

Model ID	Reference citation	Study type	Name of the developed or applied model	Regions applied	Groups applied	Number of species or diseases assessed	Total number of questions	Question type	Weighting	Rank formation - brief explanation	Rank output	Uncertainty comments	Uncertainty L&E	Policy implemented (references at the bottom of the table)	T/E/A/S/I individual components estimated (yes/no)	Model components	Notes
1	Ahmed et al (1988)	Development	Ahmed et al	No application	Fish	0	26		Inbuilt to scoring	Sum of the questions	The species with the highest score is the most potential candidate	Not necessary to answer all questions, but really intended to be used if the question is not appropriate to the situation.	NNN	The authors had no information regarding the use of the model.	Y/Y/Y/N/Y	S: Species trait (8) Ig: Survival during transport (1) r: Reproductive potential (1) IK: Population growth (3) IL: Impact (2) Management (1) Other (A) use available (related to use an historical)	The paper is not about invasive species, but employs a similar approach, and is such an early approach that it was included in the sample.
2	Bomford & Glover (2004), Bomford (2008)	Further development	Australian freshwater fish model	Australia	freshwater fish	47	5	Different types of predictor variables (continuous, categorical) related to species and environmental traits	No	One rank	Low, moderate, serious, extreme	No	NNN	The Australian Federal Government (national level) uses the models to assist decision making when assessing applications to import new exotic vertebrate species into Australia. Several State Governments in Australia (for example the Western Australian Government) also use the models to assist decision making in relation to exotic vertebrate species.	N/Y/NNN	pr(1): Probability of establishment (5)	
3	Bomford (2003)	Development	Australian vertebrate model	Australia	mammals and birds (adapted to herpetiles and fish)	44 mammals / 60 birds	19	Different types of predictor variables (continuous, categorical) related to species and environmental traits	Inbuilt to scoring	Three different ranks formed	Low, moderate, serious, extreme: determined from the various combinations of the three risk scores	No	NNN	As above	N/Y/NN/Y	S: Species trait (2) pr(1): Probability of establishment (7) IL: Impact (10)	
4	Bomford (2005, 2008)	Further development	Australian reptile and amphibian model	Australia, New Zealand	reptiles and amphibians	UK: 42; California: 56; Florida: 68; AUS: 7	3	Different types of predictor variables (continuous, categorical) related to species and environmental traits	No	One rank	Low, moderate, serious, extreme	No	NNN	As above	N/Y/NNN	pr(1): Probability of establishment (3)	
5	Bomford (2008)	Further development	Australian bird and mammal model	Australia, New Zealand	mammals and birds (adapted to herpetiles and fish)	AUS: 45 mammals / 73 birds	20	Different types of predictor variables (continuous, categorical) related to species and environmental traits	Inbuilt to scoring	Three different ranks formed	Low, moderate, serious, extreme: determined from the various combinations of the three risk scores	No	NNN	As above	N/Y/NN/Y	S: Species trait (2) pr(1): Probability of establishment (8) IL: Impact (10)	
6	Branquart (2007)	Development	Invasive Species Environmental Impact Assessment Protocol (ISEIA)	Belgium	Selected species of several taxa (vascular plants, mosses, mammals, birds, fish, amphibians, reptiles)	15	4	All answers are scored on a 3-point scale	No	Sum of the four components, no weighting	High, moderate and low environmental risk. (Black list, watch list, no list)	Uncertainty is rated for each question. If not information, no score. If poor information, score 1 (unlikely) and 2 (likely)	N/Y/Y	Applied formally and informally in Belgium	N/Y/N/Y/Y	pr(1): Probability of establishment (1) ID: Dispersal (1) IL: Impact (2)	
7	Brunei et al. (2010)	Development	EPPO prioritization process for invasive alien plants	EPPO region	Plants	0	11	Yes - no (6) and three scaled responses (5)	No	Decision tree resulting in a matrix of scores combining three levels of impact with three levels of spread potential (phase 1), and decision tree with a three scored result (phase 2)	Phase 1: lists of species of minor concern, observation list and list of invasive alien plants, Phase 2: no, lower or priority for PRA	Phase 1: uncertainty rated for each question as low, medium or high, and one overall uncertainty rating	Y/N/Y	The EPPO system is formally applied to many alien species in the EU to justify regulations. Most widely applied to non-plant pests in Great Britain.	Y/Y/N/Y/Y	OL: Organisms at uptake (2) pr(1): Probability of establishment (5) ID: Dispersal (1) IL: Impact (3)	
8	Caley and Kuhnert (2008)	Application + Further development + Comparative test	Caley and Kuhnert TREE model (A-WRA)	Australia	Plants	370	4	Yes/No binary questions	No	Classification regression tree	Weed, non-weed	No	NNN	Not applied according to the authors.	Y/Y/NN/Y	N: Propagule pressure (1) pr(1): Probability of establishment (1) ID: Dispersal (1) IL: Impact (1)	
9	Champion and Clayton (2000)	Development	The New Zealand Aquatic Weed Risk Assessment Model (AWRAM)	New Zealand	Aquatic plants	21	36	Multiple-choice questions. Questions use different scales (0-10, 0-5, 0-3 and 0-1) depending on their level of importance.	Inbuilt to scoring	Sum of individual questions	Continuous ranking	No	NNN	Formally applied to decide on which aquatic species should be banned and also to select species for eradication programmes in New Zealand	N/Y/Y/Y/Y	S: Species trait (5) r: Reproductive potential (4) IL: Environment interactions (2) ID: Dispersal (4) IL: Impact (10) Management (6)	
10	Cook and Proctor (2007)	Development	Cook & Proctor	Australia	Plant pests and diseases	10	10		Explicit	Weighted sum of all sub-categories	Continuous ranking	Formed a 'risk index' based on variability of scores by panel members	Y/N/Y	The bioeconomic models that were used to provide expert testimony on the economic impact criterion, continue to be used informally to support decision-making processes in Australia. Used internally by government biosecurity agencies to allocate scarce funds to control pests. The bioeconomic model FISK has been used to inform government policy makers (Department of Environment, Food and Rural Affairs) in the UK as regards the potential invasiveness of non-native freshwater fishes when classifying fishes for regulation under the Import of Live Fish Act (ILFA). The outcome of FISK was taken into consideration when deciding on classifications, but it was not used to classify the species. FISK scores also are part of a decision support tool for assessing management options impacts, but has been applied in a post-prior manner only.	Y/NN/N/Y	N: Propagule pressure (1) IL: Impact (8) Management (1)	
11	Copp et al (2005)	Further development	Fish Invasiveness Screening KIT (FISK) / Invasive Fish Risk Assessment (IFRA) based on A-WRA/EPPO	UK	Fish	9	49	Central components (e.g. rank formation) of FISK are based on A-WRA	Inbuilt to scoring	Equally-weighted sum of answered questions grouped in two sections, with total score compared against a set of critical values that determine the risk of the species to become invasive. Minimum number of answered questions (10/49)	Accept, evaluate (=need further evaluation), reject taxon	No	NNN		Y/Y/Y/Y/Y	S: Species trait (10) N: Propagule pressure (2) v: Transport vectors (3) OL: Organisms at uptake (1) pr(1): Probability of establishment (5) r: Reproductive potential (6) IK: Population growth (5) IL: Environment interactions (2) ID: Dispersal (1) IL: Impact (12) Management (1) Other (1: quality of the information)	
12	Copp et al (2005)	Further development	Fish Invasiveness Screening KIT (FISK) / Invasive Fish Risk Assessment (IFRA) based on A-WRA/EPPO	UK	Fish	1	34	Central components (e.g. rank formation) of IFRA based on EPPO. Use of scoring matrices for questions in which two or more variables contribute to the assessment. Explanatory guides provided with each question, and the assessor expected to provide a brief rationale for the score given for each question	No	Sum of the sums for each category	Continuous ranking	Suggest a basic approach to calculate the proportion of 'don't know' answers. Alternatively, qualify each answer on scale of confidence (1-3): low, medium; high	Y/N/Y		Y/Y/Y/Y/Y	N: Propagule pressure (1) OL: Organisms at uptake (7) OL(Tg): Transport (3) Ig: Survival during transport (1) pr(R): Probability of release (1) pr(1): Probability of establishment (7) ID: Dispersal (1) IL: Impact (10) Management (2) Other (1: quality of the information)	
13	Cowie et al (2009)	Development	Quarantine list of Cowie et al (2009)	USA	Non-marine snails and slugs	46	12	3-point scale (0, 0.5, 1) according to a series of thresholds established by the authors	No	Question scores were summed for each species or group, and then divided by the total number of attributes scores	Continuous ranking from 0 to 1 (least to greatest concern)	None	NNN	The study was a request of US Department of Agriculture, Animal and Plant Health Inspection Service, Plant Protection and Quarantine (USDA-APHIS-PPQ)	Y/Y/Y/N/Y	S: Species trait (1) OL: Organisms at uptake (1) pr(1): Probability of establishment (4) r: Reproductive potential (3) ID: Dispersal (1) IL: Impact (2)	
14	Cunningham et al (2004)	Further development	A-WRA + staged assessment	Australia	Plants	17	49 + 6	Yes/no, some [0-2], [1-3], [1-4], [1-5]	No	Additive ranking within categories, but division between potential impact and feasibility of eradication categories to obtain benefit-cost ratio. Weed risk and benefit-cost ratio not combined. B-C ratio of the species ranking highest in WRA to be observed	Continuous ranking	No, although there is one question in the potential impact category related to data reliability	NNN	Unknown	N/Y/Y/N/Y	pr(1): Probability of establishment (1) r: Reproductive potential (1) OL: Species range (2) IL: Impact (1) Management (1)	The mapping includes only questions in addition to the A-WRA. The framework is not entirely clearly explained, and therefore difficult to map precisely.
15	Daehler and Castro (2000)	Application + Further development + Comparative test	Daehler and Castro (2002) SAF	Hawaii (USA)	Plants	111	19	Different types of predictor variables (continuous, categorical) related to species and environmental traits	No	No ranking. High/low risk determined by expert-decision tree system	Low or high risk	Not necessary to answer all questions. If no answer, higher risk path assumed (system explicitly conservative)	NN/Y	Unknown	N/Y/Y/Y/N	S: Species trait (1) pr(1): Probability of establishment (6) r: Reproductive potential (5) IK: Population growth (2) ID: Dispersal (5)	
16	Daehler et al (2004)	Application + Further development + Comparative test	HP-WRA+2nd screening	Hawaii (USA)	Plants	192	49+1		Inbuilt to scoring	Higher than 6 is pest, lower than 1/0 not pest, in between (1-4) the second screening process determines whether the species is a pest or requires further evaluation	Accept, evaluate (=need further evaluation), reject taxon	Not necessary to answer all questions	NN/Y	The assessment system is currently promoted by the Hawaii Department of Agriculture, by the Hawaii Invasive Species Council, and by Hawaii Industry groups (e.g. Landscape Industry Council of Hawaii). The assessments are made by a specialist who is paid by the State of Hawaii. While the risk assessment is endorsed and encouraged by many groups, it is NOT required by law and there is no law that restricts the growth or importation of species that are ranked as 'high risk'. Instead, the risk assessment is usually related to native conservation of	Y/Y/Y/Y/Y	S: Species trait (7) N: Propagule pressure (2) v: Transport vectors (7) pr(1): Probability of establishment (6) r: Reproductive potential (9) IL: Environment interactions (2) IL: Impact (16) Management (1)	

17	Darin et al. (2011)	Development	WHPPET (Weed Heuristics: Invasive Population Prioritization Tool)	North America (California)	Plants	19	25		Explicit	Score multiplied by weight, first to produce criteria value, and then the same for total score	Continuous ranking	Uncertain information was assigned to middle points so as not to affect the results	NNN	Used by the San Francisco Bay Area Early Detection Network to prioritize hundreds of occurrences of 73 target species. Also used by the San Diego County Department of Agriculture to prioritize regionally invasive weed infestations for control/containment/eradication. Has also been taken up by United States Forest Service, United States Fish and Wildlife <i>For more information, see the following link:</i>	NNY/Y/Y	r(.): Reproductive potential (5) X: Population size (1) D(.): Dispersal (8) I(.): Impact (2) Management (7)
18	Department of Primary Industries (2008)	Development	Victorian WRA	Victoria, Australia	Weeds	2	42		Explicit	Weight multiplied by score and the scores then summed up	The species with the highest score is the most potential candidate	Each question scored for uncertainty	NNY	Applied in the State of Victoria, Australia	Y/Y/Y/Y/Y	S: Species trait (2) v: Transport vectors (1) pr(1): Probability of establishment (3) r(.): Reproductive potential (3) D(.): Population growth (4) D(.): Dispersal (1) I(.): Impact (27) Other (Current range / potential range) O(.): Organisms at uptake (2) I(.): Survival during transport (2) pr(R.): Probability of release (3) pr(1): Probability of establishment (13) r(.): Reproductive potential (1) D(.): Dispersal (3) Q(1): Species range (1) I(.): Impact (16) Management (6) Other (1 - adaptability (is the pest highly adaptable? 1 - transient populations)) v: Transport vectors (1) pr(1): Probability of establishment (2) r(.): Reproductive potential (1) D(.): Dispersal (1) Q(1): Species range (1) I(.): Impact (8) Management (1) Other (1 - facilitation by climate change)
19	EPPO (2011)	Development	EPPO computer-assisted pest risk assessment scheme (CAPRA)	Europe, North Africa, Great Britain	Plant pests including weeds	0	48	All answers are scored on a 5-point scale (3-point for impact).	No	No explicit guidance on how to combine scores of individual questions to a final score.	No ranking	Uncertainty rated as low, medium & high for each question	NNY	Formally applied to many alien species in the EU to justify regulations. Most widely applied to non-plant pests in Great Britain.	Y/Y/N/Y/Y	r(.): Reproductive potential (1) D(.): Dispersal (3) Q(1): Species range (1) I(.): Impact (16) Management (6) Other (1 - adaptability (is the pest highly adaptable? 1 - transient populations)) v: Transport vectors (1) pr(1): Probability of establishment (2) r(.): Reproductive potential (1) D(.): Dispersal (1) Q(1): Species range (1) I(.): Impact (8) Management (1) Other (1 - facilitation by climate change)
20	Essl et al. (2011)	Development	GABIS	Germany, Austria	Many taxa (fish)	31	16	Criteria are scored on a 3 to 4-point scale.	No	Ranking is based on the highest assessment criteria score, no weighting. Precautionary approach in assigning to groups.	High (=Black List), intermediate (=Grey List), low risk (=White List)	Uncertainty rated low, med, high for each question	NNY	Applied informally, but expected to influence decisions for introduction/release of species in Germany	N/Y/Y/Y/Y	pr(R.): Probability of release (1) I(.): Impact (6)
21	European Technology Platform for Global Animal Health (2006)	Development	ETPGAH	Europe	Animal diseases	30	7		No	Probably summation, but not explained.	The diseases divided into three groups (major diseases, diseases for surveillance, and neglected zoonoses)	No	NNN	Unknown	Y/NN/N/Y	pr(R.): Probability of release (1) I(.): Impact (6)
22	Fajcz et al. (2006)	Development	Bosnia and Herzegovina animal health model	Bosnia and Herzegovina	Animal diseases	7	8		Explicit	Weight multiplied by score and the scores then summed up		No	NNN	Unknown	NNN/N/Y	I(.): Impact (6) Management (3)
23	Garry Oak Ecosystems Recovery Team (2007)	Development	IAPP Species Scoring Algorithm	Canada	Plants	None	24	Yes/no	Explicit	Weighted sum	Continuous output	No	NNN	Unknown	N/Y/Y/Y/Y	S: Species trait (7) v: Transport vectors (1) pr(1): Probability of establishment (2) r(.): Reproductive potential (10) I(.): Impact (2) Management (2)
24	GB Non-native species secretariat (2011)	Further development	GB Non-native species Rapid Risk Assessment (NRA)	Great Britain	Invasive species		17	The first 7 questions and the management questions are open, the 8 scored questions are five scaled and comments to justify score need to be provided	No	No explicit guidance on how to combine scores of individual questions to a final score.	Conclusion from questions and summarising categories drawn by risk assessor and peer review	Scores in the summarising categories and the final concluding risk assessment are accompanied by a five scaled confidence score	NNY	Officially used in the UK. Have completed 46 full risk assessments and 1 rapid risk assessment. In addition have 34 full risk assessments in progress, with 21 rapid risk assessments in progress.	Y/Y/N/Y/Y	N: Propagule pressure (1) pr(1): Probability of establishment (5) D(.): Dispersal (2) Q(1): Species range (1) I(.): Impact (2) Management (4) Other (1 - facilitation by climate change + 1 - overall risk)
25	Gederaas et al. (2007)	Development	Norwegian Black List	Norway	Selected species of several taxa	217	Phase 1: 3 Phase 2: 4	Multiple choice questions	No	Phase2: high: if at least one question is answered with yes to any of the impacts, low: if all four questions are answered with no, unknown: if no answer is yes and at least one cannot be answered because of uncertainty	Phase 1 identifies species as not requiring further assessment. Phase 2 classifies species in 3 risk categories (low, high, unknown risk)	Phase1: if uncertain, need to proceed to phase 2, Phase 2: all questions have an "unknown" category	NNY	Officially used in Norway. Each species is strictly classified on the basis of the set of criteria alone. For management, the system is used informally; management decisions (by DN) are based on the risk assessment of alien species, but may also be informed by other aspects (such as economic constraints and political priorities). So far applied to about 4000 alien species documented in Norway. Was used by the Australian Government Department of Agriculture Fisheries and Forestry in defining next pest lists used to guide monitoring and management regimes. Current status unknown.	N/Y/N/N/Y	pr(1): Probability of establishment (2) I(.): Impact (6)
26	Hayes and Siliva (2003)	Development	Australian potential marine pests model	Australia	Marine invasive species	851	4		No	Did not actually rank them, only distinguished them as potential pests	Checks if survive all the criteria, if yes, identified as pests	Excluded if uncertainty	NNN		Y/NN/N/Y	v: Transport vectors (2) Q(1): Species range (1) I(.): Impact (1)
27	Habert and Stubbendorf (1993)	Development	Exotic Plant Species Ranking System	USA	Plants	71	25		No	Sum of components, control reported separately	Matrix with two dimensions (impact and control)	No	NNN	Unknown	NNN/Y/Y	S: Species trait (1) pr(1): Probability of establishment (2) r(.): Reproductive potential (4) I(.): Environment interactions (1) D(.): Dispersal (1) Q(1): Species range (2) I(.): Impact (4) Management (10)
28	IPPC/FAO (2004)	Development	ISPM11	World	Plant pests	Many	Only guidelines	Only guidelines	No guidelines	No guidelines	No guidelines	Uncertainty should be documented for transparency	NNY	International plant trade. Risk assessment procedure acknowledged by the World Trade Organization.	Y/Y/N/Y/Y	v: Transport vectors (8) r(.): Reproductive potential (8) I(.): Environment interactions (2) I(.): Impact (4)
29	Jefferson et al. (2004)	Application + Further development + Comparative test	A-WRA, modified R-H, shortened a-WRA	Chicago (USA)	Plants	40	24	Yes/No binary questions; magnitudes not considered.	Inbuilt to scoring	Equally-weighted sum of answered questions. Minimum number of answered questions (10/40)	Accept, evaluation (=need further evaluation), reject tacit	Number of questions answered used as indicator of reliability	NNY	Chicago Botanic Gardens have an invasive plant policy that states that any plant that we import from outside the country and that is new to the region (i.e. not already common in the nursery industry) must undergo risk assessment. They use a modified version of the Reichard/Hamilton risk assessment framework. Risk assessment is conducted by the <i>Confidentiality: Please do not publish collection data</i>	NNY/Y/Y	N: Propagule pressure (1) pr(1): Probability of establishment (1) r(.): Reproductive potential (2) D(.): Population growth (1) I(.): Environment interactions (1) D(.): Dispersal (2) Q(1): Species range (2) I(.): Impact (6) Management (6)
30	Johnson (2009)	Development	Southern Australian Weed Risk Management System	New South Wales, Australia	Plants	0	22	Multiple-choice questions using different scales.	No	Questions posed in separate sections to determine weed risk and feasibility of control. Weed risk calculated by averaging the scores per section and then calculating: invasiveness x impact x potential distribution. Weed risk and Control feasibility form then a matrix	Very high, high, medium, low, negligible risk.	Index based on number of unanswered questions	Y/N/Y	Applied formally regionally in Australia (New South Wales)	Y/Y/Y/Y/Y	S: Species trait (5) N: Propagule pressure (1) v: Transport vectors (3) pr(1): Probability of establishment (4) r(.): Reproductive potential (5) I(.): Environment interactions (1) I(.): Impact (21) Management (1)
31	Koop et al. (2012)	Application + Further development + Comparative test	PPQ (Plant Protection and Quarantine) WRA, based on A-WRA	USA	Plants (non-invaders, minor-invaders and major-invaders)	204	41	Yes/No binary questions, although a there are a few multi-choice questions; magnitudes not considered.	Inbuilt to scoring	Equally-weighted sum of answered questions at PPQ WRA. However, species categorized as "involute" further follow a secondary screening model (i.e. a short decision tree, similar to IP-WRA/2nd screening)	Low risk, evaluate further, high risk	Number of questions answered used as indicator of reliability. Guidance provided on how to answer questions and assessments reviewed by two team members.	Y/N/Y	Unknown	Y/Y/Y/Y/Y	S: Species trait (5) N: Propagule pressure (1) pr(1): Probability of establishment (4) r(.): Reproductive potential (5) I(.): Environment interactions (1) I(.): Impact (21) Management (1)

Note on mapping: it was not entirely clear from the paper which questions were included in the shortened WRA and the authors were not available for comment.

32	McKenzie et al (2007)	Development	Rapid Risk Analysis based on IRA by OIE	New Zealand	Animal pathogens	82	12		No	risk: $\text{risk} = \text{Pb} \times \text{Lb} \times \text{Lob}$, ranked by target population (wildlife, etc.). Also all populations can be considered by summing the individual target	Continuous ranking (also by target population)	If information not available, worst case scenario used. Significant gaps in knowledge were recorded.	Y/N/N	Has not been applied in policy.	Y/Y/N/N/Y	N: Propagule pressure (1) p(RI): Probability of release (5) RI: Impact (6)
33	Miller et al (2010)	Development	Miller et al Relative Risk Model	Nebraska (USA)	Plants	18	23	Magnitudes, converted to 4 categories (0-absent, 2-low, 4-medium, 6-high) in increments of 1/10th (from 0-absent to 1-high). Yes/no questions at the Effects filter category	No	Multiplication of individual criteria, after adding score values of question subcategories (i.e. 17 land cover categories in Habitat rank, and 4 characteristics in Effect filter)	Continuous ranking	Present median values of uncertainty distributions; assess each question by uncertainty (1=low, 2=medium, 3=high depending on data reliability)	N/N/Y	Authors have no information on any policy application.	N/Y/N/N/Y	OL: Organisms at uptake (1) p(II): Probability of establishment (18) RI: Impact (4)
34	Minnesota Department of Natural Resources (1991)	Further development	Exotic Plant Species Ranking System for Minnesota, based on Hebert	Minnesota (USA)	Plants	153	23		Inbuilt to scoring	Sum of components, control reported separately	Threat category: minimal, moderate, severe, unknown	No		Unknown	N/N/Y/Y/Y	S: Species trait (1) p(II): Probability of establishment (1) r: Reproductive potential (4) RI: Environment interactions (1) DL: Dispersal (1) QT: Species range (4) RI: Impact (2) Management (9)
35	More et al (2010)	Development	Irish animal health prioritization, based on Animal Health Strategy of the EU	Ireland	Animal diseases	13	1	One question scored on a 5-point scale	No	One criterion	[1-5]	No	NNN	The framework was applied formally as part of the decision-making process. The framework formed the basis for prioritization of activities within Animal Health Ireland, a relatively new organization with responsibility for national coordination and facilitation of all issues relating to non-indigenous animal health.	NNN/N/Y	RI: Impact (1) Other (after initial prioritization, details on impact and management are collected, but no ranking is made)
36	Morse et al. (2004)	Development	NatureServe's Invasive Species Assessment Protocol	USA	Plants	0	20	Four scaled responses plus unknown category.	Inbuilt to scoring	Answers are assigned points that are summed up for the four categories of the scheme and then combined to a final biodiversity impact score. If there is a range of answers, then min and max added separately.	Produces an I-Rank of either "high", "medium", "low", or "insignificant".	"Unknown" category in multiple choice style of four scaled responses included, if no precise answers possible, ranges can be chosen and are accounted for in the rank formation	N/N/Y	According to the paper, used to assess biodiversity impact of about 3,500 plants established outside cultivation in the USA	N/N/Y/Y/Y	p(II): Probability of establishment (3) r: Reproductive potential (1) RI: Population growth (1) DL: Dispersal (1) QT: Species range (6) RI: Impact (5) Management (4)
37	Nentwig et al. (2010)	Development	Generic Impact-Scoring System	Europe	Invasive species (mammal, birds)	34	10	All answers are scored on a 5-point scale.	Explicit	Weighted sum of scores, multiplied by proportional area.	Continuous impact ranking	Variability of scores by experts used	Y/N/Y	Applied informally to derive a black list of alien animals in Switzerland	NNN/N/Y	RI: Impact (10)
38	OIE (2011a,b)	Development	OIE codes	World	Animal pathogens	Many	Only guidelines	Only guidelines	No guidelines	Although no specific guidance is given, it is made clear that the individual invasion steps should be seen as a pathway, i.e. earlier steps need to be overcome in order to evaluate later steps.	No guidelines	For quantitative risk assessments: probability distributions or confidence intervals for overall uncertainty, sensitivity analysis	N/Y/Y	International animal trade. Risk assessment procedure acknowledged by the World Trade Organization.	Y/Y/N/Y/Y	S: Species trait N: Propagule pressure v: Transport vectors tg: Survival during transport p(II): Probability of establishment v: Population size DL: Dispersal RI: Impact Other (risk of contamination (release), management (at each step))
39	Oleinik et al. (2007)	Development	Bioinvasion Index	Baltic Sea	Invasive aquatic species	14	5	Impact questions scored on a 5-point scale, but abundance and distribution ranges on a 3- and 4-point scales respectively	No	Matrix combining three levels of impact with abundance and distribution ranges of species	Bioinvasion Level on a scale 0 (weak) to 4 (massive).	No	NNN	Applied informally in Lithuania (and possibly will be applied in Finland) for implementation of marine strategy directive	N/N/Y/Y/Y	RI: Population growth (1) QT: Species range (1) RI: Impact (3)
40	Ou et al. (2008)	Development	Chinese WRA	China	Plants	67	19	Questions structured hierarchically (primary vs. secondary indices) and scored into a continuous scale (from 0 to 100) based on the Analytic hierarchy process (AHP).	Explicit	Weighted sum of six categories (or primary indices)	Species ordered based on their invasion risk	No	NNN	According to the paper, has been applied since March 2007 in Xiamen.	Y/Y/Y/Y/Y	N: Propagule pressure (3) p(II): Probability of establishment (2) r: Reproductive potential (1) DL: Population growth (1) RI: Environment interactions (1) QT: Species range (4) RI: Impact (3) Management (4)
41	Parker et al. (2007)	Development	U.S. Weed Ranking Model	USA	Plants	249	27 (authors say 33)	Multiple-choice questions using different scales (ranging between 0-10 or 0-1 depending on the category)	Explicit	Product of the four categories, the category scores are sums of individual questions/scores. Two categories (geographic potential and entry potential) score 0-1, while the category damage potential ranges between 0.8 and 2.4 and invasive potential (the critical element of the model) can attain scores >30	Continuous score	Quality of information was ranked; model tested statistically / Use of correlations and sensitivity analysis to determine the influence of categories and factors in the final model scores, as well as comparison of these values to independent scores from previously tested scoring systems	N/N/Y	Was applied to revise plants-for-planting regulations, but also to help select species for early detection of adventive pests. The model has since been superseded by the agency's formal weed risk assessment process	Y/Y/Y/Y/Y	S: Species trait (6) N: Propagule pressure (1) p(II): Probability of establishment (3) r: Reproductive potential (4) DL: Dispersal (1) QT: Species range (1) RI: Impact (10) Management (1)
42	Pheloung 2001	Development	Hazard model	Australia	Plants	0	12		Inbuilt to scoring	Summation		No	NNN	Used in Australia by AQIS until 1997 for decision-making on the importation of plants.	N/Y/Y/Y/Y	S: Species trait (1) p(II): Probability of establishment (1) RI: Population growth (2) DL: Dispersal (4) RI: Impact (4)
43	Pheloung et al (1999)	Development	Australian WRA (A-WRA)	Australia	Plants	370	49	Yes/No binary questions; magnitudes not considered.	Inbuilt to scoring	Equally-weighted sum of answered questions. Minimum number of answered questions (10/49)	Accept, evaluation (needed further evaluation), reject (taste)	Number of questions answered used as indicator of reliability	N/N/Y	Applied formally to thousands of species in Australia and New Zealand	Y/Y/Y/Y/Y	S: Species trait (8) N: Propagule pressure (2) v: Transport vectors (7) p(II): Probability of establishment (5) r: Reproductive potential (8) RI: Environment interactions (2) RI: Impact (15) Management (1) Other (1: quality of data)
44	Reichard and Hamilton (1997)	Development	Reichard and Hamilton scheme (R-H)	North America	Plants	235	8	Different types of predictor variables (continuous, categorical) related to species' traits.	No	Statistical analysis (classification and regression decision trees)	Accept (low probability of invasiveness), reject (high probability), further analysis/monitoring needed (intermediate probability)	No	NNN	Used to reform the US risk assessment process, but also applied informally by non-governmental organizations. It has been used by a number of botanic gardens, for instance University of Washington, Chicago Botanical Garden and Missouri Botanical Garden. Also some nurseries have used it, which uses the individual use of the model as a tool to	N/Y/Y/Y/N	p(II): Probability of establishment (4) r: Reproductive potential (3) DL: Dispersal (1)
45	Risk Assessment and Management Committee (1996)	Development	The Generic Nonindigenous Aquatic Organisms Risk Analysis Review Process	USA	Aquatic species	0	7		Inbuilt to scoring	In establishment: if any sub-rating is low, then rating is low; in impact: if any sub-rating is high then impact rating is high. Final score is average, but rounded to a conservative side	Matrix providing a summary score of low, med or high	Each question scored for uncertainty	Y/N/Y	Unknown	Y/Y/N/Y/Y	v: Transport vectors (1) tg: Survival during transport (1) p(RI): Probability of release (1) DL: Dispersal (1) RI: Impact (3)
46	Sæther et al. 2010; Sandvik et al. In revision	Development	Criteria System for Ecological Risk Assessment of Alien Species in Norway	Norway	Invasive species	Assessments for 5 species are finished	4	Varies (y/N, numbers)	Inbuilt to scoring	Matrix of invasion / spread vs. effect axis	5 categories: severe impact, high impact, potentially high impact, low impact, no known impact	Not formally included, but uncertainty should be considered by applying the "precautionary principle"	NNN	Will be applied to all alien species in Norway as basis for IAS management	N/Y/N/Y/Y	p(II): Probability of establishment (1) DL: Dispersal (4) RI: Impact (1)
47	Smallwood and Salmon (1992)	Development	Smallwood and Salmon Rating System	USA	Birds and mammals	48	40	Yes/no and multiple-choice questions using different scales. (*) When multiple choices referred to the same component, each choice was considered a separate component in the fourth column.	Explicit	Weighted sum of four section scores, calculated by summing the scores assigned to each component of the section or selecting the highest ones.	Continuous ranking	Correction value attached to uncertain information (0.05 to rating with questionable information / 0.10 to rating with poor or non-documented information)	N/N/Y	Unknown	Y/Y/N/N/Y	S: Species trait (3) v: Transport vectors (2) tg: Survival during transport (1) p(II): Probability of establishment (2) RI: Impact (19) Management (13)

48	SZEID (2006)	Development	AHW Prioritisation Decision Support Tool	UK	Animal diseases	0	40	Multiple choice questions/criteria	Explicit	Weighted sum of criteria/question, independently of the section/category in which each criteria/question belongs to.	Continuous ranking	Development of the tool with the help of stakeholders and risk/epidemiology experts, as well as few questions in the questionnaire related to availability of data	N/N/Y	Has been applied informally in the UK. D2K2 outputs have been applied in an informal manner to support the decision-making process. D2R2 outputs (including graphs and tables to show the relative rankings) have been produced to inform the monthly Veterinary Risk Group meetings, the selection of diseases for International Disease Monitoring reports, and resource planning and allocation within Delta.	v: Transport vectors (1) O ₁ : Organisms at uptake (1) pr(1): Probability of establishment (1) ID ₁ : Dispersal (3) QT ₁ : Species range (3) R ₁ : Impact (21) Management (6) Other (1 - data availability / 3 - economics)
49	Tucker and Richardson (1996)	Development	Expert System for screening potentially invasive alien plants in South African fynbos (SAF)	South Africa	Woody plants	73	24	Different types of predictor variables (continuous, categorical) related to species and environmental traits	No	No ranking. High/low risk determined by expert-decision tree system	Low or high risk	Not necessary to answer all questions. If no answer, higher risk path assumed (system explicitly conservative)	N/N/Y	Applied informally in regulations for South Africa's "biodiversity act" risk assessment for invasive species	N/Y/Y/N pr(1): Probability of establishment (8) r ₁ : Reproductive potential (8) ID ₁ : Population growth (2) ID ₂ : Dispersal (5)
50	Ward et al (2008)	Development	Invasive Art Risk Assessment	New Zealand	Arts	11	32	All answers scored on a 3-point scale (0, 0.5, 1).	No	Question scores averaged by category, and the averages summed to produce an overall score. If two species had same score, one with higher consequences first	High, medium, low risk	No	NNN	Applied formally in New Zealand by the Ministry for Agriculture and Forestry and Biosecurity NZ to assess what art species would be invasive in New Zealand.	Y/Y/Y/N/Y v: Transport vectors (2) O ₁ : Organisms at uptake (4) O ₁ /Tg ₁ : Transport (2) pr(1): Probability of establishment (8) r ₁ : Reproductive potential (1) e ₁ : Carrying capacity (1) R ₁ : Impact (12) Management (2)
51	Warner et al (2003)	Development	CAL-IPC, based on Invasive Species Assessment Protocol	Arizona, California and Nevada (USA)	Plants	0	13		Inbuilt to scoring	Uses tables, where all answer combinations are listed		Not necessary to answer all questions; documentation is assessed based on reliability, but does not affect the rank but can be used to describe degree of confidence in ranking	N/N/Y	According to the paper, applied informally in California, Nevada and Arizona	Y/Y/N/Y/Y v: Transport vectors (1) pr(1): Probability of establishment (4) r ₁ : Reproductive potential (1) ID ₁ : Dispersal (2) QT ₁ : Species range (1) R ₁ : Impact (4)
52	Weber and Gut (2004)	Development	Classification key for Neophytes (WG)	Central Europe	Vascular plants	240	12	Multiple-choice questions with different scales, but always ranging between 0 and 4	Inbuilt to scoring	Sum of all individual questions	High, intermediate, low risk	No	NNN	Unknown	N/Y/Y/Y/Y S: Species trait (1) pr(1): Probability of establishment (4) r ₁ : Reproductive potential (2) ID ₁ : Population growth (1) ID ₂ : Dispersal (1) R ₁ : Impact (2) Other (1 - habitat vulnerability)
53	Weber and Gut (2005)	Development	Weed Survey	Europe	Plants	281	3	Short questions (e.g. "weediness")	Explicit	First listed by most important criterion, then by second criterion		Used the average of several respondents, but in the model as such no uncertainty included	N/N/Y	Unknown	NNN/Y/Y ID ₁ : Dispersal (1) R ₁ : Impact (1) Management (1)
54	Widrichner et al (2004)	Further development + Comparative test	modified R-H, R-H/matrix model, New CART model	Iowa (USA)	Plants	100	3	Different types of predictor variables (continuous, categorical) related to species' traits.	No	No ranking, high risk/low risk is determined in a hierarchical decision tree	Accept (low probability of invasiveness), reject (high probability), further analysis/monitoring needed (intermediate probability)	No	NNN	Iowa Department of Agriculture & Land Stewardship (IDALS) plans to use the models the next time that state weed regulations are modified. Chicago Botanic Gardens has applied the models informally as part of their internal decision-making process.	pr(1): Probability of establishment (1) r ₁ : Reproductive potential (1) ID ₁ : Dispersal (1)
55	Widrichner et al (2004)	Further development + Comparative test	modified R-H, modified R-H/matrix model, New CART model	Iowa (USA)	Plants	100	10	Different types of predictor variables (continuous, categorical) related to species' traits.	No	No ranking, high risk/low risk is determined in a hierarchical decision tree	Accept (low probability of invasiveness), reject (high probability), further analysis/monitoring needed (intermediate probability)	No	NNN	As above	N/Y/Y/Y/N pr(1): Probability of establishment (5) r ₁ : Reproductive potential (3) ID ₁ : Dispersal (2)
56	Andreu and Vilà (2010)	Application + Comparative test	A-WRA, WG	Spain	Plants	80								Agreement between the competent authorities species have been listed to an Act on Management on Invasive species in the	
57	Barney and Ditomasso (2008)	Application	A-WRA	USA	Biofuel plants	3									
58	Buddenhagen et al (2009)	Application	HP-WRA	Hawaii (USA)	Plants	80								Unknown	
59	Copp et al (2009)	Application	FISK (with uncertainty and predictive power improvements)	UK	Fish	67	49	Central components (e.g. rank formation) of FISK are based on A-WRA		Equally-weighted sum of answered questions grouped in two sections, with total score compared against a set of critical values that determine the risk of the species to become invasive. Minimum number of answered questions (10/49)	Accept, evaluation (=need further evaluation), reject taxon	Each species scored by two people, difference denotes a delta score. Each response scored for uncertainty on scale 1-4	Y/N/Y	Applied in the UK as a part of the decision-making process regarding the Import of Live Fish Act.	
60	Crosti et al (2010)	Application	A-WRA	Italy	Plants	20								Has not been applied in policy.	
61	Dawson et al (2009)	Application	HP-WRA+2nd screening	Tanzania	Plants	230									
62	Gascó et al (2010)	Application	A-WRA	Spain	Plants	197								Has not been applied in policy.	
63	Gordon et al (2008)	Application + Comparative test	A-WRA, HP-WRA+2nd screening, Caley and Kuhnert TREE model	Florida (USA)	Plants	158								Has been applied informally in Florida. The tool has been incorporated into the University of Florida's IFAS Assessment of Non-native Plants in Florida, which is the basis for extension faculty recommendations for which species to	
64	Kato et al. (2006)	Application	HP-WRA+2nd screening	Japan, Bonin Islands	Plants	130									
65	Křivánek and Pyšek (2006)	Application + Comparative test	A-WRA, HP-WRA+2nd screening, R-H	Czech Republic	Plants	180									
66	Massam et al (2010)	Application	Australian bird and mammal model	Western Australia	Invasive species	40	20			Two different ranks formed (matrix)		Based on number of references available			
67	Nishida et al (2009)	Application + Comparative test	A-WRA, HP-WRA+2nd screening	Japan	Plants	259									The Japanese government has used it applied to decide Invasive Alien Species by "Invasive Alien Species Act". However, the Invasive Alien Species
68	Pheloung et al (1999)	Application	NZ-WRA	New Zealand	Plants	291	49	Yes/No binary questions; magnitudes not considered.		Equally-weighted sum of answered questions. Minimum number of answered questions (10/49)	Accept, evaluation (=need further evaluation), reject taxon	Number of questions answered used as indicator of reliability	N/N/Y	Applied formally to thousands of species in Australia and New Zealand	
69	Randall et al (2006)	Application	Invasive Species Assessment Protocol	USA	Plants	0	20			Subranks are summed up, if range of answers, then min and max added separately	Produces an I-Rank of either "high", "medium", "low", or "insignificant".	Can answer unknown (U), can answer ranges	N/N/Y	The Department of Agriculture and Forestry is using the protocol to assess the estimated 3,500 nonnative vascular plant species that are established in the United States to create a national list	

70	Tricarico et al (2010)	Application	Freshwater Invertebrates Scoring Kit (FISK), based on FISK	Italy	Invertebrates (crayfish)	37	49	Yes/No/Don't know questions, with a level of certainty (spread over four rankings)	Sum of the section scores	Total scores range from -4 to 39, but were evaluated in light of the score thresholds used in the original A-WRA and FISK and a statistical computation (ROC)	Each response scored for uncertainty on scale 1-4 (according to the Intergovernmental Panel on Climate Change)	N/N/Y
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ID manuscript	Citation	Groups applied	Number of species assessed	Regions applied	Uncertainty examined	Model description	T/E/A/S/I component & estimated (yes/no)	Mapping	Mapping comments
1	Ahrens et al. 2011	Plant herbicide resistant (<i>A. stolonifera</i> , <i>A. gigantea</i> , <i>A. perennans</i> , <i>A. scabra</i> , <i>A. capillaris</i> , <i>A. Canina</i>)	6	USA	Yes (cross validation and kappa)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
2	Allen et al. 2001	Molluscs (mussel)	1	USA & Canada	Yes (accuracy)	Logistic regression on traffic, hydrology and water chemistry as predictors for <i>Dreissena</i> occurrence	N/Y/N/N/N	$Z_i = pr(\lambda_c E_i, V_i)$	
3	Andrew & Ustin 2010	Plants	1	USA (California)	No	Dispersal model that examines the role of propagule pressure and landscape structure on rate of spread	Y/Y/N/Y/N	$Z_{i,j} = N_{i,j} = \sum_{j=1}^J \sum_{k=1}^K f_D(\bar{E}, D_{i,j})$	Dispersal dependent on environment in pathways (corridors too), as well as general habitat suitability
4	Aragon & Lobo 2012	Insect (maize pest western corn rootworm <i>Diabrotica virgifera</i> ssp. <i>virgifera</i>)	1	Northern hemisphere	Yes (misclassification rates)	Species distribution models combined with physiological spatial projections outputs (geographical projections of temperature physiological thresholds based on key fitness components, MDE, and MD categorized as lower or higher climatic favourability) to estimate areas of invasion under both current and future climate scenarios	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
5	Arriaga et al. 2004	Plant (buffel grass, <i>Cenchrus ciliaris</i> L.)	1	Mexico	Yes (kappa)	Species distribution model derived from using climate, habitat and edaphic data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
6	Ba et al. 2010	Crustacean (decapode)	1	Global	No	Modelling invasion risks on a global scale using SDM. To assess invasion risks in major ports, dispersal distances and major ocean shipping routes were additionally accounted for.	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i, D_{i,j})$	Dij is the distance from known source and nearest coastline. Verbally indicated included it into SDM
7	Barney & DiTomaso 2010	Plant (switchgrass, <i>Panicum virgatum</i>)	1	USA	No	Bioclimatic envelope models used to estimate the potential distribution and suitable habitat based on the climate and distribution data in the native range under both current and future climate scenarios	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
8	Bartell & Nair 2004	Insect (Asian longhorned beetle, <i>Anoplophora glabripennis</i>)	1 species/1 pathway (establishment); the study mention other transitions like transport and spread, but they are not really modeled	USA	Uncertainties inherent to the estimation of model parameters that determine the risk of establishment are defined, quantified, and propagated through the population model. The approach takes into account a wide array of factors influencing establishment: (1) factors that determine the rate of pest entry into the United States; (2) the biology and ecology of the pest species; (3) the availability of suitable hosts and environmental factors that influence pest establishment in different geographical regions; (4) the population dynamics of the pest species; and (5) the implications of uncertainties on resulting estimates of risk and risk reduction.	The risk of establishment is modelled considering both the rate of entry and the population size required for persistence (i.e., threshold). Risk of establishment is modelled with a stage-based demographic model, including information on the biology and ecology of pest species, the suitability of potentially susceptible hosts, and the quality of available habitats	N/Y/Y/N/N	$X_{t+1} = f_x(X_t, r + \sigma_x)$ $Z = P_i^A = pr(X_t > threshold)$	Stage based population model more complex than demographic model mapped. Assume demographic stoch, but not entirely sure. No density dependence. Not clear whether there were any environmental variables modeled
9	Beaumont et al. 2009	Plants (hawkweeds: <i>Hieracium pilosella</i> , <i>H. aurantiacum</i> and <i>H. muronum</i>)	3	Australia	Yes (AUC and kappa)	Ecological niche models combining occurrence data with various climate variables to predict current and future species distributions	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
10	Bomford et al. 2010	Freshwater fishes	280	Global	Yes (AUC using cross-validation)	Establishment success of species based on the number of countries where introductions occurred (a measure of propagule pressure) and species' genus and family. Climate matching was also tested.	N/Y/N/N/N	$Z = P^* = f_p(E, S, N, v)$	
11	Bomford et al. 2009	Reptiles and amphibians	1995 introduction records (1028 successful and 967 failed introductions) for 596 alien species	World, but emphasis on Great Britain, California and Florida	Yes (AUC using cross-validation)	Statistical models examining the association of establishment success and range expansion with environmental conditions, range size and history of establishment	N/Y/N/Y/N	$Z = P^* = f_p(E, S)$ $Z = Q = f_D(E, S)$	E is climate match. Also looked at hierarchical models, and therefore components of stochasticity. Not shown explicitly in mapping. Also examined multiple jurisdictions for validation
12	Bouchier & Van Hezewijk 2010	Plant (Japanese knotweed, <i>Polygonum cuspidatum</i>)	1	Canada	Yes (efficiency test with independent data)	Climate matching model on predicted the occurrence of invasive species	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
13	Bradley 2009	Plant (cheatgrass, <i>Bromus tectorum</i>)	1	Western USA	Yes (model prediction comparison)	Bioclimatic model using occurrence data was constructed based on Mahalanobis Distance. Future invasion risk considered	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
14	Bradley & Mustard 2006	Plant (cheatgrass, <i>Bromus tectorum</i>)	1	USA (North-central Great Basin)	Yes (validation with field data)	Occurrence of species compared to various spatially explicit landscape variables (elevation, aspect, hydrographic channels, cultivation, roads, and power lines) to assess risk of future invasion	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i, f(O, D))$	Distance to road, cultivated areas here were included as disturbance rather than related to human transport, and thus are in E. f(O,D) is the distance from a source population of cheatgrass (1973 distribution)
15	Bradley et al. 2010	Plants (kudzu, <i>Pueraria lobata</i> ; privet, <i>Ligustrum sinense</i> and <i>L. Vulgare</i> ; cogongrass, <i>Imperata cylindrica</i>)	3	South-East USA	Yes (AUC)	Climatic habitat using occurrence data was developed using both the Maxent and Mahalanobis distance methodologies. Future invasion risk considered	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
16	Bradley 2010	Plants	1	USA	Yes (probability models)	Hierarchical risk modeling on the probability of invasion by Cheatgrass on Sagebrush	N/Y/N/Y/Y	$f_i(E_i) = pr(\lambda_{c2}^{Native} E_i)$ $Z = \sum_{i=1}^I pr(\lambda_{c2} E_i) f_i(E_i)$	Also used ensemble models, and mis-classification rates
17	Bradshaw et al. 2008	Fabaceae plants	8906	World	No	Statistical model that examines the relationship between the ecological and life-history traits and the threat and invasiveness risk of species	N/Y/N/N/N	$Z = P^* = f_p(S)$	Although environment is mentioned, it is more a feature of the species than the receiving environment per se (not spatially explicit)
18	Brickman 2006	Various	0	Canada	Yes (statistical parameter distribution, SE etc)	Introduces a RA model for ballast water dispersion (which is the carrier of transported IAS) after release in seas, including a) time integrated ballast water concentrations; b) time need for an organisms to reach an area; and c) the onshelf average concentrations.	N/N/N/Y/N	$Z_{i,j} = N_{i,j} = \sum_{j=1}^J \sum_{k=1}^K f_D(\bar{E}, D_{i,j})$	fd - from oceanographic model. Ballast water layer captured in Vj across space. N - abundance reaching location i. Also looked at time to reach area, but not included in mapping
19	Broennimann et al. 2007	Plant (spotted knapweed; <i>Centaurea maculosa</i>)	1	North America	Yes (AUC using independent data)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
20	Brooks et al. 2012	Plants	1	Argentina (native range) & USA (Florida, alien range)	Yes (statistical parameter distribution, SE etc)	Assessment of the role of the environmental envelope as compared with patterns of host-herbivore associations based on collections made in the native range.	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	

21	Buckley et al. 2005	Plants	1	New Zealand	Yes (sampling from parameter distribution)	Dispersal model incorporating life-history parameters	N/N/Y/N/Y	$X_{j,t+1} = f_X(X_{j,t}, r(E_{j,t}, S))$ $N_{i,t} = \sum_{j=1}^J f_{XD}(X_{i,t}, D_{i,j})$	Simplification - paper had stage structured population model, and had both long and short distance dispersal kernels, but ideas remain similar. Population rate of change (r) was dependent on species traits, and on grazing
22	Carrasco et al. 2010a	Insect (maize pest western corn rootworm <i>Diabrotica virgifera</i> ssp. <i>virgifera</i>)	1	Austria	Yes (maximum likelihood and predicted versus observed regression)	Spatially explicit metapopulation model (integrating both natural and human-assisted dispersal) and phenology model (driven by degree-days)	Y/Y/N/Y/Y	$N_{i,t} = \sum_{j=1}^J f_D(E_i, E_j, D_{i,j})$ $Z = Q_i = \sum pr(\lambda_c E_i) * pr(N_{i,t} > 0) * f_{attractiveness}(\text{environment})$	Included both natural and human mediated dispersal, using pop sizes as attractiveness (environment) or locations (gravity models affecting transport). Indicated population submodel, but no info on it, except to indicate effect of degree day
23	Carrasco et al. 2010b	Multiple (bacterium, insects)	3	United Kingdom	No	Bioeconomic model that considers the exclusion, detection and control of multiple NIS spreading by stratified dispersal and presenting Allee effects	N/Y/N/Y/Y	$P^{(t)} = f(M^{(t)})$ $P_{i,t+1}^{(t)} = 1 - (1 - pr(\lambda_{A_i}))^{N_{i,t}^{(t)}}$ $Q_T = \sum_{i=1}^I f(P_{i,t}^{(t)})$ $Z = \sum P_i^{(t)} * P^{(t)} * (Q_T * f_i(\alpha^* M^{(t)}) + M^{(t)})$ $P_i^{(t)} * (1 - P^{(t)}) * (Q_T * f(\cdot) + M^{(t)} + M^{(t)}) = (1 - P_i^{(t)}) * (M^{(t)} + M^{(t)})$	Also examined detection (Pe). From description in paper, not clear whether QT increased as a function of t, or increased after establishment. Also, included efficacy of control, but this was also not stated in paper
24	Carrasco & Baron 2010	Mollusc (Pacific oyster, <i>Crassostrea gigas</i>)	1	South American coast	No	Occurrence Data, Species distribution model combining occurrence data with surface seawater and atmospheric temperature	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
25	Carrete & Tella 2008	Birds	202	Spain	Yes (standard errors)	Life trade model that uses species origin (wild-caught versus captive-bred), availability on international markets and potential phylogenetic effects to explain invasion success	Y/N/N/N/N	$Z = N = f(O(S))$	Here, O is the potential for uptake, as measured by the number of stores carrying particular species ("popularity") or the number being sold. Propagate pressure was some function (f) of popularity response variable - invasion success - not clear which component, although we treat this as establishment. Also, looked at env traits, but from the perspective of tolerances of species. Thus, did not include E as a predictor
26	Castro-Díez et al. 2011	Trees (Australian acacias)	85	Australia	Yes (models compared with AIC)	Study of predictors of invasiveness among a set of 85 Australian Acacia species	N/Y/N/N/N	$Z = P^* = f_p(S, f(V))$	
27	Catford et al. 2011	Plants	441	Australia	Yes (survey gap analysis using the multivariate environmental similarity surface (MESS) algorithm; CV correlation, AUC, Deviance)	Occupancy and abundance based on environmental, biotic and propagate pressure predictors analysed by boosted regression trees	N/Y/N/N/N	$Z_i = f_{X_i}(E_i, f(V))$ $Z_i = f_{X_i}(E_i, f(V))$	Didn't look explicitly at abundance occurrence, but could do so. IXs denotes abundance of cover of all exotic species. Vproxy is a surrogate (e.g., human population size) of traffic to a location. In using it here, does not consider temporal dynamics or sources
28	Catterall et al. 2012	Plants	1	United Kingdom	Yes (posterior distributions of parameter estimates)	Habitat suitability and dispersal model built within a Bayesian framework	Y/Y/N/Y/N	$N_{i,t} = \sum_{j=1}^J f_D(D_{i,j}); Z = Q_i = \sum pr(\lambda_c E_i) * pr(N_{i,t} > 0)$	
29	Chen 2008	Insect (Yellow crazy ant, <i>Anoplolepis gracilipes</i>)	1	Global	No	Species distribution models calibrated using occurrence data and past, current and future climate data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
30	Chown et al. 2012	Plants	>100	Antarctica	No	Not a risk model per se but a quantification of propagate pressure in different climate environments	Y/Y/N/N/N	$Z = pr(\lambda_c E) \sum_v N_v$	Env is degree days, and propagate pressure is separated by vector types (v)
31	Chytrý et al. 2008	Plants	Many (> 1000)	Europe	Yes (sensitivity analysis)	Potential level of invasion (% neophytes in local communities) based on extrapolation from average occurrences at different habitat types	N/Y/N/N/N	$Z_i = f_i(E_i)$	Looking essentially at whether environment is suitable. Risk given per environment. Not sure how to deal with fraction of total species. Define Is as # of species. Sum across i - is it more insightful to keep separate?
32	Claudi & Ravishanker 2006	Not specified (everything in ballast water)		Canada	Yes (sensitivity analysis)	Probabilistic model to assess the risk of alien species introductions via ballast water, including journey duration and mortality of species	Y/N/N/N/N	$Z_{i,j} = \sum_j V_{i,j} f_g(t - t', \alpha^p M^p)$	
33	Cohen et al. 2007	Aquarium plants	138	Montreal, Canada	Yes (stochasticity and parameter uncertainty using a Bayesian approach)	Live trade-release model that estimates propagate pressure by means of Bayesian analysis. It uses a multinomial distribution to calculate the uncertainty distribution associated with all combinations of disposal pathways.	Y/N/N/N/N	$Z = N = \sum_{j=1}^J O_j(S) * pr(R S)$	Takes into account number of owners (J), number owned per individual (O), weighted by species (S) popularity, and release pr(R S) based on percent of individuals who release, and how many are released as a function of species traits. This model estimated total releases into the wild in a year, but does not consider environmental or spatial issues, which should affect probability of survival
34	Colnar & Landis 2007	Crustacean (crab)	1	USA	Yes (Monte Carlo analysis)	Ranking system for identifying exposure and impact on target taxa but does not include local spread	Y/N/N/N/Y	$N_{i,t} = \sum_v O(E_i, v) f_g(\cdot)$ $Z = \sum_i f(N_{i,t}) + \sum_{j=1}^J f_j^2(E)$	Not sure about this / Also took into account life stages - transitions between life stages and spread between patches not made clear. Considered impact on different ecosystem attributes. Conceptual model
35	Colunga-García et al. 2009	Various	0	USA (to global)	Yes (statistical parameter distribution; SE etc)	Analysis of regional freight transport to determine the likelihood of establishment of exotic species	Y/N/N/N/N	$Z = N = \sum_{j=1}^J f(V_{i,j,t})$	V actually incorporates multiple stages of transport (multiple vectors), which isn't shown in the mapping.
36	Compton et al. 2012	Plants	4	New Zealand	Yes (statistical parameter distribution; AUC, SE etc)	Statistical model examining whether variables that indirectly describe weed spread via human access and use, as well as a lake's position in the landscape, could describe the distribution of species	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i, f(V))$	Determine relative risk at different locations f(V) includes pop density and roads, integrated with environments to determine establishment
37	Cooke & Hill 2010	Fish	2	North America (Great Lakes)	Yes (statistical parameter distribution; SE etc)	Bioenergetic models that relate the metabolic requirements of invaders (under various body sizes, swimming speeds and reproductive stages) to planktonic food resources and environmental temperature to predict when and where they may survive	N/Y/N/N/N	$Z = \sum_{i=1}^I pr(\lambda_{c2} E_i, S)$	Based on bioenergetic model - not linear statistical model like most others examining E,S simultaneously

38	Copp et al. 2010	Fish	18	England	Yes (statistical parameter distribution; SE etc)	Spatial relationships between the occurrence of non-native fishes and human demographic factors associated with propagule pressure	Y/N/N/N/N	$Z = N = f(O(v), f(V))$	Here, treat O as stores (sources of species) and V proxy as human density. This is partly to remain consistent with previous mappings (e.g., Table 2). Also, looked at multiple sources (pathways?)
39	Coredeeli et al. 2010	Crustacean (copepods)	1	Pacific (coast of North America)	Yes (statistical parameter distribution, SE etc; model comparison approach, sensitivity analysis)	Statistical model that assesses areas at risk of invasion by examining the influence of various parameters on species occurrence	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
40	Costello et al. 2007	Crustacean (copepods)	Many (alien fauna & flora)	USA (San Francisco Bay)	Yes (statistical parameter distribution, SE etc, and by separating introduction and detection history)	Analytical model linking exotic species introductions and discoveries to trade volumes used to assess if trade origin and history affects new IAS numbers. The model is estimated using a novel historical data set on global trade and species introductions by region	Y/N/N/N/N	$Z_i = P_{i,t}^A = f\left(\sum_{j=1}^{J_i} (V_{i,j,t})\right)$	Examined probability of establishment as a function of trade volume from different areas
41	Cotner & Schooley 2011	Mammals	1	USA (Illinois)	Yes (statistical parameter distribution; AIC, SE etc)	Distribution model examining the relationship of site occupancy with local habitat conditions and anthropogenic landscape alterations	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
42	Crossman & Bass 2008	Tree (<i>Olea europea</i>)	1	Australia	Yes (AUC, missclassification measures)	Habitat suitability models applied to a single invasive plant	N/Y/Y/N/N	$Z = \sum_{i=1}^I f_X(E_i) pr(\lambda_i E_i)$	Also looked at Bayesian for uncertainty, and did ensemble models. As with others, did not actually combine both density and occurrence predictions, but could have
43	Crossman et al. 2011	Weeds	33	Australia	Yes (AUC, missclassification measures)	Spatially-explicit indicator of threat to biodiversity from invasive plants under human-induced climate change by combining both a pattern-based element and a process-based element (dispersal).	Y/Y/N/Y/Y	$V_{i,j,t} = V_{j,t} f_D(\vec{E}_i, \vec{D}_j, E_i, D_{i,j})$ $N_{i,t} = \alpha \sum_{j=1}^J (V_{i,j,t})$ $Q_t = \sum_{i=1}^I (1 - pr(\lambda_A))^{N_{i,t}}$ $Z = Q_t * f_f(S)$	Used a constant for propagule source sizes. Invasion history used for impact f(S). Also looked at climate change effects (not included in equation). Converted components to relative scores. In actual paper, summed across species s, to obtain a multi-species predictions per location i
44	Dangles et al. 2008	Insect (moths)	3	Ecuador	Yes (standard errors, and spatial and temporal variation)	Distributed delay models on temperature effect on life history development and demography. GLM on the effect of climate and host type on occurrence and abundance	N/N/Y/N/N	$Z = X = \sum_{i=1}^I f_X(E_i)$ $Z = X_i = f_X(X_{i-1}, r(E_{i,t}), k)$	Looked at 2 studies - experimental which basically gave population dynamics parameters, and SDM - abundance. Pop dynamics model more complex (life stages) than the one in box2
45	Davis et al. 2011	Plants	1	USA	No	WRA vs. demographic model in relation to habitat suitability. No propagule pressure or dispersal	N/N/Y/N/N	$Z = X_{i,t+1} = f_X(X_{i,t}, r(E_{i,t}), \sigma)$	Stochasticity included but not distinguished between environmental and demographic. E reported as disturbances, but seemed to be based on management (e.g., herbicide treatments). Also had seedbank and stages, not included in mapping
46	Dawson et al. 2009	Plants	142	Tanzania	Yes (sensitivity analysis)	Trait-based model that also examines role of propagule pressure on tree survival, regeneration and spread	N/Y/N/N/N	$Z = P^C = f(S, E, N, t, v, D)$	Also more finely divided establishment. t is the time that the first plot was planted, D is the minimum distance between plot and forest. Not sure whether summed across establishments (i.e., estimate Q). Also looked at linear distance moved, which could be considered Q (wave-front - not sure whether multiple plots were considered)
47	De Meyer et al. 2008	Insects (Mediterranean fruit fly, <i>Ceratitis capitata</i> ; Natal fruit fly, <i>Ceratitis rosa</i>)	2	Africa, southern Europe, and world	Yes (binomial tests of correct prediction of both presences and absences)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
48	Denslow et al. 2008	Plants	Many (174-892/island)	Pacific islands	No	Linear regression models on geographic (environmental, socioeconomic) predictors to alien plant richness in island	N/Y/N/N/N	$Z_i = f_i(E_i, V_{prop})$	V based on a surrogate of human population attributes, which may affect the number of vectors transporting aliens
49	DeVaney et al. 2009	Eurasian cyprinid fishes (Common carp, <i>Cyprinus carpio</i> ; Tench, <i>Tinca tinca</i> ; Grass carp, <i>Ctenopharyngodon idella</i> ; and Black carp, <i>Mylopharyngodon piceus</i>)	4	North America	Yes (independent data)	Species distribution models calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
50	Diez et al. 2012	Plants	>25,000	New Zealand and Australia	Yes (Bayesian posterior distributions)	Hierarchical Bayesian approach using informed prior information from Australia to derived probabilities of naturalisation in NZ	N/Y/N/N/N	$Z = P^C = f(S)$	Here S is actually establishment (naturalization) in other locations (Australia) as a predictor of establishment in NZ. Priors are also implicit in calculations of probabilities (but not shown in mapping)
51	Dorcas et al. 2011	Reptile (Burmese pythons, <i>Python molurus bivittatus</i>)	1	USA	No	Species distribution model	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
52	Drake 2005	Mammal (feral nutria, <i>Myocastor coypus</i>)	1 pathway (establishment) / 1 species	United Kingdom	Yes (model selection by least squares, AIC and BIC)	Demographic models describing the lag phase of the population, where demographic stochasticity and Allee effects play a major role in extinction; ignores expansion and saturation phases	N/Y/Y/N/N	$X_{i,t+1} = f_X(X_{i,t}, r, \kappa, \beta, \sigma_r, \sigma_d)$ $Z = P_t^A = pr(X_{i,t} > threshold)$	Considered Allee effects (Beta) in population dynamic model. Considered invasion if population size exceeded a threshold (1000)
53	Drake & Lodge 2004	Aquatic species	+1	Global	Yes (uncertainty examined using hindcasts)	Propagule pressure-establishment model that identifies global hotspots of invasion based on worldwide patterns of ship traffic and estimates rate of port-to-port invasion	Y/Y/N/Y/N	$V_{i,j,t} = V_{j,t} f_D(\vec{E}_i, \vec{D}_j, E_i, D_{i,j})$ $N_{i,t} = \alpha \sum_{j=1}^J (V_{i,j,t})$ $P_{i,t}^A = 1 - \prod_{i=1}^T (1 - pr(\lambda_A))^{N_{i,t}^{C_i}}$ $Z_T = Q_T = \sum_{i=1}^I P_{i,t}^A$	Uses vector traffic as a surrogate of propagule pressure, and calculates establishment as a function of propagule pressure. Implicitly assumes all propagules have equal probability and that no interaction occurs with environment (compare with full model, "establishment"). Note, J refers to all invaded sources, rather than all sources
54	Drake et al. 2005	Marine plankton	None	None	Yes (uncertainty indirectly associated with parameters)	Function of diffusive spread and population growth that estimates safe release thresholds	N/N/N/Y/N	$Z = P^C = f(\Delta X_i(S), \Delta Q)$	Function of diffusive spread and pop growth, which is functions of species traits. Establishment occurs if critical density exceeded, given pop growth and spread rate

55	Duggan et al. 2006	Aquarium fishes	308	Canada, USA	Yes (standard errors)	Live trade model that uses occurrence frequency in aquarium stores to explain introduction and establishment of aquarium fishes in freshwater habitats	Y/N/N/N/N	$Z = N = f(O(S))$	Here, O is the potential for uptake, as measured by the number of stores carrying particular species ("popularity") or the number being sold. Propagule pressure was some function (f) of popularity
56	Epanchin-Niell et al. 2009	Weeds (<i>Bromus tectorum</i>)	1	USA	Yes (sensitivity analysis)	Vegetation dynamic model to predict vegetation changes and management costs under different intensities and types of post-fire revegetation. Results estimate species dominance	N/N/Y/N/Y	$X_{i+1} = f_x(X_i, E_i)$ $E_{i+1} = f_e(X_i, E_i)$ $Z = f(E, M^c)$	Space not considered. Uses probability transition matrix between states (E), and has more states than presented in equations, but dynamics would be similar to interactions between species
57	Essi et al. 2010	Plants	86	Global (Temperate and subtropical countries and regions (n = 60) from five continents spanning both hemispheres)	No	Uses forest usage and forestry area across 60 global regions as a measure of possible "propagule pressure" and maps this on to regional environmental suitability.	N/Y/N/N/N	$Z = P^c = f(E, S, v)$	S is origin of species, E is regions area, and v is use in forestry
58	Fabregas et al. 2010	Animals	1 pathway (introduction)	Spain	No	Risk of potential animal escape at zoological parks by assessing the security of enclosures	Y/N/N/N/N	$Z = N = O(S) * pr(R S)$	O is the number of enclosures given S (taxonomic groups), and pr(R S) is the proportion of unsecured enclosures
59	Ficetola et al. 2007	Amphibian (American bullfrog, <i>Rana catesbeiana</i>)	1	Global	Yes (AUC and r^2)	Species distribution model calibrated with occurrence data, included hunting pressure	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
60	Fitzpatrick et al. 2012	Insect (hemlock woolly adelgid, <i>Adelges tsugae</i>)	1	Eastern North America	Yes (AUC)	Dynamic model using both abundance and dispersal data	N/Y/Y/Y/N	$X_{i,j+1} = f_x(X_{i,j}, r(E_{i,j}), \kappa(E_{i,j}), \sigma_e)$ $N_{i,j} = \sum_{j=1}^J f_{SD}(X_{i,j}, S, \sigma_{SD}) f_D(D_{i,j})$ $P_{i,j}^{A2} = 1 - (1 - pr(\lambda_{A2} E_i))^{N_{i,j}}$	Environment for k and IA was host levels, whereas for i it was winter temp. Stochasticity was incorporated at all levels. For pop dynamics, we called it environmental stochasticity, since per capita rates did not decrease with population size. Stochasticity in dispersal occurred in number of dispersers, not in the dispersal kernel
61	Floerl et al. 2005	Various (alien hull fouling species)	Many	New Zealand	Yes (statistical parameter distribution; SE etc)	Quantitative risk screening techniques that predicts the abundance and variety of organisms being transported by ocean-going yachts	N/N/N/N/Y	$Z_{i,j,k,t} = N_{i,j,k,t} = f(M^p, t'-t)$	Risk of vector k, traveling from i to j in time t, is a function of prevention (e.g., paint) and of time. In this case time is voyage history. Provides a metric of propagule pressure, but does not consider release converted to ranks
62	Floyd et al. 2006	Weeds	8	USA	Yes (ANOVA using different site categories, with and without management)	Each soil series and each vegetation community was ranked using an ordinal scale 1-3 representing low to high density of invasive non-native species, based on soil analyses	N/Y/N/N/N	$Z_i = f_{S_i}(E_i, M^c)$	
63	Follak & Strauss 2010	Plant (Horse nettle, <i>Solanum carolinense</i>)	1	Central Europe and North America	Yes (model predictions evaluated with independent data)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
64	Follak 2011	Plant (kudzu, <i>Pueraria lobata</i>)	1	Switzerland, Austria and Slovenia and parts of northern Italy	Yes (visual inspection and using independent data)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
65	Forsyth et al. 2004	Mammals	40	Australia	Yes (standard error applied to the fitted model)	GLM considering both occurrence and various explanatory variables (climate matching, life-history traits, introduction efforts) on the introduction and spread of species	N/Y/N/N/N	$Z = P^c = f_p(S, N)$	
66	Foxcroft et al. 2007	Plants	231	South Africa	Yes (r^2)	Risk assessment that alien species invade a protected area from surrounding watersheds; include species abundance or, as a proxy, species richness as a method of incorporating propagule pressure	N/Y/N/N/N	$Z = \sum_i \sum_{j=1}^J pr(\lambda_{c,i} E_i)$	Summed across species for a given location, or range for each species. Different than ones that use species richness as a response variable
67	Frank et al. 2008	Fungi (rust)	1	USA	Yes (spatial and temporal variation)	Synoptic classification meteorological data from upper (geopotential height, specific humidity, and u- and v-wind components) and surface (humidity and temperature) level used to simulate establishment likelihood	Y/Y/N/Y/N	$N_{i,j} = \sum_{j=1}^J f_D(\bar{E}, D_{i,j}, v)$ $Z_i = f(N_{i,j}) + f(pr(\lambda_B E_{i,j}))$	paper good for pr(B). Converted to scores, and then summed to determine risk according to date. Weird way of going about it
68	Fujisaki et al. 2010	Reptiles	68	USA	Yes (error rates)	Discriminant analysis, logistic regression and recursive partitioning and regression trees on species traits, live release, manageability and environmental variables to predict establishment success	N/Y/N/N/N	$Z = P^c = f(S, E, f(N))$	Authors discuss manageability, but mapped as species traits (e.g., aggressiveness) rather than related than as a management component (M)
69	Gassó et al. 2009	Plants	106	Spain	Yes (explained variance)	Joint species trait-environment model that examines occurrence data based on plant attributes and plant richness on environmental factors	N/Y/N/N/N	$pr(\lambda_{c2} E_i, S) = f(E_{i,1} + E_{i,2}... + S_{i,1} + S_{i,2})$ $Z = \sum_{i=1}^I pr(\lambda_{c2} E_i, S)$	incorporates both species traits and environmental suitability, simultaneously. Has only been done using linear models. Standard errors also provided, but not on validation set
70	Gertzen & Leung 2011	Crustacean (Spiny water flea, <i>Bythotrephes</i>)	1	Canada	Yes (stochasticity (probabilistic spread and establishment), epistemic uncertainty (missing data, modeled via simulation))	Extension of Leung et al. (2004). Extensions were to deal with unknown times of invasion, imperfect knowledge of invasion status, and multiple vectors.	Y/Y/N/Y/N	$V_{i,j,t} = V_{j,t}(v) f_D(\bar{E}, \bar{D}_i, E_i, D_{i,j}, v)$ $N_{i,j} = \sum_{j=1}^J V_{i,j,t}$ $P_{i,t}^A = 1 - \prod_{i=1}^I (1 - pr(\lambda_{A,i}))^{N_{i,t}^A}$ $Z_T = Q_T = \sum_{i=1}^I P_{i,t}^A$	Extension of Leung et al. (2004). Note that v was for fluvial dispersal and boater movement (combined into a single dispersal equation for simplicity, but could have had 2 equations)
71	Gertzen et al. 2008	Aquarium fishes	252	Montreal, Canada	Yes (stochasticity and parameter uncertainty using a Bayesian approach)	Life trade-release model that models propagule pressure using Bayesian analysis. The model was then extended to take into account specific characteristics and population size (species-specific propagule pressure)	Y/N/N/N/N	$Z = N = \sum_{j=1}^J O_j(S) * pr(R S)$	Takes into account number of owners (J), number owned per individual (O), weighted by species (S) popularity, and release pr(R S) based on percent of individuals who release, and how many are released as a function of species traits. This model estimated total releases into the wild in a year, but does not consider environmental or spatial issues, which should affect probability of survival
72	Gertzen & Leung 2011	Crustacean (Spiny water flea, <i>Bythotrephes</i>)	1	Canada	Yes (stochasticity modeled demographic and environmental)	Demographic model incorporating Allee effects, stochasticity, and mesocosm experiments. Looking at the effect of initial population size (propagule pressure) versus establishment success.	N/Y/Y/N/N	$X_{i+1} = f_x(X_i, r(E_i), \beta, \sigma_e, \sigma_d)$ $Z = P_i^c = pr(X_i > threshold)$	Demographic model incorporating Allee effects, stochasticity, and mesocosm experiments. Looking at the effect of initial population size (propagule pressure) versus establishment success

73	Gevrey et al. 2006	Insects	844	Global	No	Self-organising map used to link species assemblages globally to identify potential group membership	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	Looked at pest assemblages/co-occurrence. Could consider it a special case of E referring to biotic predictors (other pests)
74	Gillham et al. 2004	Plants (black henbane, <i>Hyoscyamus niger</i> ; hoary cress, <i>Cardaria draba</i> ; leafy spurge, <i>Euphorbia esula</i> ; perennial pepperweed, <i>Lepidium latifolium</i> ; spotted knapweed, <i>Centaurea maculosa</i>)	5	USA (Wyoming)	Yes (independent data)	Occurrence data, degree of disturbance, WISP model (automated extension within the GIS program)	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
75	Gladon et al. 2008	Plant (green alga, <i>Caulerpa taxifolia</i>)	1	USA (Florida)	Yes (*2)	Species distribution model calibrated with occurrence data, including biotic factors (native species presence, human population density)	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i, f(V))$	
76	Goodwin & Piggot 2009	Bacteria	1	USA (Florida)	Yes (comparison of model performance of three model variants)	Spatiotemporal models of the risks of citrus canker transmission	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	Not sure where the spatiotemporal aspect is - seemed to be based solely on host/acraege
77	Goslee et al. 2003	Plant (Russian knapweed, <i>Acroptilon repens</i>)	1	USA (Colorado)	Yes (model predictions compared with abundance maps)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
78	Goslee et al. 2006	Plants	1	USA	No	Demographic model applied to cells in landscape	N/N/Y/N/N	$Z = X_{i,j+1} = f_x(X_{i,j}, r(S), \kappa, \sigma)$	Not enough info provided to be sure (IBM approach). Also looked at error rates. Also looked at error rates.
79	Gravuer et al. 2008	Plants	70	New Zealand	Yes (variation explained)	Boosted-tree methods on human, biogeographic, species trait predictors on introduction, naturalization and spread	Y/Y/N/Y/N	$N = f(S, f(V), E)$ $Z = N * pr(\lambda_{c2} S, E, T, f(V)) * \Delta Q(S, E, T, V)$	Why is the spread rate only V?
80	Gray 2010	Insect	1	USA	Yes (probability models, standard errors)	Phenology model on probability of introduction and establishment	Y/Y/N/N/N	$N = O(X_i) * f_p(\bar{E})$ $Z = P_{i,j}^A = 1 - \prod_{i=1}^T (1 - pr(\lambda_i))^{y_{i,j}}$	O-predature phenology dev, fg
81	Gross 2001	Mammals (mountain goats, <i>Oreamnos americanus</i>)	1 species / 1 pathway (impact)	USA		Spatially explicit, individual-based model that forecasts population dynamics of a native and an exotic species	N/N/Y/Y/Y	$N_{i,j} = \sum_{j=1}^J f_{3D}(X_{i,j}, f_{2D}(\bar{E}, D_{i,j}))$ $X_{i,j+1} = f_x(X_{i,j}, r(E_{i,j}), \kappa(E_{i,j}), \sigma_e, \sigma_d)$ $E_{i,j+1} = f_E(X_{i,j}, E_{i,j})$ $Z_T = \sum_{i=1}^I \sum_{j=1}^J f_i(X_{i,j}, E_{i,j})$	Descriptions verbal, so am not sure. E is mainly biotic - includes competition from species of interest Big Horn Sheep and disease, as well as abiotic environmental factors. Impact is also effect on Big Horn Sheep (i.e., a function of biotic environment wrt invasive mountain goats). Dispersal causes immigration (emigration implicit in fx in our mapping)
82	Haight et al. 2011	Plant pathogenic fungus (oak wilt)	1	USA (Minnesota)	Yes (sensitivity analysis)	Spread model that combines historic data on short- and long-distance dispersal with costs of removing infected trees to simulate future development and costs of the disease	N/N/N/Y/Y	$Z_T = Y^{-A} = \sum_{i=1}^T \left(\sum_{j=1}^{i-1} (W_{i,j+1} * Y_j^B) \right)$ $Y_j^B = \sum_{i=1}^{j-1} (M_{i,j}^A + f_j(X_{i,j+1}, E_{i,j}, \alpha' M_{i,j}^A))$	W/2.12 is the probability of establishment of new pockets at time t2 (taken here as a constant rate). Damage is a function of pocket age (t3), we use X as area infested within call. E is the environment (Oaks), and MC is the management (cutting down trees). Estimates expected damages in absence of other preventative measures. Also calculated post-establishment
83	Hallstan et al. 2010	Mollusc (zebra mussel, <i>Dreissena polymorpha</i>)	1	Sweden	Yes (percentage correct prection and AUC using cross-validations)	Hierarchical partitioning and stepwise selection of water chemistry variables in a multiple logistic model using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
84	Hamilton et al. 2005	Plants	+150	Eastern Australia	Yes (validation data sets and error rates)	Species trait model that compares life-history correlates of invasion success between regional and continental spatial scales	N/N/Y/N/N	$Z = \bar{X} = f_x(S)$	Predicting average abundance
85	Hanson et al. 2008	Crustacean (crab)	1	USA	No	Verbal analysis of hidrographic and water quality variables as predictors of crab occurrence and larval development	N/Y/N/N/N	$P_{i,j}^B = 1 - \prod_{i=1}^T (1 - pr(\lambda_B E_{i,j}, S))$ $Z = P^{C,i,T} = P^{B,i,T} * pr(\lambda_c E_i, S)$	Note that prB did not include any propagule pressure component, but was included to indicate that timing issues were considered. Species traits here was determined for a specific species, using experimental studies rather than statistically across many species (as a discriminator)
86	Harris & Barker 2007	Insects (ants)	12	New Zealand	No	GIS-based modelling tool for management of biosecurity risks to New Zealand's indigenous ecosystems based on occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
87	Hartley et al. 2006	Insect (Argentine ant, <i>Linepithema humile</i>)	1	Global	Yes (concordance probability and misclassification errors)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
88	Henderson et al. 2011	Vertebrates	137	Australia	Yes (generalised linear models between the risk of establishment and vertebrate classes or detection categories)	Four-factor risk assessment based on the outcomes of historical introductions of exotic species and climate matches	N/Y/N/N/N	$Z_{i,v} = f_{i,v}(O_i * f_{g,v}(.)) * pr(\lambda_{c,i} E)$	Separated by vector type (v) and calculated for a series of individual species explicitly (not generalizing beyond species) (denoted here as fs.v), i.e., calculating risk ranks for the combination of species and vector. Used interception data (fs.v) which therefore we make a function of uptake and survival (but not release)
89	Herbert et al. 2012	Mollusc (Manila clam, <i>Ruditapes philippinarum</i>)	1	United Kingdom	Yes (independent data)	Hydrodynamic models coupled with a water salinity model and an individual behaviour model of Manila clam larvae	N/N/N/Y/N	$N_{i,j} = \sum_{j=1}^J f_D(\bar{E}, D_{i,j}, S, v, \sigma_D)$	v included because explicitly hydrodynamic model, and behavioural movement. E is salinity
90	Herborg et al. 2007	Crustacean (crab)	1	North America	Yes (misclassification rates)	Potential distribution of species by combining the use of habitat suitability models with data on the intensity of shipping activity	Y/Y/N/N/N	$Z = \sum_i f(V_i, v) * pr(\lambda_c E_i)$	f(Vi,v) here is the vector density map at each given location i. Multiple vectors examined, and a combined score generated for each location. This was multiplied by niche model output
91	Herborg et al. 2009	Tunicate	1	North America	Yes (AUC)	Establishment probability of invasive species by combining vector and environmental niche models	Y/Y/N/N/N	$Z = \sum_i \sum_v f(V_{i,v}, v) * pr(\lambda_c E_i)$	Developed a risk map, cumulating over pathways (v), and the number of vectors in that pathway Vv, weighted by risk of transport by that vector. And then standard SDM. Separated for each location i

92	Herrera et al. 2012	Plants	1	Venezuela	Yes (sensitivity analysis)	Demographic model of weeds integrated with simulations of management	N/Y/Y/N/N	$Z = P_i^A = pr(X_i > threshold)$ $X_{i+1} = f_X(X_i, r(S), \kappa; \sigma_e, \sigma_d, M')$	Looked at probability of establishment using PVA. Also considered stage structured population model, not shown in mapping
93	Hoddle 2003	Insect (glassy-winged sharpshooter, <i>Homalodisca coagulata</i>) and bacteria (grape pathogen, <i>Xylella fastidiosa</i>)	2	Global	No	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i)$	
94	Hortal et al. 2010	Plant (sweet pittosporum, <i>Pittosporum undulatum</i>)	1	Azores	Yes (cross validation)	Species distribution model calibrated with occurrence data, human impact considered	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i, f(V))$	
95	Hyvonen et al. 2010	Plants	163	Finland	Yes (explained variance)	GLMM and redundancy analysis on land-use and climate predictors on alien weed richness, occurrence and abundance in crop fields	N/Y/N/N/N	$Z_i = f_{N_i}(E_i)$ $Z_i = f_i(E_i)$ $Z_i = pr(\lambda_{c_i} E_i)$	Ei (land use) here could also be considered management options, but followed authors usage
96	Ibáñez et al. 2009	Plants (<i>Berberis thunbergii</i> , <i>Celastrus orbiculatus</i> , <i>Eucorynus alata</i> , <i>Elaeagnus umbellata</i> and <i>Rosa multiflora</i>)	5	New England, eastern North America	Yes (spatial variability and parameter uncertainty via Bayesian methods; model uncertainty via ensemble models. Validation data sets also used. Also, looked at error rates extrapolated to new areas)	Species distribution abundance model that combine species richness and species per cent ground cover with several environmental predictors (related to climate, local habitat and land cover)	N/Y/Y/N/N	$Z = \sum_{i=1}^L f_X(E_i) pr(\lambda_{c_i} E_i)$	Using same logic and techniques associated with habitat suitability models but applied to predict species abundance
97	Inglis et al. 2006	Molluscs	2	New Zealand	Yes (sensitivity analysis)	Model comparison between simple expert-based habitat suitability models and regression models	N/Y/N/Y/N	$N_{i,j} = f_D(D_{i,j}, \vec{E})$ $Z = \sum_{i=1}^L pr(\lambda_{c_i} E_i, N_{i,j})$	Included SDM and oceanographic dispersal model within regression model to predict establishment. Vector E is the ocean circulation environment, and Dij is distance to nearest source
98	Jacobs & MacIsaac 2009	Plants	1	Canada	Yes (ensemble modelling)	Passive and active dispersal models coupled with an environmental suitability model. Measures of propagule pressure incorporated both human-mediated dispersal and advective flow from invaded to non-invaded systems, while habitat suitability was forecasted by combining native and global data sets and using statistical models	Y/Y/N/Y/N	$V_{i,j,d} = V_{j,d} f_D(\vec{E}, \vec{D}_j, E_i, D_{i,j})$ $N_{i,j} = \alpha \sum_{j=1}^J V_{i,j,d}$ $N'_{i,j} = \sum_j f_D(D_{i,j})$ $Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i) (f_N(N_{i,j}) + f_N(N'_{i,j}))$	2 vectors were considered human and natural (differentiated with a 'superscript'). Propagule pressure estimates converted to probability, and combined as shown below
99	Jakubowski et al. 2010	Plant (reed canarygrass, <i>Phalaris arundinacea</i> L.)	1	USA (southern Wisconsin)	No	Species distribution model with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i)$	
100	Jarosik et al. 2011	Plants (<i>Ageratum houstonianum</i> , <i>Argemone ochroleuca</i> , <i>Chromolaena odorata</i> , <i>Opuntia stricta</i> , <i>Xanthium strumarium</i> and <i>Lantana camara</i>)	6	South Africa (Kruger National park)	Yes (cross validation to estimate relative errors)	Species distribution model with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i, f(V))$	6 species examined. F(V) was roads
101	Jokela & Ricciardi 2008	Mollusc (Eurasian zebra mussel, <i>Dreissena polymorpha</i>)	1	Canada-Quebec	Yes (r ²)	Regression model relating mean fouling intensity to environmental variables	N/N/N/Y/Y	$Z_i = f_i(E_i)$	Empirical study, looking at several sites and environmental correlates of fouling (impact)
102	Jones et al. 2010	Plants (<i>Geranium robertianum</i> , <i>Hedera helix</i> , <i>Ilex aquifolium</i>)	3	USA (Olympic Peninsula)	Yes (AUC and true skill statistic)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i)$	
103	Kadoya & Washitani 2010	Insect (bumblebee)	1	Japan	Yes (Bayesian posterior distributions)	Stochastic spatio-temporal model that incorporates immigration and establishment processes to assess spatial dynamics of invasion	Y/Y/N/Y/N	$Q_t = \sum_{i=1}^L \sum_{j=1}^J (f_D(D_{i,j}) + f(O)) * pr(\lambda_{c_i} E_i)$ $Z_t = Q_t (M')$	f(O) amount of tomatoes (source area). 2 paths considered, which in our mapping we sum. Management also modeled as eradication of N number of cells, in different configurations, and the effect of this on spread
104	Kaiser & Burnett 2010	Reptile (brown treesnake, <i>Boiga irregularis</i>)			They do not formally model uncertainty beyond that regarding the point of entry, using expected values for marginal damages, marginal costs, and an area's invasion population at a given time. The model is parameterized using data from Hawaii and Guam, and investigate across 30 years of potential snake presence on the island of Oahu to identify the net benefits of EDRR. The model assumes that dispersion is through a diffusion model (it is said that this is the best way to model it based on the invasion of Guam, but it is not validated with data from Oahu), taking into account accessibility (based on distance to closest road) but not other environmental factors (e.g. habitat suitability).	Spatial-dynamic model for optimal early detection and rapid response that consists of search activities beyond the ports of entry, where search (and potentially removal) efforts are targeted toward areas where credible evidence suggests the presence of an invader. Costs are a spatially dependent variable related to the ease or difficulty of searching an area, while damages are assumed to be a population-dependent variable. The spread and population model of the invader considers the accessibility of the terrain (distance to roads)	Y/N/Y/Y/Y	$N_{i,j} = \sum_{j=1}^J f_{3D}(X_{i,j}) f_D(D_{i,j}, E_i)$ $X_{i,j+1} = f_X(X_{i,j}, r, \kappa) + N_{i,j}$ $Z_t = \sum_{i=1}^I \sum_{j=1}^J M^i + M^j + f_1(X_{i,j}, E_{i,j}, M', M')$	E in ID is distance to road. E on II is other species. Ms is surveillance effort, which also affects II since can only apply strategies once detection occurs
105	Keller et al. 2007	Molluscs	18	Great Lakes basin and the 48 contiguous states of USA	Yes (validation data sets and error rates)	Species trait model that differentiate between characteristics of benign and nuisance species. Time since establishment used to explain negative impacts of molluscs	N/N/N/Y/Y	$Z = f_i(S)$	Species trait models have been applied to a number of the invasion stages, but never integrated. Metrics of impact questionable. Not a quantification of severity of damage, but a probability of any damage occurring.
106	Keller et al. 2008	Crustacean (rusty crayfish)	1	USA (Wisconsin)	Yes (AIC and AUC)	Predictive occurrence model applied with different management options to calculate cost benefit scenarios	N/Y/N/N/Y	$Z = \sum_{i=1}^L pr(\lambda_{c_i} E_{i,2}, f(V), \alpha^{t+2} M_{i,2}^{t+2}) * f_i$	propagule pressure. Assumed 100% effectiveness in management
107	Kelly et al. 2007	Fungi (<i>Phytophthora ramorum</i>)	1	USA	Yes (cross-validation)	Species distribution models calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i)$	
108	Kikillus et al. 2010	Reptile (red-eared slider turtle, <i>Trachemys scripta elegans</i>)	1	New Zealand	Yes (partial AUC using cross-validation)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i)$	
109	Kilroy et al. 2008	Diatom (<i>Didymosphenia geminata</i>)	1	New Zealand	No	Multivariate environmental distance using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i)$	
110	Kimberling 2004	Insects	83	USA	No	Traits-based prediction of success/failure and non-target impacts of biocontrol agents	N/N/N/Y/Y	$Z = f_i(S)$	Only considered analysis with respect to non-target effects (i.e., the risk of this introduction), rather than the management effectiveness
111	Kleinbauer et al. 2010	Plant (black locust, <i>Robinia pseudacacia</i> L.)	1	Austria	Yes (AUC and kappa using cross-validation)	Species distribution models calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_{c_i} E_i)$	

112	Knowler & Barbier 2005	Plants	1	USA	No	Cost benefits analysis of introductions into nurseries	N/Y/N/Y/Y	$N_{it} = \sum_{j=1}^J O_{ij}$ $Y^A = \sum_{t=1}^{T-d1} c Q_t$ $Z_T = \sum_{t=1}^T (f_i(N_{it}) * Y^A)$	O was # per nursery, J is # nurseries. F(N) is a hazard function, and here is the probability of invasion during time interval t1. In paper, these are modeled as continuous functions, but discretized for consistency with full model. d1=1 and is the delay to invasion, and c is the average impact. Article also looked at discounting, which we don't do here, and don't consider the cost-benefit (risk management) side
113	Koch et al. 2011	Forest insects from border interception	Not specified	USA	no	Prediction of species establishment from commodity import data	N/Y/N/N/N	$Z = f_i(V^{prox})$	Examined at 2 levels, initial introduction (Vproxy is amount of trade), and into urban areas (Vproxy is also goods into urban areas). Not combined. Looked at # species introduced
114	Koch et al. 2012	Forest insects	0 (they model the pathway)	USA	Yes (use AIC for model selection, likelihood ratio tests and standard errors)	An analysis of the dispersal potential of pest species via transport of firewood to campsite (consider travel distances, but do not include any assumptions on number of insects carried)	N/N/N/Y/N	$Z_{i,j} = N_{i,j} = \sum_j V_{i,j} f_D(D_{i,j})$	Where V is the number of campers. Then used an integrodifference model, with multiple dispersal kernels. Determine transport of firewood to location i
115	Kolar & Lodge 2002	Fishes	45+66+14	USA	Yes (jackknife validations of discriminant functions, and cross-validation of CART trees)	Statistical model examining the relationship between a series of predictors and establishment risk, followed by an additional analysis	N/Y/N/Y/Y	$Z = P^* = f_{P^*}(S)$ $Z = \Delta Q = f_Q(S)$ $Z = f_i(S)$	Also included invasion history
116	Konishi et al. 2009	Fish	1	Japan	Yes (statistical parameter distribution; SE etc)	Occurrence pattern of a species in relation to 16 environmental habitat variables	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	includes both abiotic and biotic factors
117	Kraus et al. 2012	Reptile (lizard)	1	USA	No?	Climate matching model on predicted distribution in Hawaii and Global. Impact not integrated in the model	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
118	Kriticos et al. 2003	Plant (Prickly acacia, <i>Acacia nilotica</i>)	1	Australia	Yes (fitted parameters considering the species known distribution)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
119	Kueffer et al. 2010	Plants	383	Many	Yes (standard errors/r^2)	Global analysis of alien species richness across islands using island geographic and economic predictors. Explores species traits but does not include them in the model	N/Y/N/N/N	$Z_i = f_i(E_i, f(V_i))$	i is each separate island
120	Kulhanek et al. 2011a	Fish (common carp, <i>Cyprinus carpio</i>)	1	USA (Minnesota)	Yes (spatial variability and parameter uncertainty via Bayesian methods; model uncertainty via ensemble models. Validation data sets also used. Also, looked at error rates extrapolated to new areas)	Species distribution abundance model	N/Y/Y/N/N	$Z = \sum_{i=1}^I f_x(E_i) pr(\lambda_c E_i)$	Using same logic and techniques associated with habitat suitability models but applied to predict species abundance
121	Kulhanek et al. 2011b	Aquatic invaders (especially common carp, <i>Cyprinus carpio</i>)	19	Global	Yes (explained variance)	Model used invasion history data to predict severity of impacts (biotic and abiotic). Meta-analysis of common carp impacts based biomass and time since introduction as predictive variables.	N/N/N/N/Y	$Z_i = f_i(X_i)$	Has primarily used previous observed relation between abundance and metrics of impact. Note that invasion history has been used in species trait models, as a dichotomous variable (has species caused impact anywhere)
122	Küster et al. 2008	Plants (neophytes)	388	Germany	Yes (explained variance)	Statistical models examining the effects of traits (related to morpho-physio-phenological traits, ecological performances and variables describing introduction history) on species frequency	N/Y/N/N/N	$Z = P^* = f_{P^*}(S, t)$	t is the time of first occurrence
123	Lambdon & Hulme 2006	Plants	862	Mediterranean	Yes (sensitivity analysis)	Introduction traits used to predict invasion risk categories using canonical discriminant analysis	N/Y/N/N/N	$Z = P^* = f_{P^*}(S, v, t, N)$	v is the mode of introduction, and t is the first time of first introduction
124	Lambdon et al. 2008	Plants	382	Mediterranean	Yes (elasticity analysis)	Introduction traits used to predict naturalisation using GLM	N/Y/N/N/N	$Z = P^* = f_{P^*}(S, v, t, N)$	v is the mode of introduction, and t is the first time of first introduction
125	Larson et al. 2001	Plants	38	USA	Yes (standard errors/r^2)	Statistical model assessing the effect of vegetation, anthropic and park unit as predictors for alien species distribution (occurrence) and frequency	N/Y/N/N/N	$Z_i = f_i(E_i)$ $Z_i = pr(\lambda_c E_i)$	Seems to look at total # plants (richness?)
126	Leung et al. 2006	Molluscs (zebra mussel)	1	USA (Michigan)	Yes (least square analysis)	Gravity model used to model boat traffic between lakes as variables for the spread of zebra mussel (i.e. surrogate for propagule pressure). Potential establishment is not considered.	N/N/N/Y/N	$Z_{i,j} = \sum_j V_{i,j} f_D(\bar{E}, \bar{D}_j, E_i, D_{i,j})$	
127	Leung & Mandrak 2007	Mollusc (zebra mussel, <i>Dreissena polymorpha</i>)	1	USA (Michigan)	Yes (uncertainty examined using hindcasts)	Joint propagule pressure-species distribution model that simultaneously incorporates invasibility and propagule pressure to predict invasion risk	Y/Y/N/Y/N	$V_{i,j,t} = V_{i,j,t} f_D(\bar{E}, \bar{D}_j, E_i, D_{i,j})$ $N_{i,j} = \alpha \sum_{j=1}^J V_{i,j,t}$ $P_{i,t}^A = 1 - \prod_{j=1}^J (1 - pr(\lambda_A))^{N_{i,j}^A}$ $Z = Q_t = \sum_{i=1}^I P_{i,t}^A * pr(\lambda_c E_i)$	Extend of propagule pressure-establishment model (e.g. Leung et al 2004) by also considering environmental differences between locations. pr(C E) is estimated using SDM. As opposed to SDM this does not suffer from requiring equilibrium. Note that species distribution model was termed "site invasibility" in Leung & Mandrak 2007, which we change here to be consistent with the majority of the literature
128	Leung et al. 2004	Mollusc (zebra mussel, <i>Dreissena polymorpha</i>)	1	USA (Michigan)	Yes (uncertainty examined using hindcasts)	Propagule pressure-establishment model that estimates the probability of population establishment taking into account propagule pressure and allele effects using survival analysis and maximum likelihood techniques	Y/Y/N/Y/N	$V_{i,j,t} = V_{i,j,t} f_D(\bar{E}, \bar{D}_j, E_i, D_{i,j})$ $N_{i,j} = \alpha \sum_{j=1}^J V_{i,j,t}$ $P_{i,t}^A = 1 - \prod_{j=1}^J (1 - pr(\lambda_A))^{N_{i,j}^A}$ $Z = Q_t = \sum_{i=1}^I P_{i,t}^A * pr(\lambda_c E_i)$	Uses vector traffic as a surrogate of propagule pressure, and calculates establishment as a function of propagule pressure. Implicitly assumes all propagules have equal probability and that no interaction occurs with environment (compare with full model, "establishment"). Note. J refers to all invaded sources, rather than all sources
129	Leung et al. 2002	Mollusc (Zebra mussel)	1	North America	Yes (stochasticity included in dynamic model, sensitivity analysis)	Bioeconomic model taking into account population dynamics of zebra mussel, and economic losses experienced by a power plant. Adaptation occurred via different investment into labor and capital	N/Y/Y/N/Y	$X_{i,t+1} = f_x(X_{i,t}, r, \kappa, \sigma)$ $Y^B = \sum_{j=1}^{J_{in}} (M_{ij}^B + f_j(r3, X_{i,t}, \alpha' M_{ij}^B))$ $Z_t = \sum_{i=1}^I ((1 - p_{it}^C(\alpha' M^B)) * M^B + U_{it}(\alpha' M^B))$	Bioeconomic approach. Not spatially explicit. Took into account population dynamics, and losses experienced by a power plant. Control was also in a power plant, and reduced the Zebra Mussel population size in the power plant only. Adaptation occurred via different investment into labor and capital. Was not spatially explicit

130	Liddle et al. 2006	Vertebrates	3	Australia	Yes (sensitivity analysis)	Assessment of the level of impact by an invader	N/N/N/N/Y	$\sum_{i=1}^n f_i(i, E, M')$	Looked at impact due to invader in terms of palm trees. Simulated management effects, and also looked at environment over time. Looked at several sites, but analysis not spatially explicit
131	Lindgren et al. 2010	Plants (<i>Tamarix ramosissima</i> Ledeb., <i>T. chinensis</i> Lour. and hybrids)	3	Canada	No	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
132	Lippitt et al. 2008	Insect (gypsy moth, <i>Lymantria dispar</i> L.)	1	Contiguous USA	Yes (AUC)	Species distribution model using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i, V)$	V is human traffic from source locations
133	Liu et al. 2011	Crustacean (red swamp crayfish, <i>Procambarus clarkii</i>)	1	Global	Yes (model predictions compared to null models; AUC)	Species distribution model calibrated with occurrence data, considered human impact	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
134	Liu 2009	Amphibian (frog)	1	China	Yes (test for alternative predictors)	GLMM for numerous predictors related to farming, propagule pressure, geographic and hydrology as predictors for establishment probability	N/Y/N/N/N	$Z = Q_T = \sum_{i=1}^L pr(\lambda_c E_i, O_i, D_{ij}, I_j, \alpha^p M^p)$	O frog density at source, Dij because distance from a source. Time since source pop existed also considered (ti) and enclosure type was mapped as prevention effort
135	Logan et al. 2007	Insect (gypsy moth, <i>Lymantria dispar</i> L.)	1	USA (Utah)	Yes (cross-validation)	Species distribution model with occurrence data	N/Y/Y/N/N	$X_{i,j+1} = f_{X_i}(X_{i,j}, r(E_{i,j}), \sigma)$ $Z = P_i^A = pr(X_i > threshold)$	Based on verbal description of model. Pop growth parameters function of environment, not clear from description whether K present, or relevant. Assume that sigma included, given that simulations stochastic
136	Mandle et al. 2010	Plant (<i>Impatiens walleriana</i>)	1	Global	Yes (r^2)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
137	Marchetti et al. 2004	Fishes	49 / 22 / 38	USA (California)	No	Three stage-models examining the relationships between species establishment, spread and integration and 10 life history predictor variables	N/Y/Y/Y/N	$Z = P^* = f_p(S, D, N)$ $Z = Q = f_Q(S, D, N)$ $Z = \bar{X} = f_X(S, D, N)$	D here is the distance from the nearest source
138	Maret et al. 2006	Vertebrates	introduced fish and bullfrog	North America	No	Metapopulation model to predict changes in the proportion of ponds occupied by different species	N/N/N/Y/Y	$Q_i = f(Q_{i-1}, E_{i-1})$ $\sum_{i=1}^L Q_i f_i(E_i)$	Modified to just look at major factors considered, and to be centered on the invasive species dynamics. Their functional form is just Levin's meta-pop model with parameterizations. E is fish, drying. Impact is on salamander. Because Levin's model is spatially implicit, do not consider patches i, as with other approaches
139	Marini et al. 2011	Insects (beetles)	Many	USA and 20 European countries	Yes (r^2)	Multi-model inference and hierarchical partitioning of human activity (trading) and environmental variables on species richness and establishment	N/Y/N/N/N	$Z_i = f_i(E_i, V_{primary})$	Not sure what they mean by multimodal inference - seems like a regular multiple regression
140	Mason et al. 2011	Insect (Leek moth, <i>Acrolepiopsis assectella</i>)	1	Canada and eastern USA	Yes (compared with species distribution)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
141	Mau-Crimmins et al. 2006	Plant (Lehmann lovegrass, <i>Eragrostis lehmanniana</i>)	1	Southeastern USA	Yes (misspecification error rates and r^2)	Comparison between invaded-range versus native-range dataset using species distribution models calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
142	Medley 2010	Insect (Asian tiger mosquito, <i>Aedes albopictus</i>)	1	North and South America, Europe	Yes (AUC)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
143	Mika & Newman 2010	Insect (pea leafminer, <i>Liriomyza huidobrensis</i>)	1	North America	Yes (misspecification error rates and r^2)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
144	Mika et al. 2008	Insect (swede midge, <i>Contarinia nasturtii</i>)	1	North America	Yes (misspecification error rates)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
145	Miller et al. 2007	Molluscs (bivalve and gastropod molluscs associated with oyster)	93	USA (San Francisco Bay, California)	Yes (model sensitivity and specificity, but also percent correct classification when testing the model applied to 3 different regions using a bootstrap analysis)	Statistical model employing biological attributes to determine establishment success	N/Y/N/N/N	$Z = P^* = f_p(S)$	Not sure what to do about historical abundance. Treating habitat characteristics a factor relating to species (i.e. S rather than E)
146	Miller et al. 2010	Plants	8	USA	Yes (Monte Carlo simulations)	So called Relative risk model on occurrence data for invasive and rare plants	Y/Y/N/N/Y	$f_i(S, E_i) = f_i(S) pr(\lambda_c^{Native} E_i)$ $Z = \sum_{i=1}^L f(O, d_{ij}) pr(\lambda_c E_i) * f_i(S, E_i)$	f(O) is the sources, which I didn't know how to place in notation - they have uninvasion, adjacent, to invasion, uninvasion
147	Mondor et al. 2007	Insects (aphids)	174 (96 introduced species - 78 native congeners)	USA-Hawaii	No	Morphological and ecological variables used to predict colonization success	N/Y/N/N/N	$Z = P^* = f_p(f(O), S)$	f(O) is presence in continental USA
148	Moore et al. 2010	Vertebrates	1	Australia	No	Stochastic dynamic programming used to assess the optimum allocation of resources to quarantine and surveillance	N/N/N/N/Y	$Z_T = M^P + M^S + U(\alpha^P M^P) * Y^A$ $Q = f(M^S)$ $Y^A = f(Q, M^C)$	SDP model used, and could not entirely map matrix into a small set of equations. Here MP is quarantine and Me is surveillance (which slows the spread). YA is the cost of eradication (Mc), depending on probability of extensive spread Q (a function of Me). Eradication is assumed to be successful. Space is implicit, so do not use notation for patches
149	Muirhead & MacIsaac 2005	Insect (bumblebee)	1	Canada	Yes (Monte Carlo analysis)	Vector traffic model from invaded to non-invaded lakes that assess the patterns of spread of alien species by risky activities	N/N/N/Y/N	$Z_j = \sum_i V_{ij} f_D(\bar{E}, \bar{D}_j, E_i, D_{ij})$	Double constrained gravity model, looking at the risk associated with hub lake j. Discussed management, but did not model it
150	Muirhead & MacIsaac 2011	Crustacean (spiny waterfleas, <i>Bythotrephes longimanus</i>)	1	Canada (Ontario)		Establishment probability and dispersal models	Y/N/N/Y/N	$V_{i,j,t} = V_{i,j} f_D(\bar{E}, \bar{D}_j, E_i, D_{ij}, \sigma_D)$ $N_{i,j} = \alpha \sum_{j=1}^L V_{i,j,t}$ $Z_i = Q_i = \sum_{j=1}^L f(N_{i,j})$	f(N) is a generic functional form using boosted regression trees
151	Muirhead et al. 2006	Insect (Emeral ash borer)	1	Canada	Yes (Stochasticity in establishment probabilistic; cross validation examined)	Dispersal model that considers both natural and human-assisted dispersal	N/Y/N/Y/N	$Z_i = Q_i = \sum_{j=1}^L pr(\lambda_{i,j}) f_D(\bar{E}, \bar{D}_j, E_i, D_{ij})$	Containment examined, but not explicitly modeled, and therefore not included in the mapping. Two dispersal modes considered (long-distance, human mediated) and natural dispersal. ID converted to a probability of reaching I from j, and pr(LA) is a fixed fitted probability
152	Mwebaze et al. 2010	Plant insect pests	2253 interceptions at border inspections	United Kingdom	Yes (sensitivity analysis)	Biocconomic model using the volume of imports and interception data of insect pests at border controls to estimate the number of pests coming in per volume of trade and country. This is used to model the optimal inspection effort allocation to different countries of import origins	N/Y/N/N/N	$Z_i = f_i(M^P, \sum_j f(V_{i,j}))$	MP is inspection, V is the imports from country j to country i, and fs is the number of species introduced. Also did other trade-off analyses, which would be part of risk management

153	Naddafi et al. 2011	Molluscs (zebra mussel)	1	USA, Europe	Yes (model comparisons)	GLM-GAM models on physicochemical variables determining zebra mussel density (abundance)	N/N/Y/N/N	$Z_i = f_x(E_i)$	
154	Neteler et al. 2011	Insect (Asian tiger mosquito, <i>Aedes albopictus</i>)	1	North-eastern Italy	Yes (misspecification error rates)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
155	Nietschke et al. 2008 Økland et al. 2010	Insect (<i>Scirtothrips dorsalis</i>) Pinewood nematode	1 1	USA Norway	Yes (misspecification error rates) Yes (sensitivity analysis)	Species distribution model with occurrence data Distribution model of a vector species linked to a surveillance/monitoring model to test how contingency plans work	N/Y/N/N/N N/Y/Y/Y/Y	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$ $X_{j+1} = f_x(X_{j,j}, r(E_{j,j}), \kappa(E_{j,j}))$ $N_{i,j} = \sum_{j=1}^J f_{xj}(X_{j,j}) f_D(D_{i,j,j}, \sigma_D)$ $Z = P_{i2,j}^A = pr(X_i > threshold)$ $Z_i = \sum_{j=1}^J (P_{i2,j}^C, M^C, M^S)$	Dispersal random - indicates movement to location i2, at which time establishment occurs. Risk a function of everywhere establishment occurs, given eradication and monitoring efforts. Paper actually about probability of eradication
156									
157	Olden et al. 2006	Fishes (native and nonnative)	90 (28 + 62)	USA (Colorado River Basin)	No	Statistical models assessing the relationship between life between life-history overlap and species distributional changes	N/N/N/Y/N	$Z = \Delta Q = f_Q(S)$	Looked at changes in distribution. Delta Q is the spread rate
158	Olden et al. 2011	Crustacean (rusty crayfish, <i>Orconectes rusticus</i>)	1	USA (Wisconsin)	Yes (error rates; takes into account spatial variability)	Joint suitability-vulnerability model that quantifies ecosystem vulnerability to species distribution as a function of exposure risk (i.e., likelihood of introduction and establishment based on a species distribution model) and the sensitivity of the recipient community (i.e., likelihood of impacts on native species)	N/Y/N/Y/N	$Z = \sum_{i=1}^I pr(\lambda_{c2} E_i) pr(\lambda_{c2}^{native} E_i)$	Linked habitat suitability models for invader and native species of interest. The probability of both occurring simultaneously over all sites determines the risk. Like other habitat suitability models, does not consider temporal dynamics of invasion process
159	Offert et al. 2006	Insect (swede midge, <i>Contarinia nasturtii</i>)	1	Canada	Yes (misspecification error rates)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
160	Offert et al. 2004	Insect (<i>Oulema melanopus</i>)	1	Canada	Yes (misspecification error rates)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
161	Olson et al. 2012	Plant (<i>Myriophyllum spicatum</i> L.)	1	USA (Wisconsin)	Yes (AUC and explained variance, error type I and II)	Multivariate logistic and multiple linear regressions using both occurrence and abundance data. Human impact and other biotic factors included	N/Y/N/Y/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i, f(V)) f_x(E_i)$	
162	Ofinowski et al. 2007	Plants	251	Canada	Yes (spatial variation)	Climate, habitat and species traits explaining establishment and proliferation (abundance?). Not very clear, it does not integrate climate match with native range with species traits, close to a qualitative analysis	N/Y/N/N/Y	$Z = pr(\lambda_c E) * pr(\lambda_c S) * f_i(S)$	Looked at probability of existing in region at all, and probability that traits would occur in park, in susceptible areas. Used a decision tree (convert to 0/1 instead of continuous). Each component fit independently to same data
163	Paini et al. 2011	Fungus	486	Global	Yes (validation with a virtual world, ensemble model)	Self organizing map by artificial neural network on establishment likelihood	N/Y/N/Y/N	$Z = Q = \sum_{i=1}^I pr(\lambda_{c2} E_i)$	Looked at pest assemblages/co-occurrence. Could consider it a special case of E referring to biotic predictors (other pests). Validation of theory, not consideration of real world uncertainty
164	Parker et al. 1999	All invaders	Not specified	Global	No	Conceptual model for understanding the impacts of an invader including three fundamental dimensions: range, abundance and its per-capita or per-biomass effect. No models were used	N/N/N/Y/Y	$Z_T = Q_T * \bar{X} * c$	Implicitly assumes that locations are all equal, abundances are all equal, and that consequences of abundance are linear, scaled by a per unit impact constant @
165	Peacock& Womer 2006 Perry & Vice 2008	Insect pests Reptile (brown tree snake, <i>Boiga irregularis</i>)	1 1	New Zealand Global	No No	Species distribution model calibrated with occurrence data Probability of establishment. Included propagule pressure in the model, also incorporated the likelihood of an organism entering the transportation system, avoiding detection, surviving to arrive at another location, and establishing at the receiving end	N/Y/N/N/N Y/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$ $Z_{i,j} = N_{i,j} = pr(\lambda_c) \sum_{j=1}^J \sum_{k=1}^{V_{i,j,k}} O(v_{i,j,k,j}) * f_j$	Multiplies probability of a given individual establishing, times the frequency of vectors (V) times the probability of survival transport (fg), times the probability of being taken up and remaining undetected. These are functions of the vector (transport vessel size)
166									
167	Peterson et al. 2003	Plants (<i>Hydrilla</i> , <i>Hydrilla verticillata</i> ; Russian olive, <i>Elaeagnus angustifolia</i> ; sericea lespedeza, <i>Lespedeza cuneata</i> ; garlic mustard, <i>Alliaria petiolata</i>)	4	North America	Yes (misspecification error rates)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
168	Pitt et al. 2009	Insect (Argentine ant, <i>Linepithema humile</i>)	1	New Zealand	Yes (misspecification error rates)	Occurrence and Spread Data. Spatially explicit stochastic model and uniform radial spread model	Y/Y/N/Y/N	$N_{i,j} = \sum_j f_D(D_{i,j}, v, \sigma_D)$ $Z = Q = \sum_{i=1}^I pr(\lambda_c E_i) pr(N_{i,j} > 0)$	Based on verbal description of model. Both local and long distance dispersal considered. ID seemed to yield a probability of that dispersal would reach i, so we treated this as at least one dispersal event would reach i from all infested locations
169	Potapov et al. 2011	Zooplankton	1	Canada	Yes (variation explained)	Stochastic gravity model for propagule pressure and logistic regression model on water physicochemical conditions for establishment	Y/Y/N/Y/N	$V_{i,j} = V_{i,j} f_D(\vec{E}, \vec{D}, E_i, D_{i,j}, \sigma_D)$ $N_{i,j} = \alpha \sum_{j=1}^J V_{i,j}$ $Z = Q_i = \sum_{j=1}^J (1 - pr(\lambda_i))^{v_i^j} * pr(\lambda_c E_i)$	
170	Potter et al. 2009 Pysek et al. 2009	Plant (<i>Cytisus scoparius</i>) Plants	1 Ambiguous; it includes transport, introduction, establishment and spread but the boundaries are poorly defined	Global Global	Yes (misspecification error rates) No	Species distribution model calibrated with occurrence data Effects of species' biological traits and their distributional characteristics on invasion success	N/Y/N/N/N N/N/N/Y/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$ $Z = Q = f(S, f(O), v)$	Response variable number of locations found (analogous to spread at a global level), which is a function of species traits and vector types (v). Oproxy is how widespread it is in native range, as a proxy for uptake
171	Pysek et al. 2009	Plants (neophytes)	170 (109+44+17)	Czech Republic	No	Examining the probability of introduction, naturalization and invasions by mean of statistical models using introduction data and species life-traits	Y/Y/N/Y/N	$Z = N = f_x(f(O), v, S)$ $Z = P^* = f_x(f(O), v, S)$ $Z = \Delta Q = f_O(f(O), v, S)$	N equates propagule pressure with the number of escapes, f(O) is the number of stores selling a species, v is the pathway (here the usage of the species)
172									
173	Raghu et al. 2007	Insect (beetle for biological control)	1	Australia	Yes (sensitivity analysis)	Life-cycle model for a biological control insect is combined with a model for a target and non-target plant growth model to assess potential benefits (control of target species) and risks (of non-target native species)	N/N/Y/N/Y	$X_{i+1} = f_x(X_i, r(E_i), \kappa(E_i))$ $Z = f_i(X_i, E)$	E is plants present (one for biocontrol and one a non-target effect). Population dynamics model actually had many stages and more complex than mapped. Was modelled per patch (i.e., spread not considered)
174	Raimundo et al. 2008	Terrestrial vertebrates	13	Spain	Yes (discrimination power of favourability functions; AUC, correct classification rates)	Species presence/absence related to independent variables	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i, f(V))$	

175	Raimundo et al. 2007	Plant (Siamweed, <i>Chromolaena odorata</i>)	1	Neotropics (native range) and global (introduced range) USA	Yes (jackknife evaluation and misspecification error rates)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
176	Reed 2005	Reptiles (boas, phytons, 404177 individuals, 17 genera and 40 species of snakes)	6067 shipments representing		No	Model estimating the likelihood of establishment with legal commercial imports by summing risk associated with commercial trade and risk associated with ecological variables	Y/Y/N/N/N	$N = f(O, S)$ $P^C = f_{pr}(S)$ $Z = N + P^C$	Species traits here are related to their value, and the way they were caught (wild vs raised). O is the number being traded (consider survival and release after that)
177	Reichard & Hamilton 1997	Woody plants	235	North America	Yes (validation data sets and error rates)	Species trait model that determines which traits (several structural, life history, and biogeographical attributes) characterize establishment of invasive woody plant species using discriminant analysis	N/Y/N/N/N	$Z = P^* = f_{pr}(S)$	Species trait models have been applied to a number of the invasion stages, but never integrated. Metrics of impact questionable. Not a quantification of severity of damage, but a probability of any damage occurring. Although applied separately, these approaches could be integrated for a fuller analysis of risk
178	Reino et al. 2009	Bird	1	Portugal	Yes (statistical parameter distribution; SE etc)	Survival analysis based on information available up to the first invasion peak used to predict pattern of invasion in the second peak, as well as future invasion hazards using climate change scenarios	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	Used survival analysis, but all aspects were looking at colonization using climate as a predictor
179	Reshetnikov & Ficetola 2011	Fish (<i>Perccottus glenii</i>)	1	Holarctic	Yes (AUC and cross-validation)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
180	Ricciardi 2003	Mollusc (zebra mussel, <i>Dreissena polymorpha</i>)	1	Europe and North America	Yes (explained variance)	Regression analysis of data (invasion history) from multiple invaded sites to generate empirical models of impact	N/N/N/N/Y	$Z_i = f_i(X_i)$	Has primarily used previous observed relations between abundance and metrics of impact. Note that invasion history has been used in species trait models, as a dichotomous variable (has species caused impact anywhere)
181	Richards et al. 2008	Insect (whitefly)	4	USA	Yes (use of three different models and each is made to vary according to a geometric Brownian motion, to simulate random variation in the deterministic model. The models are fitted to empirical data)	Econometric model that takes into account the spatio-temporal process underlying a particular population and apply it to the economic costs of the species damage	Y/N/Y/Y/N	$Z_{i,j} = N_{i,j} = \sum_{j=1}^J f_{XB}(X_{i,j}) f_D(D_{i,j})$ $X_{i,j+1} = f_X(X_{i,j}, r(E_{i,j}), \kappa(E_{i,j}), \sigma) + N_i$	Pricing model. Not sure what risk is. E incorporates space and time differences (could also be expressed as an exogenous matrix). Sigma causes variance, is also autoregressive
182	Richardson & Thuiller 2007	Plants	1	Global (regions similar to South Africa)	Yes (AUC)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
183	Robertson et al. 2004	Plants (<i>Lantana camara</i> ; <i>Ricinus communis</i> ; <i>Solanum mauritanium</i>) and native cicada species (<i>Capicada decora</i> ; <i>Platypleura deusta</i> ; <i>P. capensis</i>)	6	South Africa, Lesotho, Swaziland	Yes (kappa)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
184	Robinet et al. 2011	Pine wood nematode (<i>Bursaphelenchus xylophilus</i>)	1	Europe	Yes (sensitivity analysis)	Occurrence and spread data. Includes human populations	Y/Y/N/Y/N	$N_{i,j} = \sum_{j=1}^J f_{XB}(X_{i,j}) f_D(E_i, D_{i,j}, V)$ $Z = Q = \sum_{i=1}^I pr(\lambda_c E_i) pr(N_{i,j} > 0)$	Has both local and long distance. Human population size in dispersal kernel is mapped as Ei here because it is a metric of attractiveness, rather than a vector. Verbally discussed probability of establishment based on host and temperature, but not clear what relations were
185	Robinson et al. 2010	Plants	1	Australia	Yes (AUC)	Multicriteria evaluation tool coupled with a risk-adjusting technique to develop a series of alternative decision strategies for identifying the distribution of an invasive species	N/Y/N/N/N	$Z = P^* = f_{pr}(S)$	Not sure, but paper look at decisions (risk preferences, which is part of risk management), and does not affect risk itself.
186	Rodder 2009	Amphibian (<i>Eleutherodactylus johnstonei</i>)	1	Global	Yes (AUC and jackknife analysis)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
187	Rogers et al. 2007	Plant (Upland Cotton, <i>Gossypium hirsutum</i> var. <i>hirsutum</i>)	1	Australia	Yes (misspecification error rates)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
188	Rothlisberger & Lodge 2010	Plants	1	USA	Yes (AUC)	Gravity model to predict spread	Y/Y/N/Y/N	$V_{i,j,z} = V_{i,j} f_D(\bar{E}, \bar{D}_j, E_i, D_{i,j}, V)$ $N_{i,j} = \alpha \sum_{j=1}^J (V_{i,j,z})$ $Z = Q_i = \sum_{i=1}^I (1 - pr(\lambda_A))^{N_{i,j}}$	Uses vector traffic as a surrogate of propagule pressure, and calculates establishment as a function of propagule pressure. Implicitly assumes all propagules have equal probability and that no interaction occurs with environment (compare with full model, "establishment"). Note: J refers to all invaded sources, rather than all sources
189	Roura-Pascual et al. 2010	Plants	Many	South Africa	Yes (sensitivity analysis)	Spatially-explicit analysis that evaluates the sensitivity of model-based management prescriptions to changes in the relative importance assigned to different decision criteria	N/N/Y/Y/N	$Z_i = w_1 Q_i + w_2 X_i + w_3 f(t) + w_4 E_i + w_5 M_i$	risk in a given area dependent on spread, abundance, time (age of stand), environment (fire and fire risk), last management, and species type present
190	Rout et al. 2011	Vertebrates	1	Australia	No	Model optimum allocation of resources to quarantine and surveillance	N/N/N/N/Y	$Z_T = M^P + M^* + U(\alpha^* M^P) * Y^A$ $Q = f(M^*)$ $Y^A = f(Q, M^C)$	SDP model used, and could not entirely map matrix into a small set of equations. Here MP is quarantine and Me is surveillance (which slows the spread). YA is the cost of eradication (Mc), depending on probability of extensive spread Q (a function of Me). Eradication is assumed to be successful. Space is implicit, so do not use notation for patches
191	Ruesink & Colado-Vides 2006	Plant (macroalga, <i>Caulerpa taxifolia</i>)	1	Mediterranean sea	Yes (confidence intervals)	Non-spatial, discrete, linear model to describe the increase in area covered by patches of an invader	Y/Y/Y/N/N	$X_{j,t+1} = f_X(X_{j,t}, r, \sigma_x)$ $Q_t = \sum_j X_{j,t}$ $Z = \sum_i pr(\lambda_A) O(Q_i)$	We use X for Area infested, Q for fragments as a function of area and pr(A) for probability of a given individual reattaching
192	Ruesink 2005	Freshwater fishes	1 pathway (establishment) / 200 species	Global	Yes (model evaluation with independent data)	Multiple logistic regression and tree classifications to explore four classes of predictor variables (species traits, environmental traits, match between species and environment, and propagule pressure) susceptible to explain the risk of establishment	N/Y/N/N/N	$Z = P^C = f(E_1 + E_2, \dots + S_1 + S_2, \dots + V_1 + \dots)$	Linear models. All information at the country level. v are reasons for introduction (i.e., pathways such as use in fisheries, etc). Not spatially explicit

193	Sahlin et al. 2011	Plants	113	Europe	Yes (Bayesian posterior distributions)	Using Bayesian analysis to assess the cost-benefit of invasive species screening based on traits	N/Y/N/N/N	$Z = P^* = f_p(S)$	Exclude - main point is base-rate (which we can reference), but mapping will not reflect any of this
194	Sato et al. 2010	Fish (<i>Opsariichthys uncirostris</i>)	1	Japan	Yes (AUC)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
195	Sebert-Cuvillier et al. 2007	Plant (American black cherry, <i>Prunus serotina</i> Ehrh.)	1	Global	Yes (standard error or amount of explained variation applied to the fitted model)	Stage-classified matrix population model describing its population dynamics at the local scale, i.e., within a forest stand, integrating environmental stochasticity	N/N/Y/N/N	$Z = X_{i,j+1} = f_X(X_{i,j}, r, \kappa; \sigma_e)$	Stage based model, only mapping presented. Environment considered in terms of stochasticity
196	Shah et al. 2012	Plants	88	Kashmir Himalaya	Yes (r^2)	Model assessing species invasiveness from native range size	N/N/N/Y/N	$Z = Q = f(S)$	S is how widespread it is in native range
197	Sharma et al. 2009	Fish	1	Canada	Yes (standard errors/r^2)	Classification tree analysis for introduction, stepwise multiple logistic regression models for establishment and native species occurrence in lakes with risk of invasion	Y/Y/N/N/Y	$Z = \sum_{i=1}^L f(V_i) * pr(\lambda_c E_i) * f_i(E_i)$	Vproxy was human population size, and lake area. Impact - biotic environment - occurrence of species of interest. Multiplied 2 processes based on same data, but fitted independently
198	Siesa et al. 2011	Mollusc (crayfish, <i>Procambarus clarkii</i>)	1	Lombardy, northern Italy	Yes (r^2)	Species distribution model using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
199	Skarpaas & Okland 2009	Invertebrate (European spruce bark beetle)	1	Europe	Yes (sensitivity analysis)	Generic model for the first step of the invasion process for trade-imported pests, with an additional development to assess the risk of introductions and alternative management options	N/N/Y/Y/N	$X_{j,i+1} = f_X(X_{j,i}, r, \kappa(V_{j,i}), e(X_{j,i}), I(O^* \text{ * } V_{j,i}))$ $Z = N_{i,j} = f_{XB}(X_{j,i})f_D(D_{i,j})$	Also looked at management effects (but not costs) by modifying parameters (not mapped). Here V (timber imports) is also the habitat. E is the emigration rate, and I is the immigration (propagule pressure here). O is the natural pest density in wood. Special case because vector is also habitat
200	Speek et al. 2011	Plants	111	The Netherlands	Yes (r^2)	Use of plant traits, origin and human uses as predictors of regional and local frequency	N/Y/Y/N/N	$Z = P^* = \sum_{i=1}^L f_p(S, v, t)$ $Z = \bar{X} = \sum_{i=1}^L f_X(S, t)$	Dominance not abundance - not sure how to deal with relative nature. T- period of naturalization time. Human use (yes/no) denoted as a pathway of introduction v
201	Spens et al. 2007	Fish (northern pike, <i>Esox lucius</i>)	1	Northern boreal region of Sweden	Yes (misspecification error rates)	Connectivity model considering dispersal barriers	Y/N/N/N/N	$Z = N = f(E)$	E are connectivity related measures, surrogates of propagule pressure
202	Stanaway et al. 2011	Insect (whitefly)	1	Australia	Yes (Bayesian posterior distributions)	Gravity model to predict spread on plants	Y/Y/N/N/N	$N_{i,j} = \alpha \sum_{j=1}^L O(X_j) * V_{i,j}$ $P_{i,T}^A = 1 - \prod_{i=1}^T (1 - pr(\lambda_A))^{\kappa_i^D}$ $Z_T = Q_T = \sum_{i=1}^L P_{i,T}^A$ $Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	Also looked at detection (observation) error. - look at stochasticity?
203	Stephens et al. 2007	Insect (oriental fruit fly, <i>Bactrocera dorsalis</i>)	1	Global	Yes (misspecification error rates)	Species distribution model calibrated with occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
204	Stohlgren et al. 2010	Plants (<i>Linaria dalmatica</i> , <i>Carduus nutans</i> , <i>Bromus tectorum</i> , <i>Melilotus officinalis</i>)	4	USA (Yellowstone and Grand Teton National Parks, Wyoming; Sequoia and Kings Canyon National Parks, California, and areas of interior Alaska)	Yes (spatial variability and parameter uncertainty via Bayesian methods; model uncertainty via ensemble models. Validation data sets also used)	Species distribution model calibrated with occurrence data and various predictors related to topography, climate and vegetation	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	Species Distribution Models . Many different techniques (e.g. Maxent, GARP, Neural Networks). Implicitly assume equilibrium conditions, but see models in Leung & Mandrak (2007) and Williams et al (2008) for improvements. Treated as probabilities here, as closest analogue in full model, but discrepancies may exist because of biased sampling, use of pseudospecies and base-rates effects (see also Jiménez-Valverde et al. 2011)
205	Strubbe et al. 2010	Aleut bird (ringnecked parakeets <i>Psittacula krameri</i>) and native bird (nuthatches <i>Sitta europaea</i>)	2	Belgium, Brussel	Yes (AUC and cross-validation)	Abundance data using boosted regression trees. Biotic interaction considered in the model	N/N/Y/N/Y	$X_i = f_X(E_i)$ $Z = \sum_{i=1}^L f_i(X_i, E_i)$	
206	Sutherland & Maywald 2005	Insect (Red imported fire ant, <i>Solenopsis invicta</i>)	1	USA, Australia and New Zealand	Yes (misspecification error rates)	Species distribution model calibrated with occurrence data, considered nonclimatic factors	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
207	Sutherland et al. 2007	Insect (ticks: <i>Boophilus microplus</i> , <i>Boophilus decoloratus</i> ; fruit fly: <i>Ceratitis capitata</i> , <i>Bactrocera tryoni</i>)	1	Australia, Africa	Yes (standard deviation)	Species distribution model calibrated with occurrence data, considered species interactions	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
208	Theriault & Herborg 2008	Tunicate (<i>Ciona intestinalis</i>)	1	Canada	Yes (hierarchical partitioning and AUC)	Species distribution model calibrated with occurrence data and two environmental variables (temperature and salinity tolerances)	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
209	Thuiller et al. 2005	Plants (<i>Carpobrotus edulis</i> , <i>Senecio glastifolius</i> , <i>Valerophyton dealbatum</i>)	3	Global	Yes (AUC)	Species distribution model calibrated with occurrence data, considered propagule pressure (trade and tourism as a proxy)	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i, f(V))$	
210	Thuiller et al. 2006	Plants	+500	South Africa	Yes (explained variance)	Joint species trait-environmental model that explains the spatial patterns of invasive plants using a multivariate method (predictors considered related to land use, life-history traits, residence time, origin and human usage)	N/Y/N/N/N	$pr(\lambda_c E_i, S) = f(E_{i,1} + E_{i,2} \dots + S_{i,1})$ $Z = \sum_{i=1}^L pr(\lambda_c E_i, S)$	incorporates both species traits and environmental suitability, simultaneously. Has only been done using linear models. Standard errors also provided, but not on validation set
211	Trethowan et al. 2011	Plant (Pompe weed, <i>Campuloclinium macrocephalum</i>) and biological control agents (<i>Liothrips tractabilis</i> and <i>Cochylis campuloclinium</i>)	3	South Africa	Yes (AUC and misspecification error rates)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	
212	Trnka et al. 2007	Insect (European corn borer, <i>Ostrinia nubilalis</i> , <i>Hubner</i>)	1	Global	Yes (misspecification error rates)	Multi-generational phenology model based on occurrence data.	N/N/Y/N/N	$Z = f_X(E)$	E is degree days - looked at development times rather than populations per se. Not sure about dynamical aspect to model. Not sure where multi-generational component comes in
213	Vaclavik & Meentemeyer 2012	Plant pathogen (<i>Phytophthora ramorum</i>)	1	Western USA	Yes (AUC and standard deviation using jackknife cross-validation)	Potential distribution at different stages of invasion using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i, t)$	By stages he means time since invasion or nearness to equilibrium (denoted as t). - testing the issue of equilibrium distributions in SDMs
214	Vall-Isoera & Sol 2009	Birds	202	Global	Yes (misspecification error rates)	Generalized linear mixed models and hierarchical tree models to be used with occurrence and introduction data	N/Y/N/N/N	$Z = P^* = f_p(S, N)$	Actual info on propagule size, in addition to species traits
215	van klinken et al. 2009	Plants (<i>Parkinsonia aculeata</i>)	1	Australia	Yes (sensitivity analysis and r^2)	Species distribution model calibrated using both occurrence and abundance data	N/Y/Y/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i) f_X(E_i)$	
216	van Wilgen et al. 2009	Reptile and amphibians	67	USA (California and Florida)	Yes (AUC, kappa and misspecification error rates)	Climate matching model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^L pr(\lambda_c E_i)$	

217	Vanderhoed et al. 2009	Plants (Pepperweed, <i>Lepidium latifolium</i>)	1	USA (California)	Yes (likelihood ratio, Nagelkerke r ² , Hosmer and Lemeshow goodness of fit, % absent classified correctly, % present classified correctly)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
218	Veale et al. 2012	Mammals (stoats, <i>Mustela erminea</i>)	1	New Zealand	Yes (AUC)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
219	Venette & Cohen 2006	Plant pathogen (<i>Phytophthora ramorum</i>)	1	USA	Yes (sensitivity analysis and misspecification error rates)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
220	Verling et al. 2005	Zooplankton	+1	USA	Yes (standard errors)	Vector traffic model that does not simply estimate propagule pressure from total ship arrivals. Significant differences exist in (i) the frequency and volume of ballast water discharge among vessel types, (ii) the frequency of vessel types and routes (source regions) among recipient ports, and (iii) the transit success (survivorship) of zooplankton in ballast tanks among voyage routes,	Y/N/N/N/N	$Z = N = \sum_{i=1}^I \sum_{j=1}^{J_i} \sum_{t=1}^{T_{ij}} f_{ij}(t-f) * pr(R V)$	Uses amount of anthropogenic traffic in a pathway (ballast water) as a metric of risk. Takes into account mortality depending on source, and hence time. Also takes into account different releases of ballast water depending on vector (ship) type, with the assumption that ballast volume released relates to propagule pressure. The amount and type of traffic V also differs to destinations, and this is implicitly captured in different values of V _{ijk} . and pr(R v) which is specific to a given vector. No knowledge of uptake, or variability
221	Vicente et al. 2011	Native plant (<i>Ruscus aculeatus</i> L.) and introduced plant (<i>Acacia dealbata</i>)	2	North-West of Portugal	Yes (AUC and cross-validation)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
222	Villemant et al. 2011	Insect (Asian bee-hawking hornet, <i>Vespa velutina nigrithorax</i>)	1	South-Western France	Yes (AUC and cross-validation)	Species distribution models calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
223	Wang & Jackson 2011	Crustacean (<i>Bythotrephes longimanus</i>)	1	Canada (Ontario)	Yes (AUC and cross-validation)	Species distribution models calibrated using occurrence data, considered biotic interactions with predators	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
224		Plants (<i>Buddleja davidii</i>)	1	New Zealand	Yes (model evaluation with independent data)	Parameters from current species distribution were used to predict future distribution under climate change	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
225	Watt et al. 2008	Plant (broad-leaved paperbark, <i>Melaleuca quinquenervia</i>)	1	Global	Yes (misspecification error rates)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
226	Watt et al. 2011	Fungi (<i>Phaeocryptopus gaeummannii</i>) and plant (Douglas-fir, <i>Pseudotsuga menziesii</i>)	2	New Zealand	Yes (misspecification error rates)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
227	Weik et al. 2002	Plant (garlic mustard, <i>Alliaria petiolata</i>)	1	North America	Yes (Jaccard index and misspecification error rates)	Climate matching model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
228	Williams et al. 2008	Plant (orange hawkweed, <i>Hieracium aurantiacum</i>)	1	Australia	Yes (probabilities are stochastic. Uncertainty examined using hindcasts)	Dispersal constrained habitat suitability model that combines a habitat suitability index (developed from disturbance, site wellness and vegetation community parameters) with a phenomenological dispersal kernel that uses wind direction and observed dispersal distances	N/Y/N/Y/N	$N_{i,j} = \sum_{j=1}^J f_D(D_{i,j}, E)$ $Z = P_i^A = \sum_i pr(\lambda_c E_i) * pr(N_{i,j} > threshold)$	Uses dispersal kernel for probability of reaching an area in combination with habitat suitability. Like model 7, does not consider propagule pressure versus establishment relation. Here J are occupied sources, which result in temporal dynamics
229	Wilson et al. 2009	Fish	2	USA	Yes (standard errors/r ²)	Mayfield logistic regression on temperature variables and species traits (strain and mass-length ratio) on daily survival as a proxy for establishment success	N/Y/N/N/N	$Z = pr(\lambda_{c1} E_i, S) = f(E_{i1} + E_{i2}, ... + S_{i1} + S_{i2}, ...)$	Purely experimental, and looks at survival only
230	Wu et al. 2010	Mollusc (zebra mussel, <i>Dreissena polymorpha</i>)	1	USA (Wisconsin)	Yes (misspecification error rates)	Risk-based decision model based on occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	E was the habitat suitability index (HIS). Paper presented conceptual demographic model as well with management (CASM), but results were all based on HIS analysis. Also, not enough info on CASM to map
231	Yan et al. 2006	Plant (<i>Pinus radiata</i>)	1	Southwest China	Yes (misspecification error rates)	Climate matching model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
232	Yemshanov et al. 2009	Insect (<i>Sirex noctilio</i>)	1	Cross-border USA-Canada	Yes (misspecification error rates)	Bioeconomic Model based on occurrence and spread data. Propagule pressure considered	Y/Y/N/Y/Y	$N_i = \sum_{i=1}^I V_{i,j}$ $P_{i,j}^{A2} = f_D(D_i)$ $Z = \sum_{i=1}^I f(N_i) \sum_{j=1}^{T-i} \sum_{t=1}^{T-i} (P_{i,j}^{A2} - P_{i,j+1}^{A2}) * f_i(E_i)$	N used shipping as a proxy. N was rescaled to be a probability. Pr(A) was dependent on distance (D) to closest source only. It was dependent on host (Pine) characteristics, and climate. Combination of factors not explicit and was assumed. Time elements not indicated either, and assumed
233	Zalba et al. 2000	Plant (old man saltbush, <i>Atriplex nummularia</i>)	1	Argentina (Buenos Aires)	No	Species distribution model based on occurrence data and considering species germination, establishment, growth and reproduction.	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i, S)$	
234	Zenni et al. 2009	Plant (Kangaroo thorn, <i>Acacia paradoxa</i>)	1	South Africa	Yes (AUC and misspecification error rates)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
235	Zhu et al. 2012	Insect (Brown marmorated stink bug invasion, <i>Halyomorpha halys</i>)	1	Global	Yes (cross-validation, AUC and binary omission rate)	Species distribution model for both native and exotic species calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	
236	Zimmermann et al. 2011	Plant (<i>Rosa rubiginosa</i>)	1	Southern Argentina	Yes (AUC and explained variance)	Species distribution model calibrated using occurrence data	N/Y/N/N/N	$Z = Q = \sum_{i=1}^I pr(\lambda_c E_i)$	

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Appendix 3: Mapping examples of scoring RAs and quantitative RAs onto the TEASI model

Here, we walk through and compare specific examples of RAs (two scoring RAs and two quantitative RAs). We identify the general elements of RAs, and illustrate how they may be mapped onto the TEASI model structure (see Box 2 & 3 for model structure, and Table 1 for a glossary of notation). By detailing sample RAs, readers should be able to extrapolate to other papers examined, and use Appendix 1 & 2 as a summary of each of the 300+ RAs examined in this study.

Generally, each RA can be described by the components/sub-components that they include, the way that these components are combined (model structure), and the dependencies (surrogates) used to estimate them. For scoring approaches, the components and sub-components are framed in terms of the questions asked within a questionnaire, whereas in the quantitative RAs, they are framed as explicit variables included in a model. The individual questions/variables were mapped to match the closest counterpart in the TEASI model. Where relevant, we noted conditional dependencies (e.g., $F_I(S)$ is explicitly dependent on species traits, whereas $F_I(.)$ denotes unspecified dependencies). More generally, we also determined which model components (TEASI) each RA included (see Appendix 1 & 2). For model structure, in scoring RAs, the methods of combination are explained verbally (e.g., scores are summed), whereas in quantitative RAs, the model structure can be expressed more succinctly as an equation.

Example 1: EPPO Risk Assessment

The EPPO decision-support scheme (EPPO 2011) for quarantine pests is intended to be used to assess the potential importance of a particular pest for a clearly defined area. The scheme provides detailed instructions for the following stages of pest risk analysis: initiation, pest categorization, probability of introduction, potential economic consequences and pest risk management. The EPPO risk assessment firstly estimates the probability of the pest being introduced into the pest risk assessment area (its entry, establishment and spread) and secondly makes an assessment of the likely environmental and economic impact if that should happen. The evaluation is based on the replies to a series of questions, expressed as the choice of an appropriate phrase out of a set of five alternatives (e.g. very unlikely, unlikely, moderately likely, likely, very likely).

To map the EPPO RA (see Table 2 and Appendix 1, Model ID 7), we converted questions into a common notation, based on the TEASI model. The EPPO assessment starts with the identification of all pathways that allow entry or spread of a pest. For each pathway, two questions relate to the number of propagules O transported in this pathway: the probability of a pest to be associated with this pathway and the volume of movement along the pathway. Since they are pathway-specific, they were mapped as $O(v,.)$. Two questions are about population dynamics during transport, namely the likelihood of survival and the likelihood to increase in numbers during transport $f_g(t-t_L, v,.)$, and another two questions about the likelihood to be released in a suitable habitat, i.e. that the release will support entry of the pest $pr(R/E, v)$. After having answered these 6 questions, one is then asked to assess the overall probability of entry of each pathway and the overall probability of all pathways. To answer these questions, no specific guidance is given as to how combine the specific questions into an overall score or probability.

The next part of the EPPO RA assesses in 12 questions the suitability of the release site for establishment, or in other words, the overall probability of establishment in a given environment $pr(l_C/E,.)$. The questions asked for host plants and suitable habitats, alternate hosts and other essential species, climatic suitability, other abiotic factors, competition and natural enemies, the

spatial distribution of (alternate) hosts or suitable habitats. This is followed by general questions on the pest's reproductive strategy $r(\cdot)$, its adaptability (not mapped; this is not a part of the full model), and the species's history as a pest elsewhere, i.e. its distribution outside its native range, which was mapped as the general likelihood to establish $pr(l_C/\cdot)$. At the end of this part, again an overall assessment of the establishment probability should be made. Again, the connection to the previous questions is not made explicit in the EPPO scheme.

The probability of spread is assessed in 4 questions. Three questions are about the rate of spread by natural and human means, without specifying any dependencies like species traits or environmental characteristics $f_D(\cdot)$, followed by one question about the expected area invaded after 5 years $Q_T(\cdot)$.

Finally, there are 8 questions about general economic impact (crop yield and/or quality, production costs, loss in export markets, transmission of diseases), 4 questions about environmental impact (without giving any details) and 2 questions about social impact (again, without details), all without qualifiers $f_I(\cdot)$.

The EPPO RA covers all TEASI components except local abundance. Uncertainty (only epistemic) should be documented for each question (on a scale of 3: low, medium, high) for transparency and identification of future research needs. However, there is no guidance on how to incorporate uncertainty into assessing risk. Likewise, no guidance is given how to combine scores of individual questions to an overall risk score.

Example 2: Australian Weed Risk Assessment

The Australian WRA (Pheloung *et al.* 1999) includes 49 questions divided into three larger sections, which are further divided into a total of eight sub-sections. The questions are in most cases answered yes/no, which typically yield a score of 0 for no and 1 for yes, but in some questions different scales are used (e.g. 0 vs. -3, -1 vs. 1, and 0 vs. 5). In a few questions the score scale is 0-2 (low, intermediate, high). It is not obligatory to answer all the questions, but a sufficient number of questions in each category need to be answered (2, 2 and 6 subsequently). This accounts for uncertainty in the sense that if the answer to some question is not known, it is not necessarily required in order to complete the assessment. The aggregate score is achieved by summation of the individual question scores. Based on their aggregate score, the species are then categorized to three risk categories: accept, reject, or evaluate further, based on pre-determined score boundaries. Each question is also denoted as relating to agriculture, environment, or both. Therefore it is also possible to aggregate the score to assess the importance of the species as, say, an agricultural weed.

The mapping is discussed below in the order of the full invasion model (see Table 2 and Appendix 1, Model ID 43).

There are eight questions mapped as estimating the species traits (S). These do not link directly to any particular component (TEASI) of the full model --- rather they are questions that attempt to estimate whether the species in general makes a good invader. The questions relate to shade tolerance, growth on infertile soils, being aquatic, grass plant, nitrogen fixing woody plant or geophyte, toleration of mutilation, cultivation or fire, and whether the propagules are buoyant. A further two questions were mapped related to environmental interactions ($f(E)$): whether the species requires specialist pollination, and whether effective natural enemies are present in the area. These could not be directly linked to any specific invasion model component either.

The seven questions mapped as relating to the Transport component (and v more specifically) were related to propagule dispersal by people intentionally and unintentionally, as a produce contaminant, by wind, by bird and by other animals either internally or externally. Propagule buoyancy was mapped as a species trait, although it can relate to water dispersal. The dispersal questions can be interpreted as belonging to either Transport or Spread phase of the invasion – we interpreted them as covering both.

Two questions were mapped related to propagule pressure (N): Is the species highly domesticated, and does the species have a history of repeated introductions outside its natural range. These characteristics may be related to either Transport or Establishment phase. In the case of the Australian WRA we did not have to assess whether they link to one or another (or both), as there were other questions relating to both Transport and Establishment, ensuring that the model covered these components.

Five questions were directly mapped relating to establishment probability ($pr(\lambda_C)$), and therefore to the Establishment component of the model. There were two questions with unspecified dependencies: has the species become naturalised where grown, and is it naturalised beyond its native range. There was a further question with environmental dependency ($pr(\lambda_C/E)$) (Does the species have broad climate suitability), and two questions with both environmental and species traits as dependencies ($pr(\lambda_C/E,S)$) (Is the species suited to Australian climates, and is the species native or naturalised in regions with extended dry periods).

Eight questions were mapped to the reproductive potential of the species (r), and therefore presenting the Abundance component in the model. There was one question with unspecified dependency (Is there evidence of substantial reproductive failure in native habitat), and seven questions that were dependent on species traits (r/S).

The Impact component was also strongly present, as the Australian WRA includes 15 questions on impacts of the species, with 10 of those especially interpreted as being related to species traits ($F_I(S)$). The five questions that were interpreted as having unspecified dependencies ($F_I(.)$) were related to whether the species has weedy races, and whether it is regarded as a weed in different contexts (e.g. garden, agriculture, environment).

The management question, mapped as $\alpha_C M_C$ (i.e. related to control and control effectiveness of the species), was related to the ability of herbicides to control the species. This one question results in the model mapped as considering the management of the species. The final remaining question regarding the quality of climate match data was mapped as “other”.

Therefore, all the five TEASI components were considered to be covered by the Australian WRA. In comparison with the EPPO approach, the Australian WRA includes a systematic method for aggregating the scores to form a final score. The EPPO system on the other hand includes a more careful consideration of uncertainty related to the assessment, although it provides no guidance on how that should affect the final evaluation. The overall scope of the questions in both schemes was relatively similar.

Example 3: Species distribution models

Species distribution models have seen wide development within invasion biology, with numerous statistical approaches linking environmental characteristics with the probability of establishment (Table 3). While often many different predictor variables are used (e.g., hydrology and water

chemistry, Appendix 2, Model ID 2, climate, edaphic data, Model ID 4, etc), and a number of different statistical approaches are available (e.g., Maxent, GARP, Neural Networks, Generalized Linear Models), in essence they all fit environmental predictors of establishment to occurrence data, and are thus denoted as $pr(\lambda_C | E_i)$. Thus, this approach yields much more in depth predictive ability compared to the scoring approaches described above, but is much more abbreviated in scope, examining only a single subcomponent in the TEASI model (and is therefore much simpler to describe).

Given that these models are often based on GIS environmental layers, a simple extension of the fitting technique allows a prediction of spatial extent (Q), which we denote in our estimate of risk as

$Z = Q = \sum_{i=1}^L pr(\lambda_C | E_i)$. Here, E_i typically refers to abiotic environmental traits, but can also include

biotic factors such as native species presence (e.g., Appendix 2, Model ID 75). Moreover, researchers have also included aspects related to transport into these models, which thus enters our notation as well. For instance, Appendix 2, Model ID 2 included a measure of vector traffic and is denoted as $Z_i = pr(\lambda_C | E_i, V_i)$; Model ID 4 included a measure of distance from known source

locations and is denoted as $Z = Q = \sum_{i=1}^L pr(\lambda_C | E_i, D_{i,j})$. In these ways, multiple TEASI components can be included into the same statistical structure, although with subtle ramifications, as discussed in the next paragraph.

Note the presence of subscript i indicates that spatial locations were considered. Note also the absence of t , indicating that temporal dynamics were not considered (i.e., these models implicitly assume equilibrium conditions, although joint models are becoming more popular). Further note that we are treating these as probabilities, as closest analogue in the TEASI model, but discrepancies may exist because of biased sampling, use of pseudoabsences and base-rate effects.

Example 4: Species trait models

Species trait models help predict whether a given species will be able to invade a system, using a variety of statistical techniques (e.g., discriminant analysis, generalized linear models, boosted regression trees), and a variety of species traits (e.g., brain size, ecological niches breadth, Appendix 2, Model ID 214; reproductive mode, phenology, invasion history, Model ID 177). These models were mapped to the TEASI framework as $Z = P^c = f_{p^c}(S)$; P^c indicates the probability of the system as a whole will be invaded. f_{p^c} indicates that it is some function of species trait (S) (to determine P^c).

Analogously to species distribution models, these provide quantitative predictions, but are also typically more limited in scope compared to scoring methods. However, note that species traits have been used to predict other aspects of the TEASI model besides establishment, and were denoted as such (e.g., to predict average abundance, $Z = \bar{X} = f_X(S)$, Model ID 85; to predict impact, $Z = f_I(S)$, model ID 105), although they are typically not integrated. As another comparison with species distribution models, note the absence of subscripts i and t , indicating that spatial and temporal heterogeneity and dynamics were not considered. Although these differences in notation are subtle, they are important as they dramatically change the implications of the models.

As another subtle variation in notation between studies, the predictive dependencies could also extend beyond species traits, and be incorporated into the same statistical method. For instance, environmental conditions (e.g., climate matches between native and introduced ranges) could be included in the model (denoted as $Z = P^c = f_{p^c}(E, S)$, e.g., Model ID 11). As a subtle judgment call, environmental tolerances were treated as a species trait (e.g., minimum temperature, Model ID 115; temperature variability in native range, Model ID 26). Researchers also included factors related to transport (e.g., human usage factors relate to vector movements, denoted as $Z = P^c = f_{p^c}(S, f(V))$, Model ID 26). As mentioned above, these all lacked temporal and spatial resolution.

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