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Predicting plant invaders in the Mediterranean through a weed risk assessment system

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Abstract Risk assessment schemes have been developed to identify potential invasive species, prevent their spread and reduce their damaging effects. One of the most promising tools for detecting plant invaders is the weed risk assessment (WRA) scheme developed for Australia. Our study explores whether the Australian WRA can satisfactorily predict the invasion status of alien plants in the Mediterranean Basin by screening 100 invasive and 97 casual species in Spain. Furthermore, we analysed whether the factors taken into account in the WRA are linked to invasion likelihood (i.e., invasion status) or to impacts. The outcome was that 94% of the invasive species were rejected, 50% of the casual species were rejected and 29% of them required further evaluation. The accuracy for casuals is lower than in other studies that have tested non-invasive (i.e., casuals or non-escaped) alien species. We postulate that low accuracy for casual species could result from: (1) an incorrect "a priori" expert classification of the species status, (2) a high weight

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of the WRA scores given to potential impacts, and (3) casual species being prone to becoming invasive when reaching a minimum residence time threshold. Therefore, the WRA could be working as a precaution early-warning system to identify casual species with potential to become invasive.

Keywords Alien plants · Casual plants · Mediterranean region · Species traits · Weed risk assessment

Introduction

Predicting the success of alien species has been a major aim of ecological research since invasions were recognized as an important conservation issue (Rejmánek and Richardson 1996; Reichard and Hamilton 1997; Smith et al. 1999; Kolar and Lodge 2001; Pyšek 2001; Caley et al. 2006). To this purpose, many studies have focused on finding which biological traits make a species invasive and the characteristics of invaded habitats (Goodwin et al. 1999; Daehler 2003; Heger and Trepl 2003; Richardson and Pyšek 2006). Since Baker (1965), the search for invader syndromes (i.e., suites of traits and abiotic factors associated to invasiveness) aims to improve our ability to predict the invasion success of alien plants in new regions (Richardson and Pyšek 2006). For example, Thuiller

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et al. (2006) have found that the spatial regional distribution of invasive alien species in South Africa is driven by life forms, reproductive traits and human uses.

The accumulated knowledge on invader traits together with the characteristics of invaded habitats, and information in whether a species is invasive in other parts of the world, especially in areas with similar environmental conditions (Westbrooks 1981; Rejmánek 2000; Union of Concerned Scientists 2001; Thuiller et al. 2005; Richardson and Thuiller 2007), has been the basis to develop risk assessment schemes that attempt to predict the success of alien species in a given region of introduction (Mcneely et al. 2001; National Invasive Species Council 2001; Wittenberg and Cock 2001; Leung et al. 2002; Keller et al. 2007a; Gordon et al. 2008). For plants, only a small proportion of introduced alien species become invasive (Di Castri 1989; Williamson 1996) and it is important to identify them to prevent their spread and impacts.

Because risk has two components: likelihood of invasion and ecological or socioeconomic impacts, risk assessment schemes attempt to identify this small fraction of species with a high likelihood of becoming invasive, and also to prevent their spread and their damaging effects (Andersen et al. 2004). So far, there is evidence that the implementation of risk assessment protocols produces net economic benefits (Keller et al. 2007b).

The scientific literature abounds in risk assessment schemes differing in the methods used and the phase of invasion process they target. Recently, taking advantage of the rapid increase in computing tools, facilities and database availability, new screening models have been developed. The most convenient to compare across regions are the quantitative tests that offer a species score. One of the most promising as a tool for detecting plant invaders is the Australian weed risk assessment (WRA, Pheloung 1995) which has also been tested for New Zealand (Pheloung et al. 1999) and implemented in other regions (Hawaii and Pacific islands: Daehler and Carino 2000; Daehler et al. 2004; Bonin Islands: Kato et al. 2006; Czeck Republic: Křivánek and Pyšek 2006; Italy: Crosti et al. 2007). A recent review has compared the WRA accuracy in different geographic regions (Gordon et al. 2008) and has found it to be high. The WRA

scheme can be adopted as an initial screen for plant species proposed for introduction in a new region with the aim to reduce their economic and ecological impacts.

Our study explores whether the original Australian WRA scheme (Pheloung et al. 1999) satisfactorily predicts the invasion success of alien plants in the Mediterranean region. For this purpose, we have chosen Spain as a representative because of its climatic, geological, landscape and habitat heterogeneity and high species diversity (Medail and Quezel 1997). In many studies the WRA scheme has been used to evaluate "major invaders", "minor invaders", and "non-invaders" (species not-escaped from cultivation or casuals) (Gordon et al. 2008). In this study we compare the accuracy for invasive and casual (non-invasive) species. We follow Richardson et al. (2000a) and Pyšek et al. (2004) in defining invasion plant status. Invasive species are alien species with self-sustaining populations not requiring direct human intervention, that produce offspring at considerable distances from the parent plants, and thus, have the potential to spread over large areas. Casual species do not form self-replacing populations outside cultivation and rely on repeated introductions for their persistence. Differently from other studies, we did not include non-escaped from cultivation species because we found it difficult and not very accurate to randomly chose a list of non-escaped species while among the invasive and casual species sets there are several pathways of introduction (e.g., 21% of the 1,000 alien species listed in Sanz-Elorza et al. 2004 are unintentionally introduced). Furthermore, by testing casual species the screening protocol can be interpreted as an early screening system of potential invaders among casual species. Moreover, in our study we go a step further by analysing whether the factors taken into account in the WRA are those really involved in differentiating invasive from casual alien plants, considering status differences to be a consequence of invasion likelihood.

Specifically, our main questions are (1) Does the WRA have a high accuracy in identifying invasive species when adapted and applied to a Mediterranean region? and (2) What are the determining factors for an alien plant species to be considered invasive and do these correspond with the questions taken into account to calculate the WRA final score?

Methods

Weed risk assessment for Mediterranean regions

The scheme chosen for testing was the original Australian WRA (Pheloung 1995) (See "Appendix 1"). The WRA system consists of 49 questions which encompass biogeography, undesirable plant attributes and biology/ecology (Pheloung et al. 1999). A high score (>6) identifies a species likely to be of high risk (weed) and rejects it for import; a low score (<1) accepts the plant for import (non-weed) and intermediate scores (1–6) require further evaluation. The WRA score ranges from -14 (benign species) to 29 (maximum risk).

From the original WRA we modified the question 2.01 from "Species suited to Australian climates" to "Species suited to Mediterranean climates". This question and the 2.02 were answered without performing any climatic model, but following the criteria described in Daehler et al. (2004). We also modified the question 5.03: "Nitrogen fixing woody plant" to "Nitrogen fixing plant", to include the non-woody nitrogen fixing plants. These are an important component of Spanish alien flora, many of such species being very abundant in ruderal, disturbed habitats (Sanz-Elorza et al. 2004).

Screened species and information sources

We screened a plant data set that was comprised of information on 100 invasive and 97 casual alien plants recorded in the Atlas of Invasive Plant Species in Spain (Sanz-Elorza et al. 2004), including a broad range of life-forms (See "Appendix 2"). Species status was defined following Richardson et al. (2000a) and Pyšek et al. (2004). The chosen invasive species comprised all the most invasive species listed in the Atlas, while the casual species were selected randomly from the pool of 380 casual species recorded therein. To answer the WRA questions, the information was gathered from regional and local floras (Castroviejo et al. 1986-2000; Bolòs et al. 2005), weed atlases (Sanz-Elorza et al. 2004), and Internet databases such as: BioFlor (http://www.ufz.de/biolflor), Plants for a Future (http:// www.comp.leeds.ac.uk/pfaf), Hypermedia for Plant Protection—Weeds (http://www.dijon.inra.fr/hyppa), Poisonous Plants of North Carolina (http://www. ces.ncsu.edu/depts/hort/consumer/poison), species accounts from Plantas Invasoras en Portugal (http:// www.uc.pt/invasoras), USDA Plants database (http:// plants.usda.gov), International Survey of Herbicide Resistant Weeds (http://www.weedscience.org), Global Compendium of Weeds (http://www.hear.org/gcw), Global Invasive Species Database (http://www.issg. org/database/welcome), Weeds in Australia (http:// www.weeds.gov.au), and Ecological Traits of New Zealand Flora (http://ecotraits.landcareresearch.co.nz).

Data analysis

To evaluate the predictive ability of the WRA applied in Spain, we developed a receiver operating characteristic (ROC) curve (DeLong et al. 1988) because it has been found to be a useful tool for evaluating the performance of invasive species screening tests (Caley and Kuhnert 2006; Hughes and Madden 2003; Gordon et al. 2008). This analysis requires comparison of two groups: one for which rejection is the incorrect outcome and the other for which it is the correct. A ROC curve is formed by plotting the proportion of true positives (i.e., rejected invasives) against the proportion of false positives (i.e., rejected casuals) across the range of cutoff points on an indicator scale (i.e., the WRA score). The area under the ROC curve (a value between 0.5 and 1) represents the probability that a randomly chosen positive case (an invasive species) will have a higher test value (WRA score) than a randomly chosen negative case (a casual species) (DeLong et al. 1988). The closer the area under the ROC curve is to one, the better the screening tool's ability to differentiate between the two groups (Lasko et al. 2005).

A principal components analysis (PCA) was used to identify the main variables that characterize the pool of all tested alien species and to reduce the number of multivariate data for a posterior comparison between invasive and casual species characteristics. This linear method was chosen because the number of species was low (197) compared to the number of variables (30) and the length of the longest gradient from Detrended Correspondence Analysis was between 1 and 2 (Lepš and Šmilauer 2006). Ordination was performed using CANOCO for Windows 4.5.

We adopted the framework of the generalized linear mixed models (GLMM) (e.g., Blackburn and

Duncan 2001) to determine variation between species in invasion status (i.e., casual or invasive) and WRA score (i.e., two estimations of invasion risk) as a function of the 15 WRA variables best characterized by the first two principal components (Fig. 2). Both GLMM analyses incorporated taxonomy to overcome potential phylogenetic biases. Because species are linked by phylogeny (Harvey and Pagel 1991), using species as independent data points may inflate the degrees of freedom (Brändle et al. 2003) and increase the Type-I error. GLMM deals with phylogenetic effects by allowing the incorporation of taxonomic categories as nested random factors. In this way, the likely non-independence of response values of closely related species is controlled by assuming a common positive correlation between introduction outcomes for species within nested taxonomic groups (genera and family, in our case). Conversely a zero correlation is assumed between introduction outcomes for species in different groups (a variance components model). There are more sophisticated procedures that allow implementing the complete phylogenetic structure in the model (Sol et al. 2008), but we could not use such methods because there was no robust phylogenetic hypothesis available for the species studied. In addition, the approach we used helped to mitigate the problem of invasive species being a non-random subset of all species introduced (Blackburn and Duncan 2001).

We modeled invasion status with a Binomial distribution of errors (Crawley 2002), because the response variable was binary [i.e., casual(0)/invasive(1)]. Invasion risk (i.e., WRA score) followed a normal distribution and did not need transformation to achieve the requirements of parametric analysis, so we modeled it with a Normal distribution of errors. In both cases, the inspection of the residuals showed

that error structures adjusted well to our response variables.

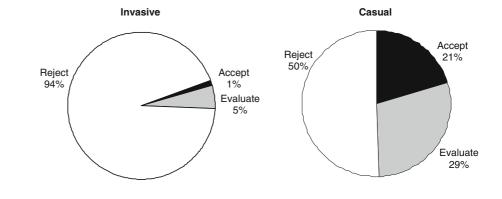
We started modeling invasion status with a full model that contained all our predictors. Using a backward selection process, we next simplified the model so as to leave only significant predictors (minimum adequate model). The model was run in the glmmPQL procedure of the MASS library on the R statistical package (Venables and Ripley 2002; R Development Core Team 2006).

Results

We were able to answer a mean of 43 questions (range 29–49 questions) out of 49, after a detailed search of at least 5 h on information for each species. Furthermore, for each species the test was answered by two different persons to avoid subjectivity or biases. While only one invasive species (*Chloris gayana*) was accepted, the system accepted 21% of casual species. Surprisingly, half of the casual species were rejected and many of them needed further evaluation (Fig. 1). The area under the ROC curve for the WRA classifying plants as invasive or casual was 0.79 (Fig. 2).

When performing the principal components analysis, the first two principal components explained 24% of the variability in the species data. Out of the 49 WRA questions, there were only 15 that were best characterized by the first two principal components, so they were the ones that better predicted the total of alien plant species variation. The PCA biplot (Fig. 3) indicates invasion-related traits, which are assembled in three different groups. The first group contains crop and gardening plants, most of them nitrogen fixing, bird dispersed, and forming dense thickets. Many species included in this cluster belong to the

Fig. 1 Results of the Australian weed risk assessment system of Pheloung et al. (1999) applied to 100 invasive and 97 casual plant species in Spain. The percentage of species rejected, accepted or suggested for further evaluation is indicated



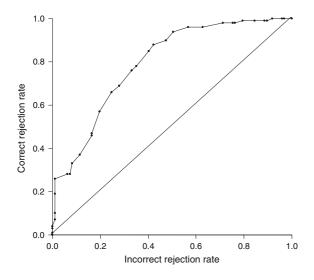
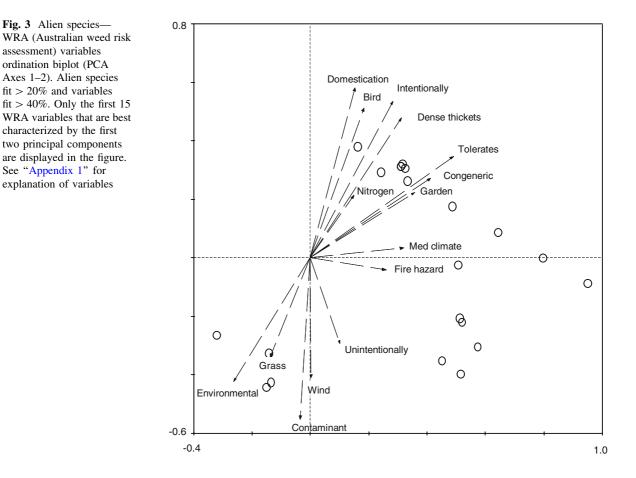


Fig. 2 Receiver operating characteristic (ROC) curve for the WRA classifying plants as invasive or casual species in Spain. The area under the ROC curve was 0.79

genus Acacia, Opuntia and Lonicera. The second group comprises species of Mediterranean origin and species that may create a fire hazard, such as *Gleditzia triacanthos* and *Eucalyptus globulus*, respectively. Finally, the last cluster is mainly composed of grasses (e.g., *Eleusine indica*, *Sorghum halepense*) and species belonging to Asteraceae and Amaranthaceae, and also species which are unintentionally or wind dispersed, or potentially dispersed as a contaminant.

There was a clear taxonomic bias in the set of all alien species analysed, with 33% of species belonging to just three families; Asteraceae (12%), Fabaceae (11%), and Poaceae (10%). Thus, clustering the species according to taxonomic relationships throught the GLMM analysis, we found that only five out of the 15 PCA predictors were significantly related with the invasion status of the species: existence of congeneric weeds, creating fire hazard, intentional



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Table 1 Minimum adequate mixed model accounting for variation in invasion status (i.e., invasive or casual species)

					-
Variable	Estimate	Standard error	DF	<i>t</i> -value	Р
Intercept	-1.249	0.653	60	-1.912	NS
Congeneric	2.476	0.519	42	4.767	< 0.0001
Fire hazard	-1.700	0.795	42	-2.137	< 0.05
Intentionally	-1.312	0.558	42	-2.349	< 0.05
Contaminant	1.557	0.544	42	2.861	< 0.01
Bird	1.214	0.583	42	2.082	< 0.05

A positive estimate value indicates a positive relationship with the invasion status. See "Appendix 1" for explanation of variables

 Table 2
 Minimum adequate mixed model accounting for variation in WRA final score

Variable	Estimate	Standard error	DF	<i>t</i> -value	Р
Intercept	-8.875	1.787	48	-4.966	< 0.0001
Domestication	-3.894	0.891	24	-4.368	< 0.001
Med climate	2.764	0.601	24	4.595	< 0.0001
Fire hazard	2.769	1.118	24	2.476	< 0.05
Dense thickets	3.516	0.755	24	4.655	< 0.0001
Garden	4.570	0.785	24	5.820	< 0.0001
Environmental	5.154	0.721	24	7.150	< 0.0001
Congeneric	6.107	0.746	24	8.185	< 0.0001
Tolerates	2.651	0.730	24	3.629	< 0.001
Unintentionally	3.644	0.804	24	4.530	< 0.0001
Intentionally	2.140	0.907	24	2.360	< 0.05
Contaminant	2.240	0.758	24	2.956	< 0.01
Wind	2.040	0.846	24	2.412	< 0.05
Bird	3.632	0.816	24	4.451	< 0.001

A positive estimate value indicates a positive relationship with WRA score. See "Appendix 1" for explanation of variables

introduction, introduced as contaminant, and bird dispersed (Table 1). In contrast, we found that the WRA score was significantly related with more predictors: 13 out of the 15 (Table 2). These variables were mainly related to domestication, climate and area of distribution, undesirable traits, dispersal mechanisms, and persistence attributes.

Discussion

as potentially suitable for a wide range of geographical regions (reviewed by Gordon et al. 2008). We also obtained encouraging results when screening invasive species through the WRA scheme for Spain as an example for a Mediterranean region: 94% of invasive species were rejected. This result is similar to that for other regions (see "Results" for "major invaders" in Gordon et al. 2008). The results for casual species (considered here as non-invasive) were somewhat unexpected, 50% of them being rejected. As some of the inconsistencies when comparing different studies are due to different ways of defining the alien status (Gordon et al. 2008), we opted for a clear and accepted definition of "casual species" as alien species that do not reproduce outside cultivation and rely on repeated introductions for their persistence (Richardson et al. 2000a; Pyšek et al. 2004), avoiding any possible conceptual confusion. However, this terminology does not take into account the impact caused by these species as other studies do (Pheloung et al. 1999; Kato et al. 2006). According to the results for non-invasive species in other regions (ranking from 2 to 23% of species rejected, Gordon et al. 2008), our results for casuals are less accurate. This difference could be due to the fact that we did not include species not escaped from cultivation in the pool of non-invasive. Křivánek and Pyšek (2006) using the same terminology than in our study, found that only a 6% of casual woody species were rejected. Probably, they found lower rates of rejection for casuals because they only tested for woody species and therefore their plant sample was more homogeneous than ours.

Pheloung et al. (1999) designed the WRA score to be a precautionary method, rejecting all serious and most of minor invaders. It is better to make the mistake of denying introduction to a species that would not manage to survive outside cultivation than to allow the introduction of a harmful invader, because the consequences of an introduction are often irreversible (Daehler et al. 2004). From this point of view, our result of rejecting 50% of casual species would not be a result of low accuracy but an indication of the invasive potential of these species. However, after calculating the area under the ROC curve, we found that the WRA test applied in our dataset is less accurate in separating invasive from casual species than the test applied in other regions (Gordon et al. 2008). As mentioned above, our lower accuracy could be a consequence of not including species not escaped from cultivation into the pool of non-invasive or a consequence of several other nonexclusive explanations concerning the status of casual species. First, as mentioned, the species rejected could be in fact potential invaders with small residence time or with long lag phases, therefore, needing more time to pass from casual to naturalized and later to invasive status (Caley et al. 2008; Crooks 2005). As other authors have demonstrated, minimum residence time is one of the most important factors that should be considered in evaluating invasion success (Pyšek and Jarosík 2005). We did not have information on minimum residence time for casual species, so more research would be needed on this direction, because according to this hypothesis, the WRA scheme could be working as an early-warning mechanism for potential invaders. Another reason for our findings could be an inappropriate classification of species as casual in the Atlas of Invasive Plant Species in Spain (Sanz-Elorza et al. 2004) when they could be considered invasive. For example, Ligustrum lucidum, classified as casual, has recently experienced a high population growth in some localities near Barcelona (Gassó, unpublished data) and could be locally classified as invasive. Finally, another explanation for the large number of casual species misclassifications could be a consequence of the high weighting that the WRA gives to some variables that are not related with invasion status but to the potential impact of the species (Table 1). In fact, risk has two components: the likelihood of invasion which is related to invasion status and impact. Other studies have shown that quantitative risk evaluation systems only based on invasion likelihood would have almost the same performance as the current WRA (Caley and Kuhnert 2006).

Characteristics related to dispersal capacity are highly related to invasion status. Dispersal as a produce contaminant and bird dispersed species are prone to becoming invasive, as previous studies have demonstrated (Rejmánek and Richardson 1996; Richardson et al. 2000b; Lloret et al. 2005). On the other hand, intentional dispersal by humans is related to casual status. Species intentionally introduced in new regions (e.g., for agricultural purposes or as ornamentals) often manage to jump into natural ecosystems and survive; becoming casual species, but their persistence depends on the constant influx of more individuals. This result is in accordance with the intention of the first question of the WRA; "Is the species highly domesticated?" If answering "yes" to this question, the final score becomes lower (i.e., the more domesticated is a species, the lower the invasion risk). This assumption considers that plants selected and bred by humans for many generations to grow in a cultivated environment have a handicap that might limit the species survival in the wild because of their reduction of competitive traits (Daehler and Carino 2000). However, when competitive traits have been selected through domestication, we considered the answer to be negative. Moreover, a species that is highly domesticated might have a higher propagule pressure, and many studies have demonstrated that propagule pressure is one of the most important factors related to invasion success (Williamson 1996; Lockwood et al. 2005). For example, Pyšek and Jarosík (2005) found frequent planting to be correlated to invasive success, because the more times the species has been introduced the higher its probability of being successfully dispersed and spread. Thus considering propagule pressure, there is a positive relationship between intentional dispersal (question 7.02) and the final WRA score. Indeed, we consider that, in certain cases, there might be a contradiction between the question regarding intentional dispersal and the one addressing domestication (question 1.01). More research is needed to elucidate the effects of domestication and propagule pressure on invasion likelihood.

We obtained a conspicuous result concerning the difference between invasion likelihood and impact: the capacity of creating a fire hazard was negatively related to invasion status. Therefore, fire risk generation does not appear to be a factor that increases the probability of spread, as there are more species with fire generation capacity among casual species than among invasive. Nevertheless, the capacity of creating a fire hazard is positively related to the WRA score, because it is an undesirable trait, especially in the Mediterranean region where fire is an important inherent disturbance and plants with this capacity could cause major impacts. Therefore, because many plants currently classified as casuals, if spread in the future, could generate significant impacts, we can highlight the importance of including impact questions in a risk assessment. However, as the additive model to integrate probability and risk could result in misunderstandings (e.g., casual species with low probability to become invasive but a high potential impact such as increase of fire hazard), we suggest that specifications on the impact and likelihood components weight on the final score should be done, or even a separate analysis to evaluate them. This measure would provide clearer interpretation and more useful information for management and decision makers.

Finally, concerning the viability of applying the WRA scheme, we would like to stress that many questions require very specific information that in many cases, especially for casual species, has not been documented yet. Those species that are widely invasive over the world have been studied considerably, thus there is generally more information about invasive species than casual species. For some characteristics, information is lacking or is very difficult to obtain for most of the taxa even if there is clear evidence of their importance in successful invasions. Examples of such traits are hybridization (Vilà et al. 2000), allelopathy (Callaway and Ridenour 2004), release of natural enemies (Colautti et al. 2004), self compatibility (Daehler 1998), and potential impacts (e.g., pest and pathogens).

In conclusion, we found that the rate of correctly identifying invaders is high, and that it could be working as an early-warning mechanism for casual species with potential to become invasive. Moreover, having detected some incongruities between invasion

2.01

2.02

2.03

2.04

2.05

likelihood questions and impact questions, we suggest that a separate analysis should be done to evaluate these two risk components, in order to provide what would provide more useful information for management. Concerning invasion likelihood, more research is needed to determine if differences between casual and invasive are due to differences in biological traits, or whether it is a matter of propagule pressure and residence time. This study represents a first step towards the construction of a Mediterranean Basin WRA. However, more research is needed to evaluate the performance of the test in identifying non-invasive species.

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Med climate

Quality data

Climate suitability

Native med climate

Repeated intro

Appendix 1

See Table 3.

et al. (1999) (WRA) to our study, and abbreviations used in our analysis						
Category	WRA code	WRA question	Abbreviation			
Domestication/cultivation 1.01		Is the species highly domesticated?	Domestication			
	1.02	Has the species become naturalized where grown?	Naturalization			
	1.03	Does the species have weedy races?	Weedy races			

Species suited to Mediterranean climate

Broad climate suitability (environmental versality)

Native or naturalized in regions with extended dry periods

Does the species have a history of repeated introductions

Quality of climate match data

outside its natural range?

 Table 3 Table of correspondences between the adapted questions from the Australian weed risk assessment system of Pheloung et al. (1999) (WRA) to our study, and abbreviations used in our analysis

Climate and distribution

Tabl	le 3	continue	d

Category	WRA code	WRA question	Abbreviation
Weed elsewhere	3.01	Naturalized beyond native range	Nature native range
	3.02	Garden/amenity/disturbance weed	Garden
	3.03	Weed of agriculture/horticulture/forestry	Agriculture
	3.04	Environmental weed	Environmental
	3.05	Congeneric weed	Congeneric
Undesirable traits	4.01	Produces spines, thorns or burrs	Spines
	4.02	Allelopathic	Allelopathy
	4.03	Parasitic	Parasitic
	4.04	Unpalatable to grazing animals	Unpalatable
	4.05	Toxic to animals	Toxic
	4.06	Host for recognized pests and pathogens	Host pathogens
	4.07	Causes allergies or is otherwise toxic to humans	Allergies
	4.08	Creates a fire hazard in natural ecosystems	Fire hazard
	4.09	Is a shade tolerant plant at some stage of its life cycle	Shade tolerant
	4.10	Grows on infertile soils	Soil
	4.11	Climbing or smothering growth habit	Climbing
	4.12	Forms dense thickets	Dense thickets
Plant type	5.01	Aquatic	Aquatic
	5.02	Grass	Grass
	5.03	Nitrogen fixing plant	Nitrogen
	5.04	Geophyte	Geophyte
Reproduction	6.01	Evidence of substantial reproductive failure in native habitats	Reproductive failure
1	6.02	Produces viable seed	Viable seed
	6.03	Hybridizes naturally	Hybridizes
	6.04	Self-compatible or apomictic	Self-compatible
	6.05	Requires specialist pollinators	Specialist pollinator
	6.06	Reproduction by vegetative propagation	Vegetative
	6.07	Minimum generative time	Min time
Dispersal mechanisms	7.01	Propagules likely to be dispersed unintentionally	Unintentionally
1	7.02	Propagules dispersed intentionally by people	Intentionally
	7.03	Propagules likely to disperse as a produce contaminant	Contaminant
	7.04	Propagules adapted to wind dispersal	Wind
	7.05	Propagules buoyant	Water
	7.06	Propagules bird dispersed	Bird
	7.07	Propagules dispersed by other animals (externally)	Animals
	7.08	Propagules dispersed by other animals (internally)	Gut
Persistance attributes	8.01	Prolific seed production	Seed production
	8.02	Evidence that a persistent propagule bank is formed	Propagule bank
	8.03	Well controlled by herbicides	Herbicides
	8.03	Tolerates, or benefits from mutilation, cultivation or fire	Tolerates
	8.05	Effective natural enemies present locally	Enemies

Appendix 2

See Table 4.

Family	Species	Score	Family	Species	Score
(A) Invasive spec	ies				
Aceraceae	Acer negundo	13	Crassulaceae	Crassula lycopodioides	14
Agavaceae	Agave americana	14	Cyperaceae	Cyperus alterniformis flabelliformis	18
Aizoaceae	Carpobrotus acinaciformis	21	Elaeagnaceae	Elaeagnus angustifolia	21
Aizoaceae	Carpobrotus edulis	22	Fabaceae	Acacia dealbata	24
Amaranthaceae	Achyranthes sicula	12	Fabaceae	Acacia longifolia	23
Amaranthaceae	Amaranthus albus	10	Fabaceae	Acacia melanoxylon	21
Amaranthaceae	Amaranthus blitoides	10	Fabaceae	Acacia saligna	22
Amaranthaceae	Amaranthus hybridus	10	Fabaceae	Gleditsia triacanthos	10
Amaranthaceae	Amaranthus muricatus	12	Fabaceae	Leucaena leucocephala	21
Amaranthaceae	Amaranthus powelli	7	Fabaceae	Parkinsonia aculeata	15
Amaranthaceae	Amaranthus retroflexus	6	Fabaceae	Robinia pseudoacacia	15
Amaranthaceae	Amaranthus viridis	10	Fabaceae	Sophora japonica	12
Anacardiaceae	Schinus molle	4	Hydrocharitaceae	Elodea canadensis	20
Asclepiadaceae	Araujia sericifera	17	Iridaceae	Chasmanthe aetiopica	15
Asclepiadaceae	Asclepias curassavica	9	Iridaceae	Tritonia crocosmiiflora	1
Asclepiadaceae	Gomphocarpus fruticosus	9	Malvaceae	Abutilon theophrasti	15
Asteraceae	Achillea filipendulina	11	Myrtaceae	Eucalyptus camaldulensis	17
Asteraceae	Ageratina adenophora	13	Myrtaceae	Eucalyptus globulus	21
Asteraceae	Ambrosia artemisifolia	18	Nyctaginaceae	Mirabilis jalapa	7
Asteraceae	Arctotheca calendula	17	Onagraceae	Oenothera biennis	9
Asteraceae	Artemisia verlotiorum	7	Onagraceae	Oenothera glazioviana	8
Asteraceae	Aster squamatus	17	Oxalidaceae	Oxalis pes-caprae	24
Asteraceae	Conyza bonariensis	15	Poaceae	Bromus willdenowii	4
Asteraceae	Conyza canadensis	12	Poaceae	Cenchrus incertus	6
Asteraceae	Conyza sumatrensis	15	Poaceae	Chloris gayana	-3
Asteraceae	Cotula coronopifolia	10	Poaceae	Cortaderia selloana	26
Asteraceae	Helianthus tuberosus	7	Poaceae	Echinochloa hispidula	14
Asteraceae	Senecio inaequidens	23	Poaceae	Echinochloa oryzicola	17
Asteraceae	Senecio mikanioides	16	Poaceae	Echinochloa oryzoides	17
Asteraceae	Xanthium spinosum	19	Poaceae	Eleusine indica	8
Asteraceae	Xanthium strumarium	17	Poaceae	Paspalum dilatatum	18
Azollaceae	Azolla filiculoides	32	Poaceae	Paspalum paspalodes	20
Buddlejaceae	Buddleja davidii	19	Poaceae	Paspalum vaginatum	12
Cactaceae	Austrocylindropuntia subulata	11	Poaceae	Pennisetum setaceum	26
Cactaceae	Cylindropuntia spinosior	18	Poaceae	Sorghum halepense	24
Cactaceae	Cylindropuntia imbricata	21	Poaceae	Spartina alterniflora	14
Cactaceae	Opuntia dillenii	22	Poaceae	Spartina patens	10
Cactaceae	Opuntia engelmannii	22	Poaceae	Stenotaphrum secundatum	13
Cactaceae	Opuntia ficus-indica	22	Polygonaceae	Fallopia baldschuanica	15

Table 4 Species used to test if the Australian weed risk assessment system (WRA) of Pheloung et al. (1999) was suitable to predict(A) 100 invasive (A) and (B) 97 casual species in Spain. The final WRA score is given

Table 4 continued

Family	Species	Score	Family	Species	Score
Cactaceae	Opuntia humifusa	22	Polygonaceae	Fallopia japonica	15
Cactaceae	Opuntia monacantha	22	Pontederiaceae	Eichhornia crassipes	23
Cactaceae	Opuntia phaeracantha	22	Simaroubaceae	Ailanthus altissima	12
Cactaceae	Opuntia stricta	22	Solanaceae	Datura innoxia	15
Caprifoliaceae	Lonicera japonica	14	Solanaceae	Datura stramonium	19
Chenopodiaceae	Atriplex semibaccata	19	Solanaceae	Nicotiana glauca	14
Commelinaceae	Tradescantia fluminensis	12	Solanaceae	Solanum bonariense	19
Convolvulaceae	Ipomoea indica	14	Tropaeolaceae	Tropaeolum majus	14
Convolvulaceae	Ipomoea purpurea	21	Verbenaceae	Lantana camara	25
Convolvulaceae	Ipomoea sagittata	17	Verbenaceae	Lippia filiformis	15
Convolvulaceae	Ipomoea stolonifera	17	Zygophyllaceae	Zygophyllum fabago	14
(B) Casual species					
Agavaceae	Yucca aloifolia	4	Liliaceae	Hemerocallis lilioasphodelus	5
Aizoaceae	Lampranthus multiradiatus	12	Liliaceae	Ornithogalum arabicum	3
	Alternanthera sessilis	20	Liliaceae	Tulipa clusiana	3
Amaranthaceae	Amaranthus caudatus	18	Liliaceae	Tulipa gesneriana	-1
Amaranthaceae	Amaranthus tricolor	6	Malvaceae	Gossypium arboreum	13
Amaryllidaceae	Narcisus × medioluteus	6	Malvaceae	Hibiscus rosa-sinensis	-5
-	Catharanthus roseus	1	Malvaceae	Hibiscus syriacus	-6
	Monstera deliciosa	6	Moraceae	Morus nigra	4
Asteraceae	Ageratum houstonianum	7	Myrtaceae	Callistemon citrinus	-4
	Calendula officinalis	-6	Myrtaceae	Eucalyptus gunnii	11
	Cynara scolymus	0	Myrtaceae	Eucalyptus sideroxylon	14
	Gamochaeta subfalcata	9	Nymphaeaceae	Nymphaea mexicana	14
	Gazania rigens	12	Oleaceae	Jasminum nudiflorum	0
	Senecio cineraria	-1	Oleaceae	Ligustrum lucidum	4
	Solidago gigantea	10	Oleaceae	Ligustrum ovalifolium	9
	Tagetes patula	0	Oleaceae	Syringa vulgaris	-2
	Impatiens glandulifera	18	Onagraceae	Oenothera laciniata	- 17
	Doxantha unguis-cati	17	Passifloraceae	Passiflora caerulea	6
-	Jacaranda mimosifolia	-1	Pinaceae	Larix eurolepis	-4
-	Cereus peruvianus	3	Pinaceae	Pinus canariensis	14
	Allocasuarina verticillata	-3	Pinaceae	Pinus ponderosa	8
	Casuarina cunninghaniana	15	Pittosporaceae	Pittosporum tobira	3
	Beta vulgaris	-3	Poaceae	Arundo donax	8
-	Convolvulus mauritanicus	-3 4	Poaceae	Lolium multiflorum	8 10
		4 10	Poaceae		10
	Sedum sexangulare			Panicum capillare	
	Lagenaria siceraria	0	Polygonaceae	Fagopyrum esculentum	8
-	Cupressus macrocarpa	4	Polygonaceae	Fagopyrum tataricum	4
	Cyperus michelianus	20	Polygonaceae	Rumex maritimus	12
	Acacia decurrens	17	Polygonaceae	Rumex patientia	5
	Acacia mearnsii	17	Rosaceae	Cydonia oblonga	5
	Acacia pycnantha	24	Rosaceae	Photinia serrulata	-6
	Acacia sophorae	6	Rosaceae	Prunus persica	-4
Fabaceae	Acacia verticilata	13	Rosaceae	Prunus serotina	12

Family	Species	Score	Family	Species	Score
Fabaceae	Bauhimia grandiflora	6	Rosaceae	Pyracantha angustifolia	19
Fabaceae	Cassia obtusifolia	17	Rosaceae	Rosa moschata	8
Fabaceae	Lathirus odoratus	15	Rosaceae	Spiraea cantoniensis	1
Fabaceae	Lathyrus sativus	10	Rutaceae	Citrus limon	5
Fabaceae	Mimosa pudica	20	Salicaceae	Populus simonii	0
Fabaceae	Phaseolus lunatus	8	Salicaceae	<i>Populus</i> \times <i>canescens</i>	7
Fabaceae	Robinia hispida	20	Sapindaceae	Cardiospermum halicacabum	15
Geraniaceae	Pelargonium peltatum	-4	Solanaceae	Datura ferox	19.5
Iridaceae	Iris albicans	14	Solanaceae	Nicandra physaloides	4
Lamiaceae	Mentha spicata	1.5	Solanaceae	Nicotiana tabacum	5
Lamiaceae	Perilla frutescens	5	Ulmaceae	Ulmus pumila	12
Lamiaceae	Salvia microphylla	13	Verbenaceae	Aloysia citrodora	7
Liliaceae	Aloe arborescens	20	Verbenaceae	Lantana montevidensis	14
Liliaceae	Aloe vera	18	Verbenaceae	Verbena canadensis	-1
Liliaceae	Asparagus setaceus	4	Vitaceae	Parthenocissus inserta	11
			Vitaceae	Vitis berlandieri	1

 Table 4 continued

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