However, 1% NaOH soluble extractives as well as hot-water soluble extractives were increased in decayed wood samples. It was found that the wood decaying fungi hydrolyze the extractives such as styrene esters and waxes into fatty acid and sterol moieties as an important source of carbon for their growth (Silk et al. 2001). In this regards, similar observations (e.g. Malakani et al. 2014) were reported.

FT-IR EVALUATIONS UNDECAYED WOOD (CONTROL)

The FT-IR spectrum of un-decayed hornbeam wood is depicted in Figure 1. A strong hydrogen bonded (O–H) stretching absorption was observed at 3500-3300 cm^{-1} (1) and a prominent C–H stretching absorption around 2896 cm^{-1} (2). On top of that, there were many well-defined peaks in the fingerprint region between 1800–500 cm^{-1} . The peaks in the fingerprint were assigned in accordance with several studies (e.g. Harrington et al. 1964, Hergert 1971, Schultz and Glasser 1986, Faix 1992, Pandey and Pitman 2003, Mohebbi 2005), the assigned peaks in the above mentioned literatures were un-conjugated at 1738 cm^{-1} (3); conjugated C–O stretching at 1594 cm^{-1} (4); aromatic skeletal vibration at 1504 cm^{-1} (5); C–H deformation in lignin and carbohydrates at wave numbers 1461 (6) and 1423 cm^{-1} (7); C–H deformation in cellulose and hemicelluloses at 1372 cm^{-1} (8); C–H vibration in cellulose and C₁–O vibration in syringyl derivatives at 1325 cm^{-1} (9); syringyl ring and C–O stretching in lignin and xylan at wave number 1238 cm^{-1} (10); C–O–C vibration in cellulose and hemicellulose at 1157 cm^{-1} (11); aromatic skeletal and C–O stretch at 1096 cm^{-1} (12); C–O stretch in cellulose and hemicelluloses at wave number 1035 cm^{-1} (13) and C–H deformation in cellulose at 892 cm^{-1} (14).

TABLE I
Average of percent chemical components in decayed hornbeam wood in sound and naturally infected sample by white-rot fungus *Trametes versicolor*.

Wood Property	Exposure condition	
	Un-decayed	Naturally decayed
Lignin	31.77 ± 2.83	26.10 ± 1.66
Cellulose	42.52 ± 1.39	32.52 ± 1.81
pH	05.31 ± 0.55	03.26 ± 0.41
Alcohol-benzene soluble extractives (%)	02.71 ± 0.84	01.98 ± 1.02
Hot-water soluble extractives (%)	05.12 ± 1.36	06.48 ± 1.73
1% NaOH soluble extractives (%)	21.36 ± 2.04	32.51 ± 2.65

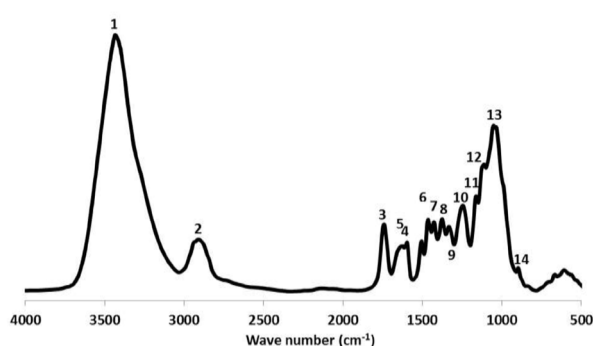


Figure 1 - FT-IR spectrum bands of un-decayed *Carpinus betulus*.

FT-IR SPECTRA OF DECAYED WOOD

The result of FT-IR spectra analyses of decayed and un-decayed wood is summarized in Table II. The FT-IR spectra of decayed hornbeam wood (Fig. 2) demonstrated changes of peaks in fingerprint regions at the naturally infected wood by *T. versicolor*. The comparison between the peaks from the decayed and un-decayed wood revealed that some peaks had severe changes; thus the subtraction of peaks demonstrated that amounts of abortion in the most picks were slightly increased in the fingerprint region. In Fig. 3 the baselines shows high differences between the un-decayed and the decayed wood; while the peaks under the baselines showed an increase of bands and above the baselines indicated prominent declined of the bands.

The FT-IR spectra of decayed and un-decayed hornbeam wood (Fig. 1) demonstrated changes of

the peaks in fingerprint regions of wood decayed by *T. versicolor*. There were many well-defined peaks in the fingerprint region of 1800–500 cm^{-1} . A severe decline took place at 1738 cm^{-1} which caused heavily reduction belonging to unconjugated groups in lignin and carboxylic acid ester hemicelluloses (Takahashi et al. 1989). This behavior was expected for this fungus; because white rot fungi use all three major chemical compositions in the cell wood wall (Eriksson et al. 1990, Eaton and Hale 1993, Schmidt 2006, Kubicek 2013), as a result of fungal activities, a considerable increase at 1594 cm^{-1} peak was seen which is related to carbon–oxygen bond (Pandey and Pitman 2003), a carbon–oxygen bond is a covalent bond between carbon and oxygen (Carey and Sundberg 2007), which was increased after the attack by white rot fungi (Fengel and Wegener 1989). Microbial cellulolysis is a very complex procedure, especially in view of that portion of the mechanism that relates to the degradation of hydrogen bond-ordered cellulose (Wood and Garcia-Campayo 1994). Therefore, cellulolysis by white-rot fungi involves the interaction of enzymes defined as exoglucanases and endoglucanases as well as β -glucosidases (Eriksson and Wood 1985). Carbon and oxygen form terminal double bonds in functional groups are collectively known as carbonyl compounds to which belong such compounds as ketones, esters, carboxylic acids and many more. Compounds with formal C–O triple

TABLE II
The molecular chemical composition changes in
hornbeam wood in sound and naturally infected sample
by white-rot fungus *Trametes versicolor*.

Peaks	Properties	Status	
		Increase	Decrease
1738	C=O un-conjugated groups in lignin and carboxylic acid ester hemicelluloses		+
1594	Conjugated C–O	+	
1504	Aromatic skeletal; Benzene ring vibration in lignin		+
1461	C–H deformation in lignin and carbohydrates		+
	CH ₃ , CH ₂ , benzene ring vibration in lignin		
1423	C–H deformation in lignin and carbohydrates		+
	Aromatic skeletal vibrations combined with C–H in plane deformation		
1372	C–H deformation in cellulose and hemicelluloses		+
	C–H bending vibration in cellulose and hemicelluloses		
1325	C–H vibration in cellulose; C _i –O in syringyl derivatives		+
1238	Syringyl ring; C–O stretch in lignin and xylan		+
1157	C–O–C vibration in cellulose and hemicelluloses		+
	Aromatic skeletal; C–C stretch		+
1096	O–H association band in cellulose and hemicelluloses		
1035	C–O stretch in cellulose and hemicelluloses		+
	C–O of primary alcohol		
892	C–H deformation in cellulose		+
	C _i group frequency in cellulose and hemicellulose		

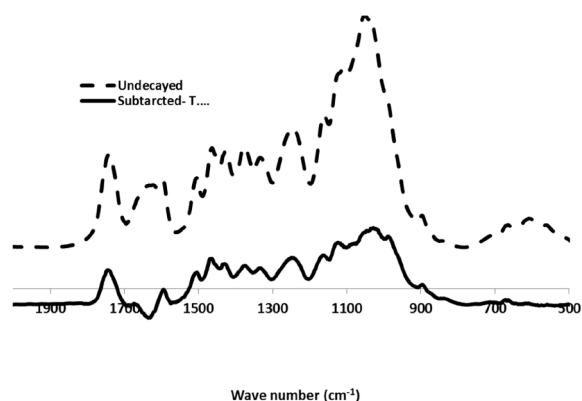


Figure 2 - Subtracted FT-IR spectra of the white-rot decayed hornbeam wood in naturally infected (*in vivo*) by *Trametes versicolor*.

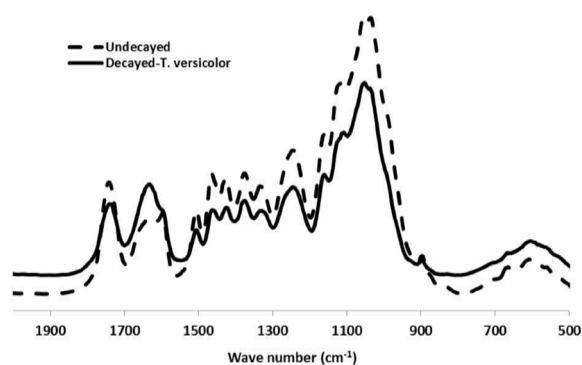


Figure 3 - FT-IR spectra of hornbeam wood naturally infected by *Trametes versicolor* (*in vivo*).

bonds do not exist except for carbon monoxide, which has a very short, strong bond (reported from the standard bond energies). The wood-inhabiting microorganisms use carbon only from enzymatically easily accessible and digestible substrates, like simply constructed sugars, peptides, or fats, or from the storage material starch in the parenchyma cells. The wood decay fungi use carbon additionally from the complex, main components of the woody cell wall, cellulose, hemicelluloses, and lignin (Fengel and Wegener 1989, Schmidt 2006, Kubicek 2013) decreases in intensity observed at around 1504; 1461; 1423; 1372; 1325; 1238; 1157; 1096 cm⁻¹ were related to aromatic skeletal; benzene ring vibration in lignin, deformation in lignin and carbohydrates as well as benzene ring vibration

in lignin; aromatic skeletal vibrations combined with C–H in plane deformation, deformation in cellulose and hemicelluloses and bending vibration in cellulose and hemicelluloses, vibration in cellulose, C₁–O in syringyl derivatives, syringyl ring C–O stretch in lignin and xylan, vibration in cellulose and hemicelluloses, aromatic skeletal and C–C stretch and association band in cellulose and hemicelluloses, respectively. This condition of attack by white-rot fungi has been reported in several studies (Pandey and Pitman 2003, Faix et al. 1993, Mohebbi 2005, Yilgor et al. 2013, Bari et al. 2015a). Different types of hydrolyzing enzyme could be liberated to alter and break linkages in the cell-wall components and release them as small molecules. During this period, many chemical changes occur in these constituents, indicating that the fungus could assimilate them as carbon sources. A strong decrease took place in the carbon–oxygen bonds which was seen at the 1035 cm⁻¹ bond that demonstrated a decrease in primary alcohol. Moreover, a slight decline was observed at 892 cm⁻¹ which belongs to the deformation in cellulose as well as C₁ group frequency in cellulose and hemicellulose (Mohebbi 2005, Bari et al. 2015a).

CONCLUSIONS

The white-rot fungus *Trametes versicolor* caused similar chemical components losses in the xylem of hornbeam wood samples at the advanced decay stages *in vivo*. FT-IR spectroscopy along with chemical analyses evaluated the qualitative and quantitative changes in lignin and carbohydrate components of infected wood by this fungus. The results showed that *T. versicolor* reduced lignin and carbohydrate nearly at the same rate, with a slight preference for lignin which was reflected in a small reduction in the lignin/carbohydrate peak area ratio as decay progressed. Nonetheless, *T. versicolor* had the ability to degrade all three cell wall polymers. Overall, this study proved that this fungus induce

simultaneous white rot decay in hornbeam wood in natural situation (*in vivo*).

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