

A SURVEY OF *ULMUS LAEVIS* IN FLANDERS (NORTHERN BELGIUM)

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ABSTRACT. -This study reports on the growth sites of *Ulmus laevis* Pallas in Flanders, Northern Belgium. The sites reported by former authors were reviewed and six newly discovered sites in Flanders are described, including population size and habitus. All together, a summarising picture of the distribution of this rare tree species in Northern Belgium is drawn. Gene conservation strategies are discussed.

KEY WORDS. — *Ulmus laevis*. conservation, Dutch elm disease, Flanders.

INTRODUCTION

Ulmus laevis Pallas, the European white elm, is a broad-leaved riparian tree species with a central and eastern European distribution and one of the three indigenous elm species in Belgium, next to *U. minor* Mill. and *U. glabra* Huds. (LAMBJON *et al.* 1998). Along with the American elm (*V. americana* L.), *U. laevis* belongs to the section *Blepharocarpus*, whereas the other two European elm species, *U. glabra* and *V. minor*, belong to the section *Ulmus*. *U. laevis* does not easily hybridise with the other European elm species, and it is self-incompatible (MJTTEMPERGER & LA PORTA 1991). It tolerates wet soils and periodic flooding, and typically occurs in damp low-lying areas and as a component of riparian forests (WHYTELEY 2004). In Western Europe, deforestation and drainage of flood plains for agriculture and industry have severely diminished the area of suitable habitat for *U. laevis*. Habitat fragmentation is a major threat for the mostly marginal populations

(COLLIN *et al.* 2000, GOODALL-COPESTAKE *et al.* 2005). Genetic studies carried out in Finland and Sweden, at the northern fringe of its range, suggest that genetic drift may have caused substantial differentiation among the small populations of *V. laevis* (MATTILA & VAKKARJ 1997, WHITELEY 2003, WHITELEY 2004).

Although *V. laevis* is susceptible to Dutch elm disease (DED), caused by the fungal agent *Ophiostoma novo-ulmi*, it is not thought to be in immediate danger from the disease (WHITELEY 2004). Experiments with elm bark beetles (*Scolytus scolytus* and *S. multistriatus*) acting as vectors for the fungal pathogen, showed that *U. laevis* is far less attractive for the beetles than *U. minor* (SACCHETTI *et al.* 1990, WEBBER 2000). In contrast to the other *Ulmus* species, no repon exists of *V. laevis* in Flanders affected by DED in natural conditions. Therefore, there is a raising interest from foresters and land managers to plant the species. Conservation of *U. laevis* is of ecological as well as economic importance. The trees

serve as habitats for other organisms. are highly valued as landscape trees and produce high-quality wood (WHYTELEY 2004).

In the atlas of the Belgian and Luxembourg flora (VAN ROMPAEY & DELVOSALLE 1972) *V. laevis* is totally absent in Flanders. LAMBINON *et al.* (1998) did not mention any growing sites either. A lack of knowledge of the species most probably is the main reason why it has been overlooked in most vegetation surveys in the past. Recently, the distribution range in Europe has been published on the web, incorporating several recently described growth sites in Flanders (<http://www.ipgri.cgiar.org/networks/euforgen>).

A global inventory of autochthonous woody plants in Flanders is being carried out under the authority of the Forest and Green Area Division of the Flemish Community since 1997 (MAES & RÖVEKAMP 1998, MAES & RÖVEKAMP 2000, MAES *et al.* 2003, ÜPSTAELE 2001, RÖVEKAMP & MAES 1999, RÖVEKAMP & MAES 2000, RÖVEKAMP *et al.* 2000, VANDER MIJNSBRUGGE 2003), following an inventory method based on MAES (1993) (see Material and methods). Because a few decades ago *V. laevis* was still unknown in Flanders (VAN ROMPAEY & DELVOSALLE 1972), these fairly recently published inventories urged the elaboration of a detailed overview of the remaining populations in Flanders.

Here, we report on six newly discovered growth sites of *Ulmus laevis*. In addition, the sites described in the inventories of autochthonous trees and shrubs in Flanders (MAES & RÖVEKAMP 1998, MAES & RÖVEKAMP 2000, MAES *et al.* 2003, ÜPSTAELE: 2001, RÖVEKAMP & MAES 1999, RÖVEKAMP & MAES 2000, RÖVEKAMP *et al.* 2000) were revisited by the authors and outlined in the same way. An overall and summarizing picture of the distribution of this rare tree species in Flanders is thus drawn. We also discuss strategies for gene conservation.

MATERIAL AND METHODS

IDENTIFICATION OF *U. LAEVIS*

For the identification of *U. laevis*, LAMBINON *et al.* 1998 was followed. The stalked flowers and nut of *U. laevis* make the species unambiguously distinguishable

from *U. minor* and *U. glaberrima* (LAMBINON *et al.* 1998). Also, the leaf shows a typical soft velvety lower surface (LAMBINON *et al.* 1998).

METHOD OF INVENTORY OF AUTOCHTHONOUS TREES AND SHRUBS

We followed the method of MAES (1993). First, old forests were located on topographical maps, using historical forest evolution maps (DE KEERSMAEKER *et al.* 2001). These maps differentiate forested and non-forested areas, but no indication of tree species can be deduced. Together with information on flora, soil condition and geography, sites were chosen to be visited in the field. Once in the field, several criteria were used to evaluate autochthony (MAES 1993). Criteria regarding the site were: (i) the site was afforested on the Ferraris map (18th century) or on other historical topographic maps (DE KEERSMAEKER *et al.* 2001); (ii) the site appeared to have remained undisturbed; (iii) the ecological conditions were similar to the conditions in the natural area of prevalence of the species; (iv) plant species indicative of old forest and ancient, undisturbed woodlands (TACK *et al.* 1993) were present in herbaceous, shrub or tree layer and (v) the site was located within the natural distribution range of the species. Important criteria regarding the tree or shrub were: (i) the tree or shrub is no cultivated variety and (ii) is old.

As the inventory lacked some details, such as the habitus of the trees or shrubs, the growth sites where *U. laevis* was described were revisited by the authors and reported in detail (Table 1). Through this inventory, knowledge on identification of *U. laevis* began to spread among botanists and other field workers. Other new growth sites, not yet published, were recorded by personal communication to the author.

RESULTS

All sites where *V. laevis* trees were discovered were visited by the authors and described in detail. The available data are summarised in Table 1.

In De Panne, the white elms grew in a tiny strip of forest on the inland edge of the dunes. The coppice stools were scattered along a small naturally meandering stream. The circumference at soil level of most stools averaged between 2 and 3 m. Because of the strong historical human influence in the dune vegetation in Flanders, the autochthony of this site is questionable.

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TABLE 1

Populations and relic individuals of *U. laevis* in Flanders (new findings from this study are in italics)

Location	n° of individuals and/or coppice stools (J)	habitat	circumference (cm)	height (m)	notes
nr. / onnr	15	edge of forest	1.5-11	1.1-1.3	<i>rhis sflrh'</i>
1 oct:ert	1	edge of forest	1.5-11	1.1-1.3	<i>rhis sflrh'</i>
8mkel	8	edge of meadow	1.5-11	1.1-1.3	<i>rhis sflrh'</i>
Smilde	27	edge of small forest in valley	0.6 m (pollard)	1 mld	Maes & Rövekamp 1998
Houten 1	1	forest in valley	2.5-11 (coppice)	1.1-1.3	Maes & Rövekamp 1998
Silft-Ur-I-clls	1	edge of forest	7 pollards and 2 normal trees	1.1-1.3	Maes & Rövekamp 1998
lrwsfI	1	in hedge	2 pollards up to 3.5 cm. coppice stool	1.1-1.3	Maes & Rövekamp 1998
Znc1:w!2	2	in forest	2.5-11	1.1-1.3	Maes & Rövekamp 1998
Zandhoven Lille	5/10	edge of forest	1 normal tree	1.1-1.3	Maes et al. 2003 Maes et al. 2001
Bertem (5)	1	large forest edge	1 stool	1.1-1.3	Opstaele 200
Hasselt	1	large forest edge	1 pollard	1.1-1.3	Opstaele 200
Heers 1	1	hedge	coppice and 1 up to 4.5 m	1.1-1.3	Opstaele 2001

Ik-ers 2	1'i	;-	small forest	low po!ards coppice and norm_11_1_tie	coppice up to 8 m normal trees up to 2 m	urn	Opstaele 2001
1 [re" 1	7	. an'	s111;1J1fnrst	coppice and nonna! trees	coppice up to 4 m normal trees up to 1, > m	11111.ug	OJ"tarle 2fl11
Hcrrs 4	l()	a/h	forest and forest edge	normal trees	Up to 2 m	um. 11g	Opstaele 2001

Notes. (1) Both an individual and a coppice stool represent one genotype. The different stems belonging to one coppice stool are not measured individually and are also not indicated as separate individuals. (2) Quality for autochthony: a. autochthonous with certainty, b. probably autochthonous. (3) Circumference is measured at breast height for normal trees. For coppice stools the circumference is measured at soil level, surrounding all individual stems belonging to one and the same stool. When more than 2 individuals and/or stools are present, only the largest circumference is indicated. (4) Presence of other elm species in immediate neighbourhood: 11111. *U. minor*; ug. *U. glaberrima*; uh. *U. x hollandica*. (5) Not found on a revisit of the site by the authors in the winter of 2003-2004.

In Ploegsteert, the white elms grew at the bottom of a forested slope, next to a road. There are no streams in the immediate neighbourhood, but the forested slope is mapped as a historical forest on the Ferraris map (DE KEERSMAEKER *et al.* 2001) and holds many indications of old and undisturbed forested sites. For instance, next to the white elm we found the extremely rare rose species *Rosa stylosa* and *R. micrantha*.

The white elm stools in Ruislede were found in an inundation area of a small stream. Although the circumference of the elm pollards indicated a reasonable age, no large stools or pollards of other woody species were present in the neighbourhood. The area is mapped as a historical forest.

In Brakel, the five recorded individuals were pollarded trees. They occurred in two clumps, with about 1 m between the stems. These two clumps possibly represent only two genotypes. The trees grew on a small slope (a difference of 1 m in altitude) that separated the inundation area of a small stream, now planted with poplars, and a higher, levelled meadow. The site is part of a relatively large historical forest.

In Sint-Lievens Houtem, white elms were found at two sites 1 km apart. One location is a humid valley of a small stream where one old coppice and a probably younger pollard were present. Other elm species were present in this valley, including the hybrid *U. x hollandica*. The second site is located along a hillside. Here, the pollarded elm were planted on a former edge between a meadow and a forest. Other species were present in this row, including *Crataegus monogyna* (not pollarded) and *Carpinus betulus* (pollarded). The meadow is now planted with poplars and the pollard row of elms is suffering from shade and the lack of maintenance. A few pollards are already torn apart.

In Zoersel, several white elms grew in a large valley that consists of a historical mosaic of forested strips, wood banks and humid meadows. Most probably, part of the area were naturally inundated on a regular basis, which makes the site a very likely natural habitat for white elm. The circumference of one of the pollarded elm (3.75 m) indicates an old age.

Zandhoven is close to Lille, a neighbouring village of Zoersel. Here the white elms grew on

the bank of a naturally meandering stream, which is part of a historical forest. It concerns large stools (circumference up to 7 m) indicating an old age.

In Lille, one old elm and several younger white elm trees grew scattered in a small historical forest that is transected by a small stream. Several of the young trees grew further than 10 m apart from the older tree. Possibly, the younger trees represent natural rejuvenation.

In Hasselt, two big pollards were recorded in the middle of a large forest complex. Both grew at a different border of a forest stand where small streams are present, possibly man-made to irrigate the stands.

In Heers white elms grew at several sites. The high number of recorded trees (68) suggests that this area is possibly the most valuable relic population in Flanders. Pollards, coppice and normal trees are present in small forest patches, scattered in between larger agricultural fields.

In all, 130 white elm trees that have a possible autochthonous status were recorded in Flanders (Table 1). They grew in forests, forest edges and wood banks that are mostly located on stream banks or in or near inundation areas. No growth site was present along the main rivers Schelde, IJzer or Maas. Human influence was prominent in many cases through pollarding (7 of 16 sites, Table 1) and coppicing (10 of 16 sites, Table 1). Coppice stools were measured with a circumference at soil level up to 8 m (Table 1). Low pollards in Heers reached a circumference of more than 4 m, whereas the pollards in Zoersel, pollarded at a height of 2 m, had a circumference at breast height of 3.75 m (Table 1).

DISCUSSION

The results of the survey clearly reveal that *U. laevis* is a rare and endangered species in Flanders. Nearly all natural riparian forests have disappeared along the main rivers resulting in the extinction of typical natural habitat. As a consequence, natural populations of *U. laevis* are mostly reduced to relic individuals. The situation in Flanders is similar to Germany, as reported by MACKENTHLEN (2004): most *U. laevis* trees were

found in restricted habitats within agricultural landscapes.

Here, the term autochthonous is used to indicate elms that regenerated spontaneously or that were reproduced with local material, counting from the last ice age (HEYBROEK 1992). Most remnant populations of *V. laevis* in Flanders are considered autochthonous in the inventory (quotation for autochthony in Table 1). Apart from other criteria to evaluate autochthony (see Material and methods), the large circumference of coppice stools of *V. laevis* in Zandhoven and Heen (Table 1) are important criteria for the autochthonous quotation. Also, spontaneous natural regeneration, another criterion that may indicate autochthony, can be inferred from the population of Lille (Table 1), where one mature tree (circumference at breast height = 1.2 m) is surrounded by several younger trees. Also, the different populations in Heers show a differentiation in circumference, including younger trees that most probably were not planted and therefore indicate possible natural regeneration. On the other hand, some characters of the populations indicate old cultural heritage rather than autochthony. Flanders is the western limit of the range and nearly all growth sites of *V. laevis* testify of human influence through coppice, pollarding and/or the presence in the immediate neighbourhood of *V. glabra* and *V. minor*, two species with different ecological requirements (Table 1). Although these arguments may question the autochthonous status in Flanders, the value of the remnant population as old cultural heritage is beyond doubt.

The risk is apparent that the extremely low numbers of trees in the remnant population in Flanders may have resulted in genetic depauperation. Therefore, gene conservation is essential (JENSEN *et al.* 1999, COLLIN 2003, GODALL-COPESTAKE *et al.* 2005). A clonal archive with grafted elms is currently being developed at the Institute for Forestry and Game Management, as a gene conservation programme. Because of the reduced numbers of trees per population, all individuals are sampled for grafting, except the large population in Heers, for which only a subsample is sufficient. The resulting living collection is a static storage of genotypes, but it opens possibilities

for several more dynamic conservation approaches. As grafting trials with summer cuttings showed reasonable success rates (data not shown), cuttings will be taken from the clonal archive as basic material for a seed orchard in the near future. The genetic variability in the offspring of the seed orchard will be used in new forestation and other plantings. It will also be possible to restock remnant populations with vegetatively propagated plants from selected clones from the archive. For example, the same fluvial basin or from the neighbouring growth sites.

Future research will include a genetic survey of the *ex situ* collection. This should give more insight in the genetic diversity, which will enable fine-tuning of *in situ* conservation measures.

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