

Kin selection, kin recognition and kin discrimination in plants revisited: A claim for considering environmental and genetic variability

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1	Kin selection, kin recognition and kin discrimination in plants revisited: a
2	claim for considering environmental and genetic variability
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21 Abstract

Considering kin selection in the study of interacting conspecific plants broadens our vision of plant behaviour and brings arguments to explain plant-plant positive interactions. These interactions are the subject of abundant research in community ecology and the role of relatedness in interactions between individuals has become a hot topic. Indeed, the past decade has seen a steady accumulation of exciting but also controversial results regarding the behaviour of plants in the presence of genetically related neighbours, i.e., kin discrimination. In addition to the existence of some methodological and semantic aspects that should be strengthened to accurately reveal the occurrence of kin recognition or selection, we claim here that some key aspects must be considered to increase our ability to detect kin recognition and selection in plants. These aspects include intraspecific variability, the fact that plant-plant interactions are context-dependent, and the lifespan of species (particularly long-lived ones). Additionally, we note that the study of the population spatial genetic structure (SGS) could be a useful tool to identify candidate settings for the study of kin recognition and selection in plants.

36 **Introduction**

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Plants are sessile organisms and cannot chose or move away from their neighbours, but are tied to them throughout their life. In this context, identifying the factors that can influence the type of interactions (positive, neutral or negative) between individuals is of prime importance. One such factor that has become increasingly popular is genetic relatedness between individuals. Recent research has shown compelling evidence that plants are able to detect their close neighbours and actively interact with them. Indeed, plants exhibit complex recognition and communication mechanisms (Bilas, Bretman & Bennett 2021). Classically studied between different plant species, these interactions also apply at the intraspecific level (Bilas, Bretman & Bennett 2021). Remarkably, plants can distinguish their conspecific neighbours based on their genetic relatedness (i.e., kin recognition) and adjust their 'behaviour' accordingly (kin discrimination) (Ehler & Bilde 2019, Anten & Chen, 2021a). Being able to identify related individuals could be advantageous to reduce competition among relatives (File, Murphy & Dudley 2012), a cooperative behaviour that can be promoted through kin selection. However, traits that reduce competition can be costly for the individual (e.g., limiting resource acquisition; Dudley, Murphy & File, 2013). Kin selection theory (Hamilton 1964) states that an altruistic trait can be favoured by natural selection when the altruist and the recipient are related. This is based on the concept of indirect fitness, i.e., indirect benefits accrued by the altruistic individual through the fitness of the recipient, as the two individuals share genes. If individuals are sufficiently closely related, altruism can favour the reproduction of the set of genes they share despite losses in direct fitness by the altruist individual (Axelrod & Hamilton 1981). Note that kin recognition and kin discrimination are not necessary for kin selection to occur (e.g., in viscous populations, cooperation with neighbours is mostly directed towards kin; West et al., 2007).

In a recent review, Anten & Chen (2021a) provide a critical and detailed overview of the conceptual bases and latest results regarding kin recognition, kin discrimination and kin selection in plants. In summary, they state that most of the studies have so far explored plant-plant interactions at the root level. A frequently observed result is that plants growing with related individuals show a reduction in belowground biomass compared to plants growing with strangers (usually seeds from different mothers), which is interpreted as a reduction in competition towards relatives, possibly leading to positive interactions (File, Murphy & Dudley 2012). The possible mechanisms of belowground kin recognition are root-exudates and mycorrhizal networks. Although some methodological concerns are discussed by Anten &

Chen (2021a), the authors conclude that there is sufficient evidence for supporting kin recognition in plants.

Anten & Chen (2021a) accomplished the best update so far regarding kin discrimination and kin selection in plants. We think, nevertheless, that some aspects were not considered in their review that are of great complementary importance to broaden the kin recognition and selection framework: intraspecific variability, environmental context dependency, plant longevity and spatial patterns of plant interactions. We expand on these underexplored aspects, which we consider essential to build a testable framework about kin discrimination and kin selection and expose two case studies that exemplify these new aspects. We then develop two additional points:(i) several papers confuse (or use alternatively) the concepts of kin recognition, kin discrimination and kin selection for which we offer a clarification; (ii) this confusion often leads to methodological issues and we propose methodological guidelines for experimental designs to detect kin discrimination and assess kin selection.

Should we all cooperate? The neglected role of genetic diversity for kin interaction mechanisms

Several published studies, including Anten & Chen's (2021a) review, have concluded that kin recognition is either present or absent in a given species. However, do we expect species to be entirely homogeneous with regard to kin discrimination or even kin selection? In other words, is it reasonable to assume that all individuals of the same species exhibit the same 'behaviour'? Probably not, as already suggested by several authors (Clark 2010; Bolnick et al., 2011; Westerband, Funk & Barton 2021). For example, it is recognised that intraspecific diversity in plants shapes biotic and abiotic interactions and plays a key role in determining the performance of individuals (Westerband, Funk & Barton 2021). Intraspecific diversity is the rule for a wide range of traits (Siefert et al., 2015), it is therefore conceivable that not all individuals within a species possess the same traits associated with kin recognition, and that not all individuals have the (same) ability to distinguish their related neighbours. Moreover, our understanding of the genetic and molecular bases underlying the natural variation of plant-plant interactions remains limited and needs to be considered in further research (Subrahmaniam et al., 2018). We assert here that intraspecific variability for kin recognition has been poorly addressed thus far, although it represents a critical issue.

Theoretical models exploring kin recognition in various organisms (mainly on animals, but the results are valid for all species capable of kin recognition) predict a wide range of intraspecific variation in kin recognition mechanisms between and within populations (see review by Penn & Frommen 2010), although other models do not predict kin recognition mechanisms at equilibrium because these mechanisms are costly (Rousset & Roze 2007). We argue that kin recognition in plants could still be an advantageous local strategy in certain conditions while not being advantageous when the population reaches equilibrium. Thus, intraspecific diversity for kin interaction mechanisms should be particularly important to consider. Anten & Chen (2021a) and also Fréville et al. (2021) discussed this point in the case of crops, which exhibit lower genetic diversity than wild species and where individuals interact mostly with kin and proposed that studies include sampling replication at the genotypic level to capture sufficient intraspecific variation in kin discrimination. Most studies have used only a few different kin groups (usually two to four) and thus captured only a small part of the existing variability in the species. Yet, some recent studies have made the effort to include a larger number of kin groups (eight-ten and up to 28 kin groups) (Semchenko, Saar & Lepik 2017; Torices, Gómez & Pannell 2018; Takigahira & Yamawo 2019). To overcome this important issue, which can lead to erroneous conclusions at the species level, we advocate the use of at least five to ten kin groups originating from several populations. The logic is as follows: the probability of detection depends on the frequency of the trait. If one in five individuals in a population can discriminate kin, using five kin groups there is still a 33% chance of missing the trait (i.e., reaching erroneous conclusions), and this value is reduced to 11% with ten kin groups. If the frequency is one in ten, the chances are 60% and 35%, respectively.

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Under which conditions is it beneficial to cooperate with relatives?

Because plants are sessile organisms, they cannot move to avoid their neighbours' effects nor environmental stress. Local adaptation (i.e., selection for traits that increase the fitness of individuals in their local environment) and phenotypic plasticity (i.e., the ability of a given genotype to express different phenotypes according to some environmental cues or circumstances) are the two common processes to overcome these threats (e.g., de Villemereuil, Mouterde, Gaggiotti & Till-Bottraud 2018; Fox, Donelson, Schunter, Ravasi & Gaitán-Espitia 2019). Phenotypic variability can thus be induced by variation in the local conditions and could influence the nature of plant-plant interactions (Callaway, Pennings & Richards 2003). As a consequence, should we expect the same type of interactions between individuals living under

different conditions? The stress gradient hypothesis (SGH, Bertness & Callaway 1994) predicts that the level of stress will influence the type of interaction between individuals shifting from positive (i.e., facilitation) in stressful environments to negative (i.e., competition) in favourable environmental conditions (Bertness & Callaway 1994). Facilitation is defined as 'an interaction in which the presence of one species alters the environment in a way that enhances the growth, survival and reproduction of a second species' (Bronstein 2009, p.1160). Interactions show 'conditionality' when costs and benefits, and thus outcomes, are affected in predictable ways by ecological conditions (Bronstein 1994). Plant interactions thus depend on the ecological settings in which they take place, and in particular, plants adjust to the level of stress (Bilas, Bretman & Bennett 2021). It has been implicitly assumed that facilitation only occurs between individuals of different species, but as shown and discussed by Fajardo & McIntire (2011), facilitation can be found between individuals of the same species and was indeed suggested in several other cases (Proença et al., 2019; Reijers, Akker, Cruijsen, Lamers & Heide 2019). Facilitation allows, or even favours, the coexistence of individuals of the same species in close proximity, i.e., within their zone of interaction (ZOI sensu Stoll & Weiner 2000). If the ZOI of related individuals overlaps this would allow kin recognition, kin discrimination and kin selection mechanisms to operate. Following SGH predictions, harsh environmental conditions should lead to stronger positive interactions (i.e., cooperation) if related individuals are present. We can thus expect contrasting responses for individuals living in stressful versus benign environments and thus, in a changing environment, the same closely related individuals could cooperate or compete depending on the level of stress.

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So far, few studies have shown evidence that kin recognition depends on the level or the type of environmental stress. In these studies, kin recognition and discrimination responses were triggered by several abiotic factors including nutrient availability, for example, in *Glycine max, Caesalpinia pulcherrima, Populus tomentosa* (Li, Xu & Liu 2018), and *Pisum sativum* (Pezzola, Pandolfi & Mancuso 2020), heavy metal concentrations in *Sorghum vulgare* (Li, Xu & Feng, 2018) and biotic factors such as the presence of another species in *Medicago* genera (Ehlers, David, Damgaard, & Lenormand 2016). On the other hand, water availability, did not trigger kin recognition responses in *Glechoma hederacea* (Goddard, Varga, John & Soulsbury 2020). Thus, it is possible that kin recognition is only expressed in a given kin group under certain ecological conditions, which can only be identified by modulating the type of stress and its intensity. We note then that it is essential to include both several stressors and several

genotypes (kin groups) in any experimental effort to elucidate kin recognition/selection in plants.

Kin discrimination also depends on whether cooperation is obligate or facultative. In the case of obligate cooperation, kin recognition is ultimately not useful as individuals always cooperate whatever the identity of their neighbour is and there is no discrimination. In contrast, facultative cooperation implies that individuals exhibit phenotypic plasticity for cooperation and that they are able to distinguish relatives (i.e., present kin recognition). Both phenotypic plasticity (Ghalambor et al., 2007) and kin recognition (Rousset & Roze 2007) have associated costs. To identify cases where obligate or facultative cooperation is favoured requires thus to estimate the costs and benefits of facultative cooperation (plasticity) in addition to those of Hamilton's equation (Hamilton 1964), weighted by the probability of interacting with a relative. Therefore, in the case of facultative cooperation, the question is not only with whom but also under what conditions should individuals cooperate? In a stable environment, the phenotype is constant and obligate cooperators would be favoured. In a changing environment, SGH suggests that interactions shift between positive and negative and being a facultative cooperator would be advantageous. In the latter case, cooperation will depend on the relatedness between individuals. The important point here is that environmental conditions in which the experiment takes place, but which are seldom considered in kin studies, condition the detection of kin recognition/selection.

Cooperating throughout life? Interactions in long-lived species

For practical reasons, most of the studies on kin recognition thus far have focused on annual or short-lived plants (Subrahmaniam, Roby & Roux 2021). Few studies have investigated long-lived species such as perennial herbs, shrubs or even trees (Till-Bottraud, Fajardo & Rioux 2012; Fajardo, Torres-Díaz & Till-Bottraud 2016; Pickles et al., 2017), and mostly addressed a short period of their lives (e.g., the seedling stage; Takigahira & Yamawo 2019; He et al., 2021). Some authors reported positive interactions between mature related individuals following herbivory, through discrimination of volatile chemicals emitted from relatives (Karban et al., 2014; Hussain, Rodriguez-Ramos & Erbilgin 2019). Other showed that related seedlings grow better in the presence of kin (Assay et al., 2020) and benefit more from the mycorrhizal network shared with related neighbours (Pickles et al. 2017).

However, long-lived species exhibit several interesting characteristics that have not been much considered: (i) the long lifespan means that an individual will likely experience different environmental situations during its life; (ii) different life stages may have very different ecologies (ontogenetic stages, e.g., seedling vs. mature tree, related to the regeneration niche, Grubb 1977), kin discrimination and kin selection might thus be advantageous at a given developmental stage but not at others. For example, at early stages when seedling mortality is important because they are more sensitive to harsh conditions, positive interactions between related individuals would ensure better survival of related genotypes; (iii) some long-lived species, particularly trees, strongly affect their physical environment by improving their living conditions (i.e., the ecosystem engineering effect sensu Jones, Lawton & Shachak 1994), modulating the level of stress and improving resource availability through their effects on the local environment (McIntire & Fajardo 2014). As this effect is incremental through time it will be all the more important in long-lived species (Hortobágyi, Corenblit, Steiger & Peiry 2018). As a given individual can experience different conditions through its lifetime, the chance of detecting kin recognition could strongly depend on the life stage considered. This particular point is essential to consider when conducting field studies where local conditions can change or even be improved by engineering species. All these three characteristics are linked to the SGH, although the gradient here is temporal as long-lived species interact with their neighbours through decades or even centuries in a fluctuating or changing micro- and macro-environment.

Moreover, in seasonal environments, the physiological activity is reduced or potentially halted by dormancy periods. Is this interaction between two long-lived individuals just one absolute interaction or is it repeated interactions (e.g., in the different growing seasons), which would lead to radically different outcomes (Axelrod & Hamilton 1981)? This is still an open question, for which evolutionary game theory (Maynard Smith 1982) could provide a theoretical framework.

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Spatial patterns of plant interactions: the added value of spatial genetic structure

One potentially valuable tool to search for positive interactions among kin in natural populations is the spatial genetic structure (SGS). The SGS, defined as the non-random spatial distribution of genotypes, characterizes the relationship of relatedness estimates between pairs of individuals and their physical proximity (Loiselle, Sork, Nason & Graham 1995; Hardy 2003). The genetic composition of a population has an influence on the type of biotic interaction

that can occur among immediate neighbours (Fajardo, Torres-Díaz & Till-Bottraud 2016). Using a spatial genetic approach can thus offer complementary insights, particularly for in situ studies. A significant SGS indicates that related individuals coexist close to each other, and thus, the possibility of kin interactions increases. This makes SGS an interesting tool to detect the presence of related individuals and to consider potential positive interactions in the context of kin recognition and selection. Moreover, this SGS approach seems particularly relevant to us as kin selection can evolve in viscous populations without the need for kin recognition or discrimination (e.g., obligate cooperation) (Waldman 1988, Gardner & West, 2004). However, the fact that individuals are "close to each other" does not mean that they will necessarily interact. Plant-plant interactions are inherently local in nature. They mostly occur among immediate neighbours, in which case an individual's direct effect on another (negative or positive) will have a delimited influence in space, i.e., the ZOI (Stoll & Weiner 2000). The ZOI will grow as individuals grow because individuals, in particular trees, will occupy a larger area with their aerial and belowground organs. Thus, over their lifetime, individuals will interact with other individuals located further away as their ZOI expands, and thus allowing more or different individuals to interact through time. Biotic interactions will be influenced by the local SGS (i.e., the presence of related individuals within the ZOI) but can also significantly influence the SGS within the ZOI of individuals through the survival of related individuals in local patches, which can be monitored through time. However, very few studies have linked the SGS and the ZOI when interactions between related individuals take place (Till-Bottraud, Fajardo & Rioux 2012; Segovia, Vásquez, Faugeron & Haye 2015; Fajardo, Torres-Díaz & Till-Bottraud 2016).

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The relationship between the SGS and intraspecific interactions: two case studies in long-

lived species

To illustrate how all the aspects we have outlined above could help us assess the ultimate importance of kin selection, we describe here two examples of positive interactions between related individuals in tree species.

First, in *Nothofagus pumilio* (Nothofagaceae) second-growth stands located in the Chilean Patagonia, multi-stemmed trees are formed by related individuals. They are restricted to forest edges where stress (wind and desiccation) was very high at the regeneration stage, and

where facilitation increased seedling survival (Fajardo & McIntire 2011) and growth in multistemmed adult trees (Fajardo & McIntire 2010; McIntire & Fajardo 2011). Trees from different multi-stemmed clusters are unrelated, and no group dispersal mechanism is reported for this species (Till-Bottraud, Fajardo & Rioux 2012; Fajardo, Torres-Díaz & Till-Bottraud 2016) indicating better survival when related seedlings happen to grow together. The ecological mechanism involved here may be an altruistic or cooperative fusion of trunks, which would be physiologically and biomechanically selectively advantageous for resource acquisition or stability against wind (Tarroux & DesRochers 2011; Tarroux, DesRochers & Tremblay 2014; Fajardo, Torres-Díaz & Till-Bottraud 2016), but fusion may also involve some costs (e.g., resource sharing, pathogen transmission). Stem merging scales up the level of organization to the group and therefore competition does not occur among individuals but among groups of trees (McIntire & Fajardo, 2011). In a kin selection framework, fusions would be selected and would only occur between related individuals, as indirect fitness gains might offset these costs.

The other example is the Black Poplar (Populus nigra L. Salicaceae) in a riparian context. At the establishment phase, P. nigra seedlings grow in high densities on exposed alluvial bars of rivers, and facilitation is the major occurring interaction (Corenblit et al. 2018). Young individuals form dense patches and mechanically protect each other from destruction (Corenblit et al., 2014). During colonisation, *P. nigra* is known to enhance its local conditions (i.e., ecosystem engineer species) when growing at high seedling densities by trapping fine sediments and thus increasing nutrient availability and decreasing drought stress in its natural river reach habitat (Corenblit et al., 2014; Hortobágyi, Corenblit, Steiger & Peiry 2018). The modification of the environment by individuals can prevent, delay, or accelerate changes in the types of interactions between individuals (Corenblit et al., 2018). Based on the evolution of the SGS in a natural population, a shift in the type of interactions from positive to negative has been proposed in *P. nigra* stands along riparian corridors (Mazal et al., 2021; Mazal, Corenblit, Fumanal & Till-Bottraud 2022) (Fig. 1). The case of *P. nigra* shows that, in long-lived species, shifts in the type of interaction are possible and could be monitored by studying SGS (Mazal, Corenblit, Fumanal & Till-Bottraud 2021). A cohort monitoring approach from establishment to maturity in long-lived species could be an efficient approach to follow the evolution of interactions between individuals and ultimately to estimate the survival component of fitness (not just reproduction). In addition, cohorts should also be followed in contrasted conditions, if possible along a gradient of environmental conditions, as the type of interaction may vary with life stages and environmental conditions. An increase in SGS over time could be an indication

of positive intraspecific interactions between relatives and thus would warrant further validation.

These examples open new research avenues to consider the impact of genetic variability, environmental conditions, life-span and SGS in kin selection studies. However, we believe that some semantic and methodological aspects need clarification to accurately address kin recognition, discrimination or selection in such studies.

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Kin recognition, kin discrimination and kin selection

Kin recognition, kin discrimination and kin selection are often used alternatively in articles, however, they correspond to different concepts (see the review on kin recognition definitions by Penn & Frommen 2010). The confusion between kin recognition and kin discrimination probably comes from the fact that the only way to detect kin recognition in plants is through its expression (trait or behaviour variation between kin and non-kin neighbour treatments), i.e., kin discrimination. Indeed, many studies that showed the expression of a different phenotype depending on the genetic identity of the neighbours used the term kin recognition but de facto showed kin discrimination (Dudley & File 2007a, 2007b; Milla, Forero, Escudero & Iriondo 2009; Murphy & Dudley 2009; Lepik, Abakumova, Zobel & Semchenko 2012; Semchenko, Saar & Lepik 2014; Yang, Li, Xu & Kong 2018; Li, Xu & Liu 2018; Goddard, Varga, John & Soulsbury 2020). Although kin discrimination can be captured by measuring any kind of trait that shows plasticity between kin and non-kin neighbours, its implication for kin selection can only be inferred if it translates into fitness differences between the two treatments (Ehlers & Bilde 2019). However, fitness is very rarely or only approximatively measured in "kin recognition" studies and some authors tend to extrapolate their results on kin recognition or discrimination and conclude without proof that there is evidence of kin selection in plants (Crepy & Casal 2015; Zhang et al., 2016; Pezzola, Pandolfi & Mancuso 2020).

Studies using biomass-related metrics (Fig. 2a) often suggest that reducing below-ground biomass is a cooperative behaviour indicative of kin selection. Increasing below-ground biomass when growing with neighbours (a competitive trait) diverts resources from producing more above-ground biomass. If the above-ground biomass is a good proxy for fitness (Younginger, Sirová, Cruzan & Ballhorn 2017), a cooperative decrease in below-ground biomass of all interacting individuals could thus increase their fitness (both direct and indirect if kin are involved). However, the relationship between biomass and fitness is not always

correct, especially in long-lived species such as trees, where biomass provides little information on how individuals will reproduce or may reflect many confounding factors, making their interpretation difficult (see the methodological issues discussed below and in Fig.2). In fact, results based on the observation of biomass differences can be interpreted in opposite ways (Subrahmaniam, Roby & Roux 2021). To our knowledge, the only article that came close to demonstrate a possible gain in fitness for closely related individuals (through the study of reproductive traits) is Torices et al. (2018). They showed that Moricandia moricandioides (Brassicaceae) individuals growing with kin produced larger floral displays than those growing with strangers. The production of a larger floral display is costly but attracts more pollinators that will also benefit their neighbours. The authors argue that individuals compensate for the decrease in direct fitness caused by the production of large flowers with a gain in indirect fitness by attracting more pollinators who will visit their related neighbours (Torices, Gómez & Pannell 2018). In this case, related neighbours benefit from the attraction of pollinators that will pollinate their stigmas and export their pollen. These results were further supported by a second study which showed that plants growing with kin invest more in petal pigments, nectar volume and sugar content (Torices, DeSoto, Narbona, Gómez & Pannell, 2021).

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Following the approach proposed by Ehlers & Bilde (2019), we recommend to move on to the next step and to study kin selection per se generalizing the use of Hamilton's rule (rB > C), and the measurement of the cost (C) and benefit (B) components of fitness separately on each plant that take part in the interaction (note that, contrarily to what is stated in Ehlers & Bilde (2019), only interacting related individuals should be included for the estimation of indirect fitness), as well as their true (realized) relatedness (r), which can now be reliably estimated using next-generation sequencing techniques (de Villemereuil, Gaggiotti, Mouterde & Till-Bottraud 2016). Two particular cases need to be detailed: if r = 0, no indirect fitness can be obtained. Conversely, if r = 1, Hamilton's rule is no longer needed to explain the outcome of the interactions because each interactant is actually a clone of the same individual and the indirect fitness is actually a direct fitness. This is the case in studies where clones are used as kin groups (Yamawo, Sato & Mukai 2017). The case of highly self-pollinating species (i.e., same accession or variety) (Simonsen, Chow & Stinchcombe 2014; Yang, Li, Xu & Kong 2018; Pezzola, Pandolfi & Mancuso 2020) is slightly different because although very close to 1, r is actually slightly lower than 1. In a literature survey Subrahmaniam et al. (2021) showed that 44% and 10% of the studies that investigated interactions between related and unrelated individuals were conducted in self-pollinating or clonal species, respectively. Can we extend the results of these studies to other species that are not self-pollinated, and how general is a mechanism if it is only shown to work for clonal or self-pollinating species? Although the cases where *r* is very close to 1 are convenient to study, we think that they are rather extreme cases and that they are therefore difficult to generalize to other species. In natural populations without inbreeding, the expected relatedness full-sibs is 0.25, and values measured for ants and bees where kin selection has been demonstrated range from 0.5 to 0.7 (Gadagkar et al. 2019; Kay, Lehmann & Keller 2019). We argue that results obtained from highly-selfing or clonal species should be interpreted in the same way as studies on self-recognition (i.e. recognition of individual genotype between two ramets, be they connected or disconnected) (see also Bilas, Bretman & Bennett 2021).

As underlined by Hamilton's rule, the level of relatedness between individuals is critical to predict the outcome of the interactions. In a local population, varying levels of genetic relatedness can co-exist and thus patterns of interactions between individuals can vary based on their relatedness (Subrahmaniam, Roby & Roux 2021). Following Hamilton's rule, the potential benefits of interaction should decrease linearly or display a threshold (Till-Bottraud & de Villemereuil 2016) beyond which it is no longer advantageous to cooperate (C > rB). However, the gradient of relatedness between individuals is rarely addressed in studies and this raises the question of the generalized use of studies that tested interactions between individuals originating from a single population (Subrahmaniam, Roby & Roux 2021). Few studies thus far have used different levels of relatedness between individuals. Karban et al. (2014), for example, used a gradient of relatedness and found that Artemisia tridentata individuals responded more effectively to volatile cues from close relatives than from distant relatives when subjected to herbivory. Another study using rice cultivars with different levels of relatedness showed that plants altered their root behaviour and biomass allocation in response to relatedness level (Yang, Li, Xu & Kong 2018). Moreover, using a relatedness gradient, Xu et al., (2021) showed that plants discriminate their neighbours (related or unrelated) and adjust their growth, competitiveness and chemical defence accordingly to improved weed suppression in rice (through allelopathic processes) (see also commentary from Anten & Chen 2021b). Following on the authors' discussion, we also believe that a relatedness gradient is an important factor that can potentially trigger responses when plants interact with neighbouring individuals and thus must be considered in future studies.

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Methodological challenges in studying kin recognition and selection

Whether studies did (see the review by Anten & Chen 2021a) or did not find evidence of kin recognition (e.g., Milla, Forero, Escudero & Iriondo 2009; Lepik, Abakumova, Zobel & Semchenko 2012), much caution in their interpretations must be taken. When comparing biomass of plants growing with kin or strangers, alternative hypotheses and several confounding factors could explain the results and should be tested. A common difficulty to find evidence of kin recognition in experiments is the lack of appropriate control treatments (Fig.2b). In pot experiments, the use of a control treatment where plants are grown alone is mandatory to determine if plants do compete or not (compared to plants grown alone). This control treatment is also necessary to detect a potential genotype that has a better intrinsic growth. Genotypes with better intrinsic growth capacities can lead to size inequality in groups of plants (Simonsen, Chow & Stinchcombe 2014; Li, Xu & Liu 2018). Another confounding factor is competitive ability, i.e. the relative growth of individuals grown in competition compared to when grown alone (Masclaux et al., 2010; Zhang et al., 2019). Although it is usually argued that a species can win either through being better at suppressing the performance of others (effect) or being better at tolerating competition (response), Hart et al. (2018) showed that only the competitive response is relevant to a species' competitive ability. Following early works on kin recognition and discrimination in plants, several papers encouraged testing for competitive ability (Masclaux et al., 2010; Chen, During & Anten 2012). However, this is still not a standard procedure across studies and the lack of appropriate control treatments remains common in the literature (Semchenko Saar & Lepik 2014; Crepy&Casal 2015; Murphy et al., 2017; Murphy, Swanton, Van Acker & Dudley 2017; Li, Xu & Feng 2018; Pezzola, Pandolfi & Mancuso 2020). Here, we would like to emphasise that this lack of control treatment may result not only in misinterpretation but also in the failure to reveal kin discrimination in studies (that have therefore not been published). To test for competitive ability, it is necessary to set all the kin groups in competition with each other and test whether a kin group always performs better than the others prior to wrongly conclude to kin discrimination.

When plants grown with relatives were found to exhibit greater belowground biomass than plants grown with strangers, the results were mostly interpreted using the niche partitioning framework (File, Murphy & Dudley 2012; Subrahmaniam, Roby & Roux 2021). Niche

partitioning predicts that relatives overlap more in their niche use and compete more with each other compared with unrelated conspecifics (Young 1981; Milla, Forero, Escudero & Iriondo 2009). However, stronger competition between relatives can be a by-product of forced competition for a few dimensions of the niche that occurs when plants are grown at high density in pots. The niche is a multidimensional concept (Hutchinson 1957; González, Dézerald, Marquet, Romero & Srivastava 2017) with each component (e.g., light, nutrient, water, wind, space, microbiota) occupying one dimension. Pots strongly constrain the growth of roots for example in space or for the types of nutrients available. In pots, individuals share and compete in a restricted number of dimensions and there may be no possible compensation in other dimensions whereas in nature, individuals can compete in some dimensions of their niche but not in others (Barker et al., 2017). Pot size can thus obscure the effects of positive interactions between individuals (Hess & de Kroon 2007; Chen, During & Anten 2012) because root growth responds to the available volume (McConnaughay & Bazzaz 1991), but also to nutrient amounts (O'Brien, Gersani & Brown 2005) and concentrations (Maina, Brown & Gersani 2002). Therefore, experiments conducted in restricted conditions (i.e., few niche dimensions) will bias the conclusions towards niche partitioning versus kin discrimination and would not be relevant in natural conditions where individuals growing with relatives can exploit all the dimensions of their niche. On the other hand, when plants are supplemented with nutrients and resources are not limiting, one can expect to find little or no difference between the stranger and kin treatments because every individual has access to sufficient resources for its growth. Finally, kin cooperation (kin selection) and competition (niche partitioning) can occur simultaneously, possibly acting on different traits and thus different niche components (File, Murphy & Dudley 2012). Thus, depending on the observed traits and growth conditions it is possible to observe one process but to miss the second. Therefore, although in situ experiments are challenging, we share Anten & Chen's (2021a) appreciation that in situ experiments could solve some of the problems associated with experiments under controlled conditions, in particular the constrained niche aspects.

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Conclusion and final notes

Over the last decade the scientific literature has continuously explored the role of relatedness in the interactions between plants. However, we assert that some points have been generally neglected and are worth considering in kin recognition/selection studies. The major aspects are the underestimated role of intraspecific diversity for kin recognition and kin selection traits and

the fact that local conditions can influence the outcome of interactions. The concept of "conditionality" and the "stress gradient hypothesis" suggest that intra-specific interactions, and in particular interactions between relatives, are dependent on environmental conditions, especially under various stress types. The case of long-lived species, where individuals face different environmental conditions and stresses throughout their life, is especially promising. In addition, studying the spatial genetic structure of populations for *in situ* studies would be of great help in pointing out settings in which different types of interactions might take place. We also discuss the challenges of distinguishing kin discrimination from kin selection, which are to accurately measure the costs and benefits of the interaction as well as the relatedness of the interacting individuals, and the importance of appropriate control treatments to test for all alternative hypotheses (Fig. 2). We also strongly recommend several key-points to be considered in future studies (Fig. 2b): to capture a large intraspecific diversity of the studied species, to include different environmental conditions and to include a relatedness gradient between individuals. This would help to take care of various confounding effects that could interfere with the interpretation of the results.

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Figures

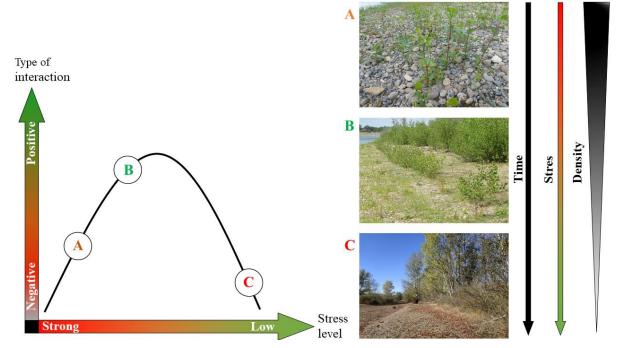


Figure 1. The stress gradient hypothesis (SGH, Bertness & Callaway, 1994) applied to the case of *Populus nigra*. A: Seedlings establish in densities on bare surfaces along rivers and experience harsh conditions (mechanical and hydric stresses). Facilitation occurs due to the 765 high densities of individuals and reduces the stresses for each individual. At this stage no spatial genetic structure (SGS) is found and the main direct interactions between individuals are neutral (commensalism). B: Patches of young individuals that survived now exhibit SGS. The effect of *P. nigra* on the environment enhance the local conditions (soil moisture and nutrients). Facilitation still occurs because of the protective effect of individuals located at the upstream part of the stand (Mazal et al., 2021). Main direct interactions shift to positive (potentially cooperation) and could be directed towards related individuals (as SGS is significant at this stage). C: At older stages, individuals no longer benefit from the presence of their neighbours and the local conditions are no longer stressful. Stem density is highly reduced through self - thinning processes (i.e., competition) (Yoda et al., 1963). The main interactions between individual are either neutral (commensalism) or negative (competition).

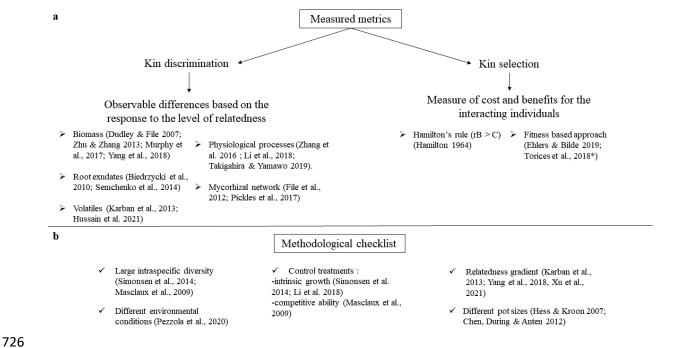


Figure 2. Suggested guidelines for the design of experiments on kinship studies in plants. (a) Measured metrics commonly used in the conceptual frameworks studied: kin recognition discrimination or kin selection. (b) Checklist for the design of future studies outlining key points to be controlled. For both (a) and (b), we have mentioned the studies that in our opinion have taken some of the points above into account and that could be used as a reference. *As discussed in the text, although this study did not measure the fitness of individuals, it is the closest evidence to date for kin selection in plants.