



Journal of Forest Research

ISSN: 1341-6979 (Print) 1610-7403 (Online) Journal homepage: https://www.tandfonline.com/loi/tjfr20

An examination on the site—family interaction of tree height of open-pollinated families of hinoki (*Chamaecyparis obtusa*) plus trees in the Kansai Forest tree breeding region

Hiroo Yamada & Tomiyasu Miyaura

To cite this article: Hiroo Yamada & Tomiyasu Miyaura (2004) An examination on the site—family interaction of tree height of open-pollinated families of hinoki (*Chamaecyparis obtusa*) plus trees in the Kansai Forest tree breeding region, Journal of Forest Research, 9:1, 81-84, DOI: <u>10.1007/s10310-003-0048-9</u>

To link to this article: https://doi.org/10.1007/s10310-003-0048-9



Published online: 01 Feb 2004.

_	_
ſ	
L	0
_	

Submit your article to this journal \square

Q

View related articles 🗹

views: 2

SHORT COMMUNICATION

Hiroo Yamada · Tomiyasu Miyaura

An examination on the site-family interaction of tree height of openpollinated families of hinoki (*Chamaecyparis obtusa*) plus trees in the Kansai Forest tree breeding region

Received: July 22, 2002 / Accepted: May 30, 2003

Abstract Stability analyses to evaluate site-family interaction for tree height at 15 years old were conducted in trials of open-pollinated families of hinoki (Chamaecyparis obtusa Endl.) plus trees established in the Honshu and Shikoku area of the Kansai Forest tree breeding region, Japan. The Honshu area included 103 sites (226 families), and the Shikoku area included 27 sites (62 families). Analyses of variance indicated that the interaction between site and family was not significant in the Honshu area, whereas in the Shikoku area the interaction was significant. The stability analyses regressed the family-site means against site means. The linear regression analysis was carried out using the data for 164 families that were tested at five or more test sites from the 130 test sites. Approximately 93% of families had values of linear regression coefficient that were not significantly different from unity. These results suggest that almost every open-pollinated family of hinoki plus trees have average stability and are equally well adapted to good and poor sites. The differences in the reaction of a family to site productivity are not the main causes of the interaction.

Key words Genotype–environment interaction · Height · Hinoki plus tree · Stability

Introduction

Genotype-environment interaction is the differential response of genotypes to different environments. Such inter-

H. Yamada (🖂)

Forest Tree Breeding Center, 3809-1 Ishi, Juo-cho, Taga-gun, Ibaraki 319-1301, Japan Tel. +81-293-32-7000; Fax +81-293-32-7306

e-mail: hirooy@affrc.go.jp

T. Miyaura Ryukoku University, Ohtsu, Japan actions complicate testing and selection in tree improvement programs and reduce overall genetic gains. The genotype-environment interaction can be diminished in two ways (Shelbourne 1972): (1) by creating groups of essentially homogeneous environments and selecting cultivars suited to each environment (Shelbourne and Low 1980; Kurinobu 1984; Clair and Kleinschmit 1986; Nishimura and Tajima 1993; Miyaura 1994) and (2) by developing stable cultivars which perform consistently over a range of environments (Finlay and Wilkinson 1963; Owino 1977; La Farge and Kraus 1981; Yeiser et al. 1981; Clair and Kleinschmit 1986; Li and McKeand 1988; McKeand et al. 1990; Yamada and Miyaura 2002).

Many studies in Japan were conducted based on the first method of diminishing interaction (Kurinobu 1984; Nishimura and Tajima 1993; Miyaura 1994). Finlay and Wilkinson (1963) reported that a linear regression coefficient (a_i) of growth traits against site productivity indicates stability and relative adaptability of a tree variety to a site. They considered a variety with a coefficient value near unity to be of average stability and equally adapted to good and poor sites. A variety with a value greater than unity is unstable and better adapted to good sites. A variety with a value less than unity is stable and better adapted to poor sites. They expected that varieties having similar values of a_i have less fluctuation in the ranking of height among forest sites. Yamada and Miyaura (2002) used the linear regression coefficient of growth traits against site productivity to examine the stability and relative adaptability to different environments within a breeding district by clonal lines and open-pollinated families of sugi (Cryptomeria japonica D.Don) plus trees. However, the examination of genotypeenvironment interactions of the growth traits of hinoki plus trees (Chamaecyparis obtusa Endl.) has not been conducted in the Forest Tree Improvement programs in Japan.

The objective of this study was to examine the interaction between family and site concerning to the tree height of open-pollinated families of hinoki plus trees in the Kansai Forest tree breeding region. We used the techniques of Finlay and Wilkinson (1963) and Yamada and Miyaura (2002).



Fig. 1. Locations of progeny test sites of hinoki plus trees

Materials and methods

Test sites and families

The analysis was carried out using the height data at 15 years old of 287 open-pollinated families. The number of the analyzed test sites was 130 (Fig. 1). The tests were established in randomized complete block designs of mostly three (up to nine) replications. The number of families per test site ranged from 5 to 45. Overall, 257 families out of 287 were tested at two or more test sites.

The Kansai Forest tree breeding region is administratively divided into six breeding districts: Kinki, Setouchi, East Nihonkai, West Nihonkai, North Shikoku, and South Shikoku. A total of 160 families were measured only in the breeding district from which they had been selected, 102 families were measured in both the selection district and a neighboring district, and 25 families were measured in three or more districts. Although 127 families out of 287 were tested in two or more districts, only one family was tested in both the Honshu area (Kinki, Setouchi, East Nihonkai, and West Nihonkai breeding districts) and the Shikoku area (North Shikoku and South Shikoku districts). Therefore, each area was analyzed separately.

Analysis of variance

SPSS (2001) was used to conduct an analysis of variance of the plot mean of tree height. A linear model for the analysis is:

$$y_{ijk} = \mu + \alpha_i + \alpha \beta_{ij} + \gamma_k + \alpha \gamma_{ik} + \varepsilon_{ijk}$$

where μ is the population mean across the tests, α_i is a fixed effect of the *i*th test, $\alpha\beta_{ij}$ is a fixed effect of the *j*th replication in the *i*th test, γ_k is a random effect of the *k*th family, $\alpha\gamma_{ik}$ is a random effect of the interaction between the *i*th site and the *k*th family, and ε_{ijk} is a random plot error of y_{ijk} . We adopted typeII sum of squares of SPSS (Searle 1987). That is equivalent to the sum of squares derived by the



Fig. 2. Distribution of test site productivity in the six breeding districts

Henderson's method 3 (Henderson 1953; Kurinobu et al. 1984).

Regression analysis

We examined the relationships between site productivity and height of the families using the height data for 164 families tested at five or more test sites from 287 families. Linear regression equations of height against site productivity were estimated with the ordinary least squares method:

$$y_{ii} = a_i X_i + b_i$$

where y_{ij} is the mean height of the *j*th family at the *i*th test site, X_i is the site productivity of the *i*th test site, a_j is the linear regression coefficient of the *j*th family, and b_j is a constant specific to the *j*th family. Site productivity was evaluated as the mean height in each of the 130 test sites. We assumed that the site productivity shows the total effect of environment factors on the tree growth, such as climate, soil type, and management (Finlay and Wilkinson 1963). The confidence interval at the 95% significance level of a_j for each family was calculated to examine whether or not the value of a_i is significantly different from unity.

Results

Difference of site productivities

The test site productivity of each breeding district is shown in Fig. 2. There was a considerable range in productivities among the 130 test sites: from 2.37 m to 9.78 m. Each breeding district had test sites with productivities ranging from high to low.

Analysis of variance

The results of analysis of variance are shown in Table 1. The height was significantly different among families in both areas. There was no significant interaction between test site and family in the Honshu area, whereas in the Shikoku area the interaction was significant.



Fig. 3. Linear regression coefficients and their confidence interval at the 95% significance level plotted in ascending order of the value of the regression coefficient within each area. *Black dots* and *vertical bars* represent the linear regression coefficients and their confidence intervals, respectively

Table 1. Analyses of variance for the plot mean of tree height for families at the 103 sites in the Honshu area and the 27 sites in the Shikoku area

Source	Honshu area		Shikoku area	
	df	Mean square	df	Mean square
Site	102	75.65***	26	59.41***
Replicate (site)	198	5.01***	67	5.17***
Family	225	0.67***	61	0.79*
Site \times Family	1322	0.52 ^{NS}	279	0.50*
Error	3040	0.49	980	0.42

*** Significant at P < 0.001, * Significant at P < 0.05, NS nonsignificant

Regression analysis

Linear regression analysis showed a significant correlation (P < 0.05) between height and site productivity for 160 families out of 164 (r = 0.740 to 0.999). The values of the linear regression coefficient a_j of these 160 families ranged from 0.643 to 1.355 (Fig. 3). Eleven families out of 160 had values of a_j that were significantly different from unity at the 5% significance level (the Honshu area: 9 out of 130, the Shikoku area: 2 out of 30). The value of a_j of seven of these families was significantly lower than unity, and that of four families was significantly greater than unity.

Figure 4 shows the relationship between the confidence interval (95% significance level) of the linear regression coefficient and the range of productivity of the test sites within which each family was tested. The confidence interval tends to decrease as the range of site productivities increases.

Discussion

The growth traits of each plus tree are evaluated by leastsquare estimates from the data of multiple progeny tests in each breeding district, and the plus trees are ranked in the growth order in the Forest Tree Improvement programs of Japan. This ranking is used for the purpose of roguing seed



Fig. 4. Relationship between confidence interval at the 95% significance level and the range of productivity of the test sites within which each family was tested

orchards and choosing parents for the second generation breeding population. In the present study, each breeding district had test sites with considerable variations in productivity (Fig. 2). Each breeding district will include heterogeneous environments. This fact suggests that the investigation of genotype–environment interaction is required in order to apply the ranking to the Forest Tree Improvement programs. If there is strong genotype–environment interaction within a breeding district, the ranking may show great fluctuation among forest sites. In this case, the stability and relative adaptability to different environments must be examined for each plus tree together with the ranking.

We obtained several important facts concerning the existence and magnitude of any genotype-environment interactions of open-pollinated families of hinoki plus trees. There was no significant interaction between test site and family in the Honshu area, whereas in the Shikoku area the interaction was significant (Table 1). Only 11 families out of 160 had values of the linear regression coefficient a_i that were significantly different from unity at the 5% significance level (Fig. 3). Approximately 93% of families had values of *a_i* that were not significantly different from unity. These results suggest that almost all families had average stability and were equally well adapted to good and poor sites (Finlay and Wilkinson 1963) in spite of the fact that there was significant interaction in the Shikoku area. We expect that the open-pollinated families of hinoki plus trees show little fluctuation in the ranking of height among forest sites.

The confidence interval at the 95% significance level of the linear regression coefficient tended to decrease as the range of site productivities increased (Fig. 4). The accumulation of data from many test sites with different productivities is now in progress. The analysis using the accumulated data in the future may reveal any other families having a value of a_j significantly different from unity. Moreover, the analyzed data in this study included considerable differences in testing frequencies among the families. Further study is required to examine the existence and magnitude of any genotype–environment interaction. The 11 families showing values significantly different from unity must be cautiously treated. Since these families are either too sensitive or fairly stable to different site productivities, these families may have great fluctuations in the ranking of height among forest sites (Finlay and Wilkinson 1963). The value of regression coefficient a_j , as an index of stability and relative adaptability of tree varieties to sites, suggests an important indicator to prevent the reduction of overall genetic gains.

Acknowledgments We are grateful to the staff of the Prefectures and Forest Agency for their kind help in collecting data, and to Y. Ando, K. Nishimura, and M. Kubota for their kind help with analyzing the data. We also would like to express our thanks to the two anonymous reviewers for their many invaluable comments which helped us to improve the manuscript.

Literature cited

- Clair JB, Kleinschmit J (1986) Genotype–environment interaction and stability in ten-year height growth of Norway spruce clones (*Picea abies* Karst.). Silvae Genet 35:177–186
- Finlay KW, Wilkinson GN (1963) The analysis of adaptation in a plant breeding programme. Aust J Agric Res 14:742–754
- Henderson CR (1953) Estimation of variance and covariance components. Biometrics 9:226–252
- Kurinobu S (1984) A methodological study on the analysis of family trial plantations of Japanese larch (in Japanese with English summary). Bull For Tree Breed Inst 2:1–60
- Kurinobu S, Kaneko T, Shingai Y, Ohba K (1984) A classification of the optimum breeding zone based on the unbalanced data of prog-

eny trials conducted on several sites (in Japanese). J Jpn For Soc 66:109–112

- La Farge T, Kraus JF (1981) Genotype \times environment interaction of loblolly pine families in Georgia, USA. Silvae Genet 30:156–162
- Li B, McKeand SE (1988) Stability of loblolly pine families in the southeastern US. Silvae Genet 38:96–101
- McKeand SE, Li B, Hatcher AV, Weir RJ (1990) Stability parameter estimates for stem volume for loblolly pine families growing in different regions in the southeastern United States. For Sci 36:10– 17
- Miyaura T (1994) On the subdivision of the Kanto breeding area based on the analysis of data in 10-yr-old clonal tests of sugi (*Cryptomeria japonica*) (in Japanese with English summary). For Tree Breed 170:8–12
- Nishimura K, Tajima M (1993) Classification of tree breeding districts in the Kyushu region by the growth of sugi (*Cryptomeria japonica* D. Don) plus trees (in Japanese with English summary). J Jpn For Soc 75:493–500
- Owino F (1977) Genotype \times environment interaction and genotypic stability in loblolly pine II. Genotypic stability comparison. Silvae Genet 26:21–26

Searle SR (1987) Linear models for unbalanced data. Wiley, New York

- Shelbourne CJA (1972) Genotype–environment interaction: its study and its implication in forest tree improvement. IUFRO Genetics– SABRAO joint symposia, Tokyo. B-1:1–28
- Shelbourne CJA, Low CB (1980) Multi-trait index selection and associated genetic gains of *Pinus radiata* families at five sites. N Z J For Sci 10:307–324
- SPSS (2001) SPSS 11.0J for Windows, user's guide. SPSS, Chicago
- Yamada H, Miyaura T (2002) Relationships between stand productivity and height growth for clonal lines and open-pollinated families of sugi (*Cryptomeria japonica* D. Don) plus trees selected from Kansai Forest Tree Breeding Region (in Japanese with English summary). J Jpn For Soc 84:50–53
- Yeiser JL, van Buijtenen JP, Lowe W (1981) Genotype \times environment interactions and seed movements for loblolly pine in the western gulf region. Silvae Genet 30:196–200