RESEARCH ON SIZE AND FREQUENCY VARIABILITY OF SOME STANDING TIMBERS' DEFECTS UNDER INFLUENCE OF THEIR BIOLOGICAL ORIGIN

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Abstract

The paper presents results of measurements and observations made on Turkey oak (Quercus cerris) species in the sample areas located in the forest area of Boboștea (Bihor county) and in some other forests (Tășnad Forest District and Dumbrava - Beliu Forest District), which allowed some conclusions on the variability of size and frequency of defects in standing timbers, under the influence of their biological origin.

Key words: timber defect, Turkey Oak tree, biological origin.

INTRODUCTION

Regeneration as other collective processes taking place in the living community life of the forest, takes place under the influence of two groups of factors namely: the biological characteristics of trees to multiply, and environmental conditions where the process takes place. The regeneration process, in general, varies in content and duration depending on the nature of species, overall structure of the forest, origin and provenance of stand, stationary conditions, silvo-technical interventions, etc. (Negulescu et al., 1973).

In relation to the material used for regeneration in the establishment of forest cultivated, one differentiates between seed based regeneration and vegetative regeneration (regeneration from sprouts). Natural seed based regeneration is based on the ability of the trees to produce, for a certain period of time (after the transition to maturity stage), the seed which, once reaching favourable environmental conditions, may generate new samples. Regeneration from sprouts is the characteristic of some wood timber species to form sprouts on the stem or stump after the cutting. If at first, sprouts, have a thrifty stand in size than seed based plants due to the fact that they benefit from the snag rootlet system, after approx. 12-15 years they are surpassed by seed based samples and remain inferior in terms of quantity and quality of the achieved output (Florescu and Nicolescu, 1996).

MATERIAL AND METHOD

Grand total

16B

62A

72C

The research was carried out in the period 2004-2010 in the several locations in the country (Bihor, Arad, Satu Mare). Sampling for assessing the quality of Turkey oaks (*Quercus cerris*) wood was the result of a large number of desk and on field operations.

Investigations were conducted in 14 forest stands of Boboștea forest section, of the Boboștea VII Management Unit. Also, investigations were extended (as a basis for comparison) to the Tășnad Forest District (V Supur Management Unit) in two stands, and Grove - Beliu Forest Unit (I Beliu Management Unit), respectively. Total number of Turkey oaks (*Quercus cerris*) samples measured in the 18 sample areas was 742 (see Table 1).

No	Forest	Subcompart	Station	Forrest	Consisten	Age	Area	Assessed
	District/	ment	type	type	cy	(years)	(m^2)	standing
	Management					-		timber
	Unit							
1	1	3D	6143	7432	0.9	70	2200	30
2	1	5A	6143	7432	0.9	70	2000	47
3	1	6C	6143	7432	0.8	70	2000	45
4	1	8D	6143	7432	0.8	80	2000	43
5	1	34B	6142	7411	0.8	75	2000	46
6	1	55C	6153	7513	0.6	100	2400	30
7	1	69B	6143	7432	0.8	75	2000	50
8	1	77B	6143	7432	0.8	100	2000	46
9	1	83A	6143	7432	0.8	105	2000	58
10	1	87A	6143	7432	0.7	100	2000	46
11	1	87C	6143	7432	0.8	90	2000	44
12	1	87D	6143	7432	0.6	90	2200	30
13	2	124A	8321	7421	0.8	85	2000	51
14	2	128A	8321	7421	0.8	80	2000	47
15	3	9A	6143	7111	0.8	85	2000	40

Sampling in stand where sampling areas were located in

Table 1

Remark for Forest district /Management Unit: 1-Sfânta Maria Forest District/VII Boboștea Management Unit; 2-Oradea Forest District/VIII Mihis Management Unit; 3-Dumbrava Forest District/I Beliu Management Unit; 4-Tășnad Forest District/V Supur Management Unit.

0.7

0.7

0.5

 The location of sample plots was established by using the electronic hypsometer (Vertex IV) to determine slope and crosscut allowance of the side located toward the highest slope. Sample plots have rectangular shape of 2000-2400 m², according to homogeneity in terms of stationary conditions, stand, and the number of constituent trees (at least 30 trees

counted/sampling area), (Dinulică, 2008). The exception is the sample plot area located on the 72C subcompartment (Tăşnad Forest District, Management Unit V. Supur) where, due to silvo-technical interventions, the number of remnant tress was lower (21 samples/sampling area).

Starting from the sorting system (dimensional and qualitative) of oak raw wood according to European standards (Balleux, 2004; Beldeanu, 2008), in the case of Turkey oaks (*Quercus cerris*) samples surveyed we also defined three quality areas on the trunk height, as follows: the 1st quality area includes the first 6m of trunk height, 2nd quality class ranges from 6m up to the tree crown, and the 3rd quality contains the trees' crown. Due to the incidence of quality defects on both quality areas (1st quality area I and 2nd quality area), we defined an intermediate area (see Fig. 1).

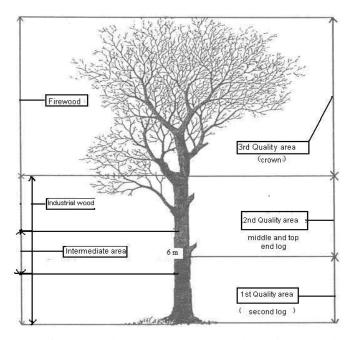


Fig. 1. Quality areas of Turkey oak (*Quercus cerris*) samples reviewed - (adaptation by Oak raw wood, according to European standards): 1st Quality area (0-6 m), 2nd Quality area (ranging from 6m to the crown), 3rd Quality area (crown), intermediate area

In relation to the shape of the trunk, associated with other defects, we defined the following types of the assessed Turkey oak (*Quercus cerris*) samples: form **1a**-straight cylinder without defects, form **1b**-straight cylinder, with other defects, form **2a**-straight, slight curved, without other defects, form **2b**-straight slightly curved, with other defects, form **3a**-sinuous, curved shape without defects, and form **3b**-sinuous, curved with other defects.

Desk work consisted in computing and interpreting data collected from the field; thus they were centralized by source provenance and diameter classes. The making of charts was performed by using MS Office Excel software.

RESULTS AND DISCUSSIONS

In order to capture existing variations between the biological origins of the trees identified in the surveyed area (seed and sprouts), regarding the distribution of some of defects, we used the bon-parametric Kruskal-Wallis one-way analysis of variance by ranks.

• The statistical significance of differences among the biological origins of trees as regards the appearance and localization of exterior rot, pruning height, height of first green branch, on tree trunk shape, quality class, the root-sweeling presence, undercovered knots height on the trees height, respectively, show the biological origins of trees and represent highly significant statistical factor with impact on trees (see Tables 2, 3, 4, 5, 6, 7, and 8).

Table 2

Statistical significance of the influence of trees biological orig	gin
on rot occurrence and localization	

Kruskal-Wallis anal	Kruskal-Wallis analysis results: H=28.56046***, N = 742 trees, f=1 degree of freedom,				
	p<0.1%				
Transg	Transgression probability matrix on the individual variations				
	caused by biological origin of trees				
	Seed origin Sprout origin				
Seed origin 0.012257					
Sprout origin	0.012257				

Table 3

Statistical significance of the influence of trees biological origin on pruning height

Kruskal-Wallis anal	Kruskal-Wallis analysis results: H=18.28200***, N = 742 trees, f=1 degree of freedom,			
	p<0.1%			
Transg	Transgression probability matrix on the individual variations			
_	caused by biological origin of trees			
	Seed origin Sprout origin			
Seed origin 0.000019				
Sprout origin	0.000019			

Table 4

Statistical significance of the influence of trees biological origin on the height of first green branch

Kruskal-Wallis anal	Kruskal-Wallis analysis results: H=18.28200***, N = 742 trees, f=1 degree of freedom,			
	p<0.1%			
Transg	Transgression probability matrix on the individual variations			
	caused by biological origin of trees			
	Seed origin Sprout origin			
Seed origin	Seed origin 0.000019			
Sprout origin	Sprout origin 0.000019			

Table 5

Statistical significance of the influence of trees biological origin on trunk shape				
Kruskal-Wallis anal	Kruskal-Wallis analysis results: H=18.02437***, N = 742 trees, f=1 degree of freedom,			
	p<0.1%			
Transg	Transgression probability matrix on the individual variations			
	caused by biological origin of trees			
Seed origin Sprout origin				
Seed origin 0.000050				
Sprout origin	0.000050			

Table 6

Statistical significance of the influence of trees biological origin on quality class of trees				
Kruskal-Wallis anal	Kruskal-Wallis analysis results: H=29.19033***, N = 742 trees, f=1 degree of freedom,			
	p<0.1%			
Transg	Transgression probability matrix on the individual variations			
	caused by biological origin of trees			
	Seed origin Sprout origin			
Seed origin 0.000004				
Sprout origin	0.000004			

Table 7

Statistical significance of the influence of trees biological origin on incidence of root-sweeling of the trees

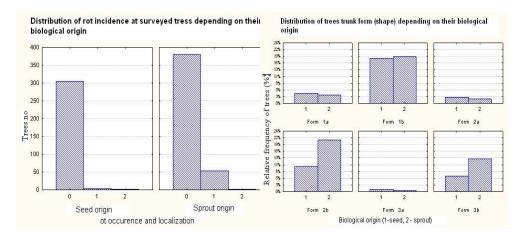
Kruskal-Wallis anal	Kruskal-Wallis analysis results: H=20.69181***, N = 742 trees, f=1 degree of freedom,			
	p<0.1%			
Transg	Transgression probability matrix on the individual variations			
_	caused by biological origin of trees			
	Seed origin Sprout origin			
Seed origin 0.007908				
Sprout origin	0.007908			

Table 8

Statistical significance of the influence of trees biological origin on the localization of undercovered knots by height of trees

Kruskal-Wallis anal	Kruskal-Wallis analysis results: H=23.10836***, N = 742 trees, f=1 degree of freedom,			
	p<0.1%			
Transg	Transgression probability matrix on the individual variations			
-	caused by biological origin of trees			
	Seed origin Sprout origin			
Seed origin 0.000072				
Sprout origin	0.000072			

Figure no. 2 shows that the exterior rot occurs in a higher percentage at trees of sprout biological origin (in 53 cases in 1^{st} Quality class area, an in 1 case in 2^{nd} Quality class), while the seed origin trees are less affected by this defect (3 cases for the 2^{nd} Quality area and 2 cases for the 2^{nd} Quality area).



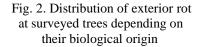


Fig. 3. Distribution of tree trunk shape depending on their biological origin

As regards the shape of Turkey oaks (*Quercus cerris*) trunk (see Figure 3), one may observe the existence of valuable shapes **1a**, **2a** and **3a** respectively, in a higher proportion in case of seed related biological origin. Sprout origin samples dominate in the shapes **1b**, **2b** and **3b**, in those with different combinations of defects.

Seed based samples are better pruned (H pruning=15.94 m, on average) compared to those with sprouts/shoots of biological origin (H pruning=14.53 m, media), the difference being of roughly 1.5 m.

Regarding the differences between quality classes of trees, depending on their biological origin (see Figure no 4), seed based samples remain almost at the same percentage as in the 1^{st} Quality class (20.35%) and the 2^{nd} quality class (19.27%), while the sprouts based samples maintain different percentages i.e. 40.43% at 2^{nd} quality class and 15.76% in 1^{st} quality class.

The root-swelling affects, in larger proportions, the seed based samples (60 cases) compared with sprout based ones (36 cases).

Regarding the occurrence of undercovered knots (see Fig. no 5), by biological origins, sprouts based samples are much affected as seed based ones. Higher frequencies appear within the intermediate area (21.83%) and the 1^{st} Quality class area (1.88%), in case of sprouts based biological origin of tree.

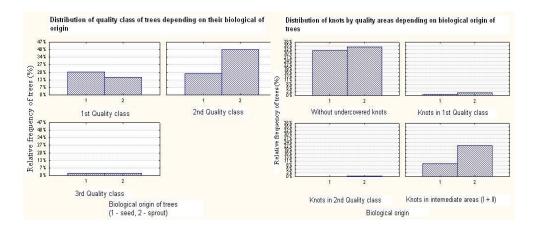
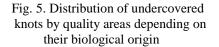


Fig. 4. Distribution of quality class of trees depending on their biological origin



CONCLUSIONS

One may conclude that the biological origin of the Turkey oak trees (*Quercus cerris*) within the surveyed area represents a **statistical highly significant factor** as regards: the occurrence and localization of trees' exterior rot, pruning height, height of first green branch, trees trunk shape, quality class of trees, root-swelling presence and localization of undercovered knots, respectively.

Seed based biological origin has many advantages in comparison with the sprout one, as follows:

• A reduced incidence of exterior rot by quality areas of trees (1.61% versus 12.47%, in the case of sprout based ones);

• Existence of **1a**, **2a** and **3a** valuable forms (as regards trees trunk), in a larger proportion at seed based biological origin samples, to the detriment of poor forms **1b**, **2b** and **3b**, which are in a larger proportion a sprout biological origin;

• Seed samples are better pruned (mean=15.94m) compared with the sprout based ones (mean=14.53 m);

• Seed origin samples remain almost at the same percentage both in the 1st Quality class (20.35%) and 2nd Quality class (19.27%), while those of sprout origin remain in different percentages, 40.43% in 2nd Quality class, and 15.76% at 1st Quality class;

• Regarding the occurrence of undercovered knots, of biological origin, seed origin samples are less affected compared to sprouts origin, higher frequencies occurring within the intermediate area

(21.83%) and the 1st Quality class (1.88%), in case of sprouts biological origins excrescences;

Sprout biological origin show the following advantage, compared to those of seed biological origin:

• A reduced root-swelling incidence 8.31%, as against 19.41%, in the case of seed origin samples.

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