Natural hybridization between cultivated poplars and their wild relatives: evidence and consequences for native poplar populations

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Abstract – It is recognized that introgressive hybridization and gene flow from domesticated species into their wild relatives can have a profound effect on the persistence and evolution of wild populations. Here, we review published literature and recent data concerning introgressive hybridization involving numerous species of the genus Populus. First, we briefly refer to some concepts and terminology before reviewing examples of natural and anthropogenic hybridization. Second, we examine whether natural genetic barriers could limit introgressive hybridization. Threat and possible consequences of anthropogenic hybridization are discussed in order to finally suggest conservation strategies for native poplar populations.

Résumé – Hybridations entre peupliers cultivés et sauvages : exemples et conséquences pour les populations naturelles. L’introgression et le flux de gènes d’espèces domestiquées vers les espèces sauvages peuvent avoir un rôle déterminant sur la dynamique des populations naturelles. Les objectifs de cette revue sont de faire le point de cette problématique chez les peupliers, à partir de données bibliographiques et de données récentes sur l’introgression et l’hybridation chez les espèces du genre Populus. Dans un premier temps, nous préciserons concepts et terminologie avant de présenter des exemples d’introgressions naturelles ou d’origine anthropogénique entre espèces autochtones et introdites. Nous décrirons ensuite les barrières reproductives qui peuvent limiter les flux de gènes entre espèces. Enfin, nous discuterons des possibles conséquences de l’hybridation et suggérerons des stratégies de conservation pour les populations naturelles menacées.

1. INTRODUCTION

The issues of natural hybridization and introgression (cf. Glossary) between cultivated plants and their wild relatives have been extensively studied in agricultural systems. However, introgression is of particular concern for forest trees because they can have large effects on ecosystem processes and biological diversity [26]. The genus Populus L. (poplars, Salicaceae) is considered as a model forest tree in plant biology [16, 93]. A large collaborative research community is dedicated to its study and this has resulted in a wealth of information on poplar biology and ecology, linking physiology, quantitative genetics (cf. Glossary) and genomics [16, 93]. In particular, Populus nigra L. (European black poplar) has been considered as a model tree in the study of gene resource conservation of wild relatives of cultivated plants [45, 55]. Much theoretical work on conservation of P. nigra has been done in Europe at national and international scales (EU-funded research project EUROPOR, cf. Glossary) [95] and a combined conservation strategy is applied at the European scale, supported by the Populus nigra EUFORGEN (cf. Glossary) network [55]. Despite the many excellent studies of recent years, many questions about hybridization in Populus remain unresolved. It is assumed that the presence of poplar artificial plantations pose a severe potential threat for the diversity and the regeneration of native indigenous poplars. For example, in Europe, the European black poplar is threatend by euramerican (P. × canadensis Moench.) and interamerican (P. × generosa Henry) hybrid poplar and P. nigra varieties such as the male Lombardy poplar (Populus nigra cv. ‘Italica’ Duroi) [3, 17, 45, 55]. Similarly, in North America, there is potential for extensive gene flow from plantations of cultivated hybrids between P. trichocarpa T. & G. and P. deltoides Marshall to native P. trichocarpa populations in the Pacific Northwest [25, 94]. Also, as hybrid poplar plantations may soon include genetically engineered trees, a major concern is that genetic resources of wild relatives will be altered through transfer of the transgene by hybridization [25, 31]. Moreover, the observed habitat fragmentation of some native poplar species due to human activities like agriculture and urbanisation of floodplain areas, increases the opportunities for...
contact with cultivated poplar plantations. This increasing con-
tact provides greater opportunities for the production of hybrid
seeds, which is produced at the expense of conspecific (cf. Glossary) seed [57]. Habitat reduction followed by hybridiza-
tion can lead to the extinction of a rare plant species [32], (for a
review see [57, 75]). As a consequence, some authors consider
native poplars as the most threatened forest tree species of old
natural floodplain forests in the temperate zones [55].

In sum, poplar offers valuable opportunities to address the
natural and anthropic effects of hybridization because of its
wide spread distribution in the Northern Hemisphere and the
numerous cultivars that have been intentionally introduced.

The aim of this review is to examine whether isolation mech-
anism (pre and post zygotic) are able to limit hybridization in
poplars, the extent and differences between natural and anthropo-
genic hybridization, and if current approaches to evaluate the
threat of hybridization are adequate or not. We discuss if cul-
tivated poplars pose a risk for the conservation of native poplar
populations with particular reference to the European black
poplar.

2. CONCEPTS AND TERMINOLOGY

The process of introgression occurs in many steps that
involve several hybrid generations (F1, F2, BC1, BC2 and so on)
(cf. Glossary). Thus, hybridization could lead to introgression if
the cultivated F1-hybrids backcross (cf. Glossary) to native
parental species. Concerning Populus, it is unlikely that natural
poplar hybrids are the result of frequent repeated backcrosses
due to the recent expansion of poplar cultivation world-wide.
In Europe, the intensive poplar cultivation started during the
1940s–1960s [23, 37] and natural hybrids are therefore assumed
to represent first generation-hybrids. Therefore we prefer to use
the term ‘hybridization’ rather than ‘introgression’.

The threat of a numerically superior species to rare cross-
compatible congeners most often springs from human activities
[57]. It is therefore important to differentiate between hybrid-
ization among native species in the absence of human activities,
and ‘exotic’ events brought about by human disturbances and
introductions (i.e. anthropogenic hybridization, cf. Glossary)
[2]. For example, major human disturbances have enabled the
European white poplar (Populus alba L.), native to Southern
Europe, to initiate naturally occurring hybrids on three contin-
te: the grey poplar (P. × canescens = P. alba × P. tremula)
in Europe, Rouleau’s poplar (P. × roullautiana = P. alba ×
P. grandidentata) in eastern North America, and the Chinese
white poplar (P. × tomentosa = P. alba × P. adenopoda) in
China [62]. In each area, the European white poplar hybridized
with a native aspen. In North America, North Europe and
China, the European white poplar itself was not native, but was
introduced as an ornamental tree and is extremely vigorous.

3. HISTORICAL CONTEXT OF POPULAR
CULTIVATION

The genus Populus is tremendously varied with 22 to about
85 species (depending on the interpretation of a ‘species’) dis-
tributed throughout the Northern Hemisphere, in both the tem-
perate and subtropical zones [30, 35]. Intercrossability among
species is one of the foundations of breeding work in the genus
Populus (e.g. [30, 86, 111]). Crossing programmes have
revealed both the wide extent of potential intercrossability and
the very real limitations on it [86, 111]. A list of natural and
artificial poplar hybrids is presented in Table I. The impressive
sylvicultural qualities of the cultivated hybrids (fast growth,
good form and easiness of vegetative propagation) have led to
widespread production of cultivated poplar plantations mainly
in Europe, North-America and China. The use of hybrid poplars
expanded from application in windbreaks to producing wood,
fibre and fuel products. In Europe, poplar species frequently
used in breeding programmes in order to produce hybrids are
the European black poplar (Populus nigra L.) and the North
American cottonwoods P. deltoides and P. trichocarpa. The
unit of cultivation and breeding in poplars is a clone, and indi-
cidual cultivars are normally represented by a single clone [73].
Poplar plantations usually represent one single cultivar or clone
in order to minimise the variability in growth and wood quality
within the plantation. They represent a very narrow genetic
base spread on a very wide scale and may contribute to a large
extent to pollen and seed pools. So, the main risk here relates

4. DISPERSAL AND ESTABLISHMENT

Populus species are predominantly dioecious and thus oblig-
atory outcrossers (cf. Glossary). Both sexes flower in early

spring prior to leaf initiation [15]. At maturity, the fruit capsules
split and release seeds embedded in significant quantities of
pappus (i.e. long white, silky hairs attached to the seed). In addi-
tion to being wind-pollinated and obligatory outcrossing, the
Table I. Some natural and artificial hybrids among species in the genus Populus. All hybrids except the ones indicated as artificial hybrids originating from controlled crosses, have spontaneously formed in areas where the natural range of species overlap or where exotic taxa have been planted near naturally growing poplar trees.

<table>
<thead>
<tr>
<th>Hybrid parents</th>
<th>Hybrid designation</th>
<th>Notes</th>
<th>Ref.</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>P. alba</em> L. × <em>P. adenopoda</em> Maxim.</td>
<td><em>P. × tomentosa</em> Carr.</td>
<td>May also be a tri-hybrid containing genes from <em>P. tremula</em></td>
<td>[24]</td>
</tr>
<tr>
<td><em>P. alba</em> L. × <em>P. trumula</em> L.</td>
<td><em>P. × canescens</em> Sm.</td>
<td>Gray poplar</td>
<td>[24]</td>
</tr>
<tr>
<td><em>P. angustifolia</em> James × <em>P. balsamifera</em> L.</td>
<td><em>P. × brayshawii</em> Boivin</td>
<td>Brayshaw’s poplar</td>
<td>[24]</td>
</tr>
<tr>
<td><em>P. angustifolia</em> James × <em>P. balsamifera</em> L. × <em>P. deltoides</em> March.</td>
<td>None</td>
<td>Trihybrid</td>
<td>[24]</td>
</tr>
<tr>
<td><em>P. angustifolia</em> James × <em>P. nigra</em> L.*</td>
<td>?</td>
<td></td>
<td>[24]</td>
</tr>
<tr>
<td><em>P. angustifolia</em> James × <em>P. trichocarpa</em> Torr. &amp; Gray.*</td>
<td>?</td>
<td></td>
<td>[24]</td>
</tr>
<tr>
<td><em>P. balsamifera</em> L. × <em>P. deltoides</em> Marsh.</td>
<td><em>P. × jackii</em> Sarg.</td>
<td>Jack’s hybrid poplar or heartleaf balsam poplar; also known as <em>P. balsamifera</em> var. subcordata or <em>P. candelicans</em></td>
<td>[24]</td>
</tr>
<tr>
<td><em>P. deltoides</em> × <em>P. maximowiczii</em>**</td>
<td>?</td>
<td>Artificial hybrid; Eridano is a clone of this hybrid</td>
<td>[34]</td>
</tr>
<tr>
<td><em>P. deltoides</em> × <em>P. yunnanensis</em>**</td>
<td>?</td>
<td>Artificial hybrid; Kawa is a clone of this hybrid</td>
<td>[34]</td>
</tr>
<tr>
<td><em>P. laurifolia</em> Ledeb. × <em>P. nigra</em> L.</td>
<td><em>P. × berolinensis</em> Dippel</td>
<td>Berlin or Russian poplar; syn. <em>P. × rasmusovskaya</em> Schneid. or <em>P. petrowskaya</em> Schneid.</td>
<td>[24]</td>
</tr>
<tr>
<td><em>P. maximowiczii</em> × <em>P. berolinensis</em>**</td>
<td>?</td>
<td>Artificial hybrid; Geneva, Oxford are clones of this hybrid</td>
<td>[34]</td>
</tr>
<tr>
<td><em>P. nigra</em> × <em>P. maximowiczii</em>**</td>
<td>?</td>
<td>Artificial hybrid; Rochester, Maxein, Maxwyo, Maxvier, Maxfünf are clones of this hybrid</td>
<td>[34]</td>
</tr>
<tr>
<td>(<em>P. deltoides</em> Marsh. × <em>P. nigra</em> L.) × <em>P. balsamifera</em> L.</td>
<td><em>P. × rollandii</em> Dode</td>
<td>Trihybrid; very similar to <em>P. × jackii</em></td>
<td>[24]</td>
</tr>
</tbody>
</table>

Pappus of the seeds promotes wind dispersal over great distances resulting in high rates of migration and high gene flow (cf. Glossary) and genetic diversity [39, 56]. Poplars are prolific seed producers. Old trees can produce over 50 million of seeds in a single season [62]. Seedlings colonise moist, recently exposed soil (due to previous flooding in winter) along gravel bars, sandbars and riverbanks within riparian corridor. Poplar seeds are small and rapidly lose viability [15]. Continuous moisture is required for seed germination. If river levels decline too rapidly, seedlings succumb to drought stress. Seedlings that establish on moist soils at lower river levels are subject to later removal or damage by the scouring of floodwaters [15]. Collectively, these environmental constraints contribute to the infrequent establishment of seedlings, on the order of every 10 to 20 years depending on climatic conditions and channel morphology [15]. The age of reproductive maturity varies among native species from five to ten years, yet in some natural populations may not occur until the trees are 15 to 20 years [15, 85], limiting introgression events. In contrast, hybrid poplars grown in well-maintained plantations commonly attain reproductive maturity in four years [94]. Poplar species are also capable of asexual or vegetative reproduction, as an alternative to regeneration from seed. Asexual reproduction is promoted only by flood disturbances when through extended periods of submergence and/or mechanical damage to parent plants, dormant primordia (cf. Glossary) in
roots and shoots are stimulated to produce new shoots and roots [8, 9]. Thus, genotypes can persist on sites for long time periods beyond the longevity of single trees [89]. There is also evidence of cladoptosis (cf. Glossary), in which short shoots abscise and can be carried long distances on watercourses and subsequently take root [38].

5. REPRODUCTIVE ISOLATION MECHANISMS

Several authors have reported on mechanisms that limit (in whole or in part) gene flow between the hybrids and the parental species. Reproductive isolating mechanisms are generally divided into two categories based on whether they act before or after fertilization (Fig. 2). Mechanisms that act to prevent mating are referred to as prezygotic, whereas those that act to reduce the viability or fertility of the hybrid zygote or later-generation hybrid offspring are referred to as postzygotic [77].

5.1. Prezygotic barriers

Pollen competition (Figs. 3 and 4) seems to be an important process in limiting the production of hybrid progeny in Populus. Rajora [63] observed competition among pollen of P. deltoides, P. nigra and P. maximowiczii A. Henry in fertilizing P. deltoides ovules. The low frequencies of interspecific matings relative to conspecific matings in pollen-mix controlled crosses suggest conspecific pollen advantage [63, 99]. As a result, the relative fertilisation success of a pollen species depends upon the species constitution of the pollen mix. Similar results were obtained by Benetka et al. [11] and Vanden Broeck et al. [99] with crossing experiments of P. nigra females: when P. × canadensis pollen was used in mixtures with P. nigra pollen, most of the seedlings were fathered by P. nigra. Pollen competition was also studied in detail for Louisiana irises (Iris fulva and Iris hexagona). These studies have focussed on conspecific pollen advantage, including measurements of pollen tube lengths in conspecific and heterospecific styles and examination of hybrid seed production following different types of controlled pollinations (for a review see [77]). In common with introgression studies in irises [77] studies on gametophytic competition in Zea mays [60] and Dianthus chinensis [59] also suggest that faster growth of conspecific pollen tubes might be acting as a barrier to interspecific hybridization (cf. Glossary) [59, 80]. Prefertilisation incompatibility (cf. Glossary) barriers can occur on the stigmatic surface or during the growth of the pollen tube in the style, in the ovary, in the ovules or in the embryo sac [50]. Prefertilisation incompatibility barriers in Populus sp. are known to be located primarily in the stylar tissues [102]. Special emphasis has been made on pollen tube-pistil interaction in P. nigra, revealing precise growth of pollen tube within the ovary [101], role of ABA in flower pedicel abscission due to zygotic interspecific incompatibility [52] and role of β-galactosidase activity associated with conspecific pollen tube growth [103]. Interspecific mixtures of viable incompatible and killed compatible (mentor/recognition) pollen have been used to achieve incompatible mating with variable success (reviewed in [102]).
5.2. Postzygotic barriers

Common postmating reproductive barriers include hybrid sterility, hybrid weakness or inviability, and hybrid breakdown (cf. Glossary) in which first generations (F1) hybrids are vigorous (hybrid superiority due to heterosis), robust and fertile, but later generation hybrids are weak or inviable [77]. Poplar hybrids are characterised by reduced fertility relative to parental species, and significant lower pollen production and seed viability in F1 hybrids [86, 94]. This has been observed in poplar hybrids growing in commercial plantations and natural zones of hybridization [94]. However, recent analyses have found that poplar hybrids are not uniformly unfit, but rather are genotypic classes that possess lower, equivalent or higher levels of fitness relative to their parental taxa. Schweitzer et al. [81] found that F-generations of *P. fremontii* S. Watson × *P. angustifolia* James and backcross generations can be just as fit as the parent taxa. F-generations produced as many viable seed as *P. angustifolia* and backcross genotypes produced as many viable seeds as both parent taxa. Moreover, hybrids produced nearly two and four times as many ramets from root sprouts as *P. angustifolia* and backcross genotypes produced as many viable seeds as both parent taxa. Therefore, extreme low fertility or viability of early-generation hybrids (e.g. F1, F2, BC1) does not necessarily prevent extensive gene flow [5]. Detailed studies of the genetic basis of hybrid breakdown suggest that hybrid weakness appears to result from the break-up of co-adapted gene complexes (cf. Glossary) that affect fitness traits [6, 58, 75, 77]. If many genes contribute to fitness traits, then much of the genome may be resistant to introgression because of linkage (cf. Glossary). These results also suggest that species genomes are often differentially permeable to introgression, where certain portions of the genome are open to the incorporation of alien alleles, but introgression is restricted in other parts of the genome [58, 76]. Introgression may therefore be an important mechanism for the transmission of ecologically functional traits [6]. Different selection pressures for different genomic regions were also observed in *Populus* [6, 58]. The results of Martinsen et al. [58]

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**Figure 2.** Reproductive isolation mechanisms in *Populus*.

**Figure 3.** Pollen tube competition within the stigma of a *Populus nigra* flower. After germination of the numerous pollen grains on the stigmatic surface, the pollen tubes grow within the stigmatic tissues and compete to reach the stylodium (bottle neck) before reaching the ovarian cavity.

**Figure 4.** Diagrammatic representation of pollen tube competition on a *Populus nigra* flower. po: pollen grain, pt: pollen tube, st.l: stigmatic lobes, st: stylodium, ov: ovule. Adapted from [103].
indicate that the hybrid zone between natural populations of *P. fremontii* and *P. angustifolia* (*P. × hinckleyana* = natural hybrid between *P. fremontii* × *P. angustifolia*) located near the river Weber (Utah, USA), acts as an evolutionary filter, preventing the introgression of most genes but allowing others to introgress. Different introgression rates (localized introgression, slow-dispersed and fast-dispersed introgression) among genetic markers suggest that there are different selection pressures for different genomic regions. A few well dispersed, negative acting genes could impede the majority of the genome from introgression [58]. Because large DNA segments are more likely to carry such negative genes, backcrosses carrying large DNA segments, would be selected against. Consistent with this hypothesis, selective filtering of the genome appears to be most intense in the very earliest backcross generations (e.g. at the hybrid zone boundary) where the average introgressing segment is likely to be 25 cM or larger [58]. The arrangement of genes and their action will have a major impact on the genomic pattern and rates of introgression [58]. If hybrid populations act as evolutionary filters, there are several important implications [58]. First, species barriers are maintained in the face of hybridization. Second, a strong filter should prevent the introgression of deleterious genes while allowing introgression of beneficial ones. Finally, a filter help explain the existence and long-term persistence of hybrid zones [58].

### 6. DETECTION OF HYBRIDIZATION

Based on morphological traits, it is not always possible to unambiguously detect introgressed genes in the offspring of a particular species. Individuals from hybrid swarms (cf. Glossary) that obtained most of their genes from one of the parental taxa are often morphologically indistinguishable from that parental taxon [2]. Furthermore, environmental influences on morphology and differences between juvenile and mature characters make it difficult to determine accurately the taxon of young poplar seedlings on the basis of morphological characters alone [2]. Molecular markers now provide helpful tools to assess introgression in plant populations. Over the years, the most conclusive evidence for introgression in *Populus* sp. has come from molecular data (Tab. II): initially from isozyme (cf. Glossary) work [10, 47, 48, 64, 67] and more recently also from DNA data (e.g. [20, 41, 42, 45, 65, 88, 100, 104]). Finding true diagnostic alleles for each species require extensive sampling from both species’ natural ranges, as alleles common in one species may be rare, but still present in the other [45]. Chloroplast and mitochondrial cytoplasmic markers (Tab. II) are maternal inherited in *Populus* [68] and are a useful complement to nuclear markers to study the direction of introgression [20]. F₁-hybrids can be reliably identified with nuclear markers but it is more difficult to distinguish between F₂, backcrosses and later-generation hybrids [13]. However, in *Populus*, interspecific natural hybridization among native species is geographically restricted and parental populations still exist [27, 29, 36, 94]. In contrast with interspecific natural hybridization, anthropogenic hybridization and introgression might have begun only recently so that to date, advanced generations of backcrosses among domesticated and native *Populus* taxa are unlikely. However, an enlarged set of nuclear and/or biochemical markers is desirable to estimate parent’ species contributions to single clones and plants more accurately [45].

Several authors agree about the usefulness of the diagnostic isozyme systems LAP, PGM, PGI to differentiate between *P. deltoides*, *P. nigra* and *P. × canadensis* [10, 11, 67, 74, 100]. The enzyme loci Aco-3 and Mdh-3 were found diagnostic for differentiating between *P. alba*, *P. tremula* and their hybrids (*P. × canescens*) [69]. Species-specific markers in *Populus* have also been described based for ribosomal DNA [22, 33], mitochondrial DNA [7, 66], chloroplast DNA (e.g. [20, 45, 51, 70, 83, 104]) and the STS nuclear gene marker win3 [41]. Recently, a microsatellite marker (WPMS09) [96] was found useful to differentiate between genes of *P. deltoides* and *P. nigra* [37, 100] (cf. Glossary).

Hybridization of native *P. nigra* with the cultivated *P. nigra* cv. ‘Italica’ and *P. nigra* cv. ‘Thevestina’ is more difficult to detect as they belong to the same species. Imbert & Lefèvre [46] found for each of 6 SSR loci studied, alleles in natural populations of *P. nigra* along the river Drôme that were shared with the Lombardy poplar, although the frequency of these alleles was highly variable. They considered the most frequent alleles as common alleles of the species whereas the rarest alleles were considered to provide an upper estimate of possible

### Table II. Some molecular markers used in studies of introgressive hybridisation in *Populus*.

<table>
<thead>
<tr>
<th>Marker type</th>
<th>Mode of transmission</th>
<th>Mode of gene action</th>
<th>Level of variability</th>
<th>Ref.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Isozymes</td>
<td>Biparental</td>
<td>Codominance</td>
<td>Low to medium</td>
<td>[10, 11, 47, 48, 64, 65, 67, 74, 88, 98]</td>
</tr>
<tr>
<td>Chloroplast microsatellites (cpDNA)</td>
<td>Maternal Nuclear</td>
<td>Codominance</td>
<td>Medium to high</td>
<td>[21, 22, 33]</td>
</tr>
<tr>
<td>Restriction Fragment Length Polymorphisms (RFLPs) of ribosomal RNA genes</td>
<td>Biparental Nuclear</td>
<td>Codominance</td>
<td>Medium to high</td>
<td>[21, 22, 33]</td>
</tr>
<tr>
<td>Nuclear microsatellites (SSR)</td>
<td>Biparental Nuclear</td>
<td>Codominance</td>
<td>High</td>
<td>[37, 92, 97]</td>
</tr>
<tr>
<td>Amplified Fragment Length Polymorphism (AFLP)</td>
<td>Biparental Nuclear</td>
<td>Dominance</td>
<td>High</td>
<td>[3, 18, 97, 108]</td>
</tr>
<tr>
<td>Sequence Tagged Site marker win3 (STS)</td>
<td>Biparental Nuclear</td>
<td>Codominance</td>
<td>Medium</td>
<td>[41, 92, 98]</td>
</tr>
</tbody>
</table>
Hybridization with the Lombardy poplar [46]. The SSR locus WPMS09 [96] showed the lowest frequency of the two alleles shared with *P. nigra* cv. 'Italica' (< 0.05) [46].

### 7. NATURAL HYBRIDIZATION AMONG NATIVE POPLAR SPECIES

Despite the numerous barriers, hybridization is common in *Populus*. Natural poplar hybrids are regularly found wherever different species come into contact with one another [28, 29, 36, 58]. However, the extent of introgression can vary substantially depending on the species involved, the environment of the hybrid zone and the portion of the genome examined [58]. Hybridization between species of different sections is comparatively restricted. Only section Aigeiros and section Tacamahaca are freely intercrossable [28, 29]. These two sections, like others in the genus, differ somewhat in habitat as well as in range. Balsam poplars (Tacamahaca) are trees of boreal and montane habitats, while cottonwoods (Aigeiros) occupy lower elevations of middle latitudes. They are largely allopatric in Eurasia but are broadly sympatric in North America, with overlapping ecological preferences. They contact each other and hybridise primary where topographic diversity brings their distinctive habitats into proximity. Such localities, although widespread, are restricted (about 10–15 km) and populations containing hybrids represent a small fraction of all populations of each parent species. Most hybrids in the field appear to be first-generation crosses [28] although advanced-generation hybrids were also observed [58, 94]. The hybrids show little tendency to invade the characteristic habitats of either of their parents but tend to remain in the hybridised habitat [28, 29]. Natural hybrid zones have existed between sympatric native poplar species for millennia, yet these species have retained their identities, with remarkably little introgression [28, 58]. Rhymer and Simberloff [75] discuss that lack of fitness of F1 hybrids, later generations, or backcrosses are evident in many stable hybrid zones. This is a form of outbreeding depression (cf. Glossary) and may result from chromosomal differences, the breakdown of co-adapted gene complexes or both. Hybrid zones also form at boundaries of narrow gradients between distinct habitats, so that each habitat favours one parental taxon, and hybrids are selected against in both [40]. There are exceptions to this ecological barrier to hybridization; in local areas the habitats of cottonwood and balsam poplar parents may be virtually indistinguishable and thus not limiting to the hybrids. This is probably the case with *P. balsamifera* L., *P. deltoides*, and *P. × jackii* Sarg. (natural hybrid of *P. balsamifera* and *P. deltoides*), in part of their broad range of sympathy [78]. In North America near the Weber River in Utah, *P. fremontii* hybridises with *P. angustifolia* where their distinctive habitats come into proximity. In the 13-km hybrid zone introgression is unidirectional; F1 hybrids only backcross with pure *P. angustifolia* (absence of hybrid × hybrid and hybrid × *P. fremontii* crosses) [49]. This restricted hybridization process means that the hybrid zone is not self-perpetuating, and will presumably go extinct in the absence of a *P. angustifolia* population [49].

Whitham et al. [107] and Rotach [79] illustrated that hybrid poplar zones are dynamic centres of ecological and evolution-ary processes for plants and their associated communities. The intermediate genetic differences between the parental species will result in the greatest genetic variation in the hybrid zone, which in turn will have a positive effect on biodiversity [79, 107]. These natural zones of hybridization are unique and worthy of special efforts to promote their conservation and protection [58, 107]. This is particular the case in China, where ten different hybrid swarms have been registered so far [106].

### 8. ANTHROPOGENIC HYBRIDIZATION

Numerous exotic poplar species have been introduced into Europe as well as into the USA, Canada and China for the establishments of shelterbelts, windbreaks and wood production in urban, suburban and agricultural landscapes. As no published information exists concerning anthropogenic hybridization of *Populus* in China, we had to focus on examples from Europe and North America.

#### 8.1. North America

Examples of non-native poplars introduced in North America are *P. nigra*, *P. alba*, *P. × canadensis* and hybrids between *P. deltoides* and *P. trichocarpa* (i.e. *P. × generosa*) [25, 94, 110] and *P. × petrowskyana* (a hybrid of *P. laurifolia* Lede. × *P. deltoides*) [78]. *P. alba* L., the European white poplar, was introduced from Europe to North America in the early 18th century, and hybridises naturally with the native American aspens, *Populus grandidentata* Michaux and *P. tremuloides* Michaux. [84]. However, only F1 hybrids resulting from natural hybridization between introduced *P. alba* and native *P. grandidentata* or *P. tremuloides* were found and no advanced generation hybrids or backcrosses were detected [84].

A female clone of *P. × petrowskyana* (often referred to as “Russian” poplar) was introduced and widely distributed in western Canada about 100 years ago [78]. This clone hybridises occasionally with the native species *P. balsamifera* and more frequently with *P. deltoides* cv. *occidentalis* and *P. × jackii* [78]. Eckenwalder [27] reports on a single locality in California at which native *P. fremontii* has apparently hybridised successfully with a female introduced *P. nigra*. The Lombardy poplar is widely planted in North America. It is quite fertile and capable of siring seeds with *P. trichocarpa* females even when the trees are separated by large distances [25]. However, despite a long history of extensive cultivation in proximity to native populations, there is no evidence for extensive introgression of Lombardy poplar in native populations in North-America [25, 94]. This paucity of hybrids may be attributed to the sex of the introduced *P. nigra*. Its pollen may not be able to compete with pollen of *P. deltoides* or *P. fremontii* in fertilizing ovules of these species in natural populations (Baker, 1951, in [94]).

Occasionally, in North America non-native poplars colonise adjacent riparian corridors via asexual propagation, yet there are no reports of these species displacing native cottonwood populations [94]. Because of the limited size of plantations compared to wild stands in most area, plantation-derived
propagules are usually greatly diluted with propagules from wild stands, including those located a short distance from plantations [25]. Other barriers like conspecific mating preference may also play a role in limiting anthropogenic hybridization [99].

8.2. Europe

The most common non-native poplars in Europe include: \( P. \times \) canadensis, \( P. \times \) generosa (cultivated hybrids between \( P. \) deltoides Marshall and \( P. \) trichocarpa), \( P. \) trichocarpa, \( P. \) deltoides, \( P. \) alba (non-native in North Europe), \( P. \) nigra cv. ‘Italica’ (Lombardy poplar) and \( P. \) nigra cv. ‘Thevestina’.

In Europe in the eighteenth century, spontaneous hybridization between the introduced \( P. \) deltoides and the native \( P. \) nigra occurred, giving rise to the widely cultivated \( P. \times \) canadensis [27]. While the activities on poplar breeding and cultivation rise in Europe, human activities in floodplain areas including agriculture, urbanisation and hydraulic engineering resulted in the habitat destruction and modification of the native black poplar. Moreover, clones of \( P. \times \) canadensis were massively planted for wood production to replace the autochthonous black poplar resources on alluvial floodplains in many countries of the temperate region. This resulted in a severe reduction in population size and habitat fragmentation of the European black poplar all over Europe. The situation in the United Kingdom and in Belgium illustrates the risk of species extinction in the margins of its range area. In these countries, black poplar is considered to be one of the rarest native trees [19, 100]. However, only recently evidence for introgressive hybridization of \( P. \times \) canadensis in the offspring of black poplar females was found [98]. In Belgium, genes of \( P. \) deltoides were detected in the open pollinated offspring of a black poplar female, surrounded by \( P. \times \) canadensis hybrids and in the absence of conspecific males. This is the only study reporting on introgression in the offspring of black poplar and the results are in contrast with other studies, were no introgression of \( P. \) deltoides genes in the offspring of \( P. \) nigra females was found, even when flowering male trees of \( P. \times \) canadensis were present in the vicinity [10, 41, 92]. The Belgian study [98] differs from the other studies by the fact that the female black poplars investigated were in the absence of conspecific males. In a mixed pollen cloud, pollen of \( P. \) nigra may be more successful than that from \( P. \times \) euramericana in pollinating female black poplar [92, 100]. This hypothesis could be confirmed by controlled crossing experiments with interspecific pollen mixes (i.e. a mix of \( P. \) nigra and \( P. \times \) canadensis pollen) suggesting conspecific pollen advantage [99]. However, if no pollen of the own species is present, \( P. \) nigra females can be pollinated successfully by pollen of \( P. \times \) canadensis. Therefore, low levels of introgression from \( P. \times \) canadensis are expected in natural populations of \( P. \) nigra where there are several male black poplars close to the female trees. This is confirmed by the results of the European research project EUROPOP were low levels of introgression was found in black poplar populations along the borders of six river systems (Danube, Drôme, Ebro, Rhine, Ticino and Usk) [54]. Also in France low levels of introgression of genes of \( P. \) deltoides in natural black poplar populations were found along the rivers Garonne [61] and Loire (Villar, pers. observations), and no alleles of \( P. \) deltoides and \( P. \) trichocarpa were detected in natural black poplar populations along the Drôme [46]. In contrast, genes of \( P. \) deltoides were detected in young poplar seedlings colonizing the banks of the river Meuse in Belgium [99], the rivers Waal and Meuse in the Netherlands [12], the river Ebro in Spain [1] the river Danube in Austria [44, 51] and Hungary [14] and in natural black poplar populations located in the Czech Republic [10]. It is most likely that these introgressed seedlings originate from hybrid × hybrid crosses or from open pollinated \( P. \times \) canadensis females [1, 11, 12, 14, 43]. These findings indicate that, although low levels of introgression are detected in natural black poplar populations, cultivated poplars are reproductive along several river systems in Europe and that they may compete with the native species in colonising new habitats. Some of the introgressed seedlings seemed to be well adapted as they survived the river dynamics over several years.

9. CONSERVATION IMPLICATIONS

Cases of hybridization appear to be relatively frequent in plants, and many taxa have been influenced by hybridization at some point in their history [105]. The numerous barriers to hybridization including hybrid breakdown, reduced hybrid fertility, and post-zygotic reproduction barriers are paradoxical in light of the widespread occurrence of natural hybrids among native poplar species. This is due to the fact that successful establishment of hybrid populations is determined by events that, in many organisms, occur only rarely [4]. This establishment may initially be nonadaptive, or even maladaptive for the hybridising pairs, but may lead to adaptive evolution (cf. Glossary) through the production of hybrid genotypes that are more fit than their parents in the parental or novel habitats [4]. Therefore, hybridization with or without introgression, even when occurring rarely, can have important consequences for evolution and conservation biology of native poplar populations. This is particularly true for anthropogenic hybridization and for rare species that come into contact with other species that are more abundant [75]. Rhymer and Simberloff [75] have concluded that the severity of this problem has been underestimated by conservation biologists. The increasing pace of the three interacting human activities that contribute most to increased rates of hybridization (introductions, fragmentation and habitat modification) suggests that this problem will become even more serious [75]. This is particularly the case for Populus, a forest tree of considerable economic importance [93]. Several studies have demonstrated that domesticated poplars are capable of spontaneously mating with their wild relatives under field conditions and that hybridization can vary considerably with the specific populations involved [10, 44, 46, 61].

When wild poplar stands are very small compared to widespread hybrid poplar plantations, the wild populations may go extinct through genetic assimilation. Individuals in small, isolated populations in contact with other taxa are much more likely to hybridise if only because of the difficulty of finding mates of the same species. This is the case for isolated native populations of \( P. \) nigra at the Northern margin of its geographical distribution area [92, 100] and at its Eastern margin in China where \( P. \) nigra freely hybridise with \( P. \) laurifolia (Villar, pers. observations, [82]). There, introduced hybrid poplars may
be able to outperform native poplar populations in the short term (a few generations). Once hybridization has begun, it is difficult to stop, especially if hybrids are fertile and mate both among themselves and with parental individuals [2], as seemed to be the case in *Populus* [99]. Eliminating the domesticated poplars is not an option because of the economic importance of cultivated hybrid poplar plantations. In this case, effort should focus on maintaining and expanding the remaining pure native populations [2]. Probably, the best long-term solution here is habitat restoration in combination with reforestation programmes.

When natural populations are large compared to cultivated poplar plantations, as is the case in France along the studied rivers Loire, Allier and Drôme anthropogenic introgression seems to be restricted ([46], Villar, unpublished data). However, in this case, monitoring of the process of hybridization is of crucial importance in order to avoid that the proportion of hybrid individuals increases progressively. The competitive ability and initial frequency are factors that have a strong effect on risk of extinction through hybridization [109]. As for most pioneer species, competition for resources becomes important soon after germination and thus selection might occur at a very early stage [46]. Population size, effective population size and potential introgression should be monitored across multiple generations in order to estimate extinction risk. They should be considered in management and conservation plans.

10. CONCLUSION AND FUTURE RESEARCH NEEDS

There is still much work to be done before we fully understand the risks that are associated with gene flow between cultivated poplars and their wild relatives. We know that anthropogenic introgressive hybridisation occurs albeit it is limited by natural genetic barriers and its frequency depends on the local population size and the abundance of the cultivated poplars. In summary, the risk of natural hybridization between cultivated poplars and their wild relatives must be analysed on a case-by-case basis. Situations where cultivated poplars outnumber very small native poplar populations are likely to be most problematic. The issue of introgression in *Populus* will continue to receive much attention as it is also important for assessing the potential impacts of transgenic poplar plantations [90, 91]. More diagnostic, species specific molecular markers and knowledge on their genomic distribution are needed to allow a more accurate and rapid assessment method to quantify introgression. To help understand the consequences of introgression, studies on reproductive properties like fertility, colonisation capacity and competitive ability of cultivated poplars compared to their wild relatives are needed. When hybrid seedlings cannot effectively compete in the field with seedlings of the native species, introgression will not occur. Also, a better understanding of the genomic distribution and function of genes that contribute to fitness traits and the detection of regions in the genome that are less prone to introgression would allow a better understanding of introgression and subsequent consequences for the native poplar populations [87]. The data and skills accumulated regarding these questions may also prove helpful in addressing the issue of transgene flow in *Populus*.

Glossary*

Adaptive evolution: the process of change in a population driven by variation in reproductive success that is correlated with heritable variation in a trait.

AFLP: abbreviation for amplified fragment length polymorphism. A type of DNA marker, generated by the PCR amplification of restriction endonuclease treated DNA. A small proportion of all restriction fragments is amplified in any one reaction, so that AFLP profiles can be analysed by gel electrophoresis. This has the important characteristic that many markers can be generated with relatively little effort.

Anthropogenic hybridization: the movement of genes between species or taxa caused or produced by human action.

Backcross (BC1, BC2): the offspring of a cross between a hybrid and one of its parents. The subscript number represents the number of generations that have been crossed in this fashion.

Chloroplast DNA (cpDNA): the DNA present in the chloroplast.

Cladopsis: the shedding of twigs by abscission.

Co-adapted gene complexes: particular combinations of genes at multiple loci that interact to confer higher fitness relative to other genotypes.

Conspecific: belonging to the same species.

Dioecious: organisms in which the male and female sex are in separate individuals.

EUFORGEN: European Forest Genetic Resources Programme.

EUROPOP: Research project “Genetic Diversity in River Populations of European Black Poplar for evaluation of biodiversity, conservation strategies, nature development and genetic improvement”. This project was carried out with financial support from the Commission of the European Communities, Agriculture and Fisheries (FAIR) specific RTD programme, PL97-3386

F1 (2): the first (second) generation of a cross.

Gene flow: exchange from genes among populations because of successful reproduction by migrants.

Genetic swamping: rapid increase in frequency of an introduced genotype (or introduced allele) that might lead to replacement of local genotypes; caused by a numerical and/or fitness advantage.

Hybrid breakdown: reduction in fitness of hybrids relative to parents caused by disruption of co-adapted gene complexes via recombination.

Hybrid swarm: a population consisting of hybrids and various types of backcrosses between two or more intercrossing sympatric species.

Hybridization: the process of forming a hybrid by cross pollination.

Incompatibility: selectively-restricted mating competence, which limits fertilization, such as lack of proper functions by otherwise normal pollen grains or certain pistils, a condition that may be caused by a variety of factors.

Interspecific hybridization: the movement of genes between species.

Introgression: the infiltration of germplasm from one species into another through repeated backcrossing of the hybrids to the parental species.

Isozyme: a genetic variant of an enzyme. Isozymes for a given enzyme share the same function, but may differ in level of activity, as a result of minor differences in their amino acid sequence. LAP, PGM, PGI: abbreviations for the isozymes leucine amino peptidase, phosphoglucomutase and phosphogluco isomerase, respectively.

Linkage: the greater association in inheritance of two or more nonallelic genes than is to be expected from independent assortment.

Microsatellite or Simple Sequence Repeat (SSR): a segment of DNA characterized by a variable number of copies (typically 5–50) of a sequence of around 5 or fewer bases.

Mitochondrial DNA: a circular DNA found in mitochondria.

Outbreeding depression: reduction in mean population fitness resulting from hybridization between genetically distinct individuals or populations of the same species; detected in F1 or subsequent generations.

Outcrosser: an individual that must be cross-pollinated to successfully complete generative reproduction.

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