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The conservation value of germplasm stored at the Millennium Seed Bank, Royal Botanic Gardens, Kew, UK

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Abstract The Millennium Seed Bank (MSB) Partnership, developed and managed by the Royal Botanic Gardens, Kew (RBG Kew), conserves propagules primarily from orthodox seed-bearing wild vascular plants. It is the largest ex situ conservation programme in the world, currently involving 96 countries and territories. Where possible, seeds are collected and conserved in the country of origin with duplicates being sent to RBG Kew's MSB for storage. In this paper we assess the conservation value of the germplasm stored at the MSB using both quantitative and qualitative methods. The MSB holdings represent a high quality, rich biological resource. Substantial and unique taxonomic diversity exists amongst the collections, representing 365 families, 5813 genera, 36,975 species and 39,669 taxa conserved. The collections cover a wide geographic range, originating from 189 countries and territories, representing all nine bio-geographic regions and all 35 biodiversity hotspots. The collections possess significant natural capital and population value: 32% of taxa, representing 49% of collections, have at least one identified use to humans; and 74% of taxa, representing 78% of collections, are either endemic, endangered (nationally or globally) and/or have an economic, ecological, social, cultural or scientific value. While 10% of taxa, representing > 8% of collections, are either extinct, rare or vulnerable to extinction at the global and/or national level, 20% of taxa, representing 13% collections, are endemic at the country or territory scale. Over the 17-year period since 2000 at least 11,182 seed

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samples, representing 12% of taxa and 8% of collections, have been distributed globally for conservation, research, education and display. This analysis highlighted collection gaps in MSB holdings in relation to their geographic representativeness, the taxonomic diversity of large families and genera of angiosperms, and coverage of threatened taxa. Further analysis across the entire MSB Partnership is required to underpin future collection activities and maximize the usefulness of collections.

Keywords Ex situ conservation · Seed banking · Vulnerability to extinction · Rarity · Uniqueness · Irreplaceability · Natural capital value · GSPC

Introduction

The long-term storage of germplasm in the form of seeds is central to an integrated in situ and ex situ conservation strategy, and together with botanic gardens is one of the most widespread and valuable approaches to ex situ plant conservation. Crop germplasm has been conserved in seed banks for over 60 years and the Royal Botanic Gardens, Kew (RBG Kew) has followed this work since the late 1960s, adapting the techniques for the conservation of 'wild' species. The Millennium Seed Bank (MSB) Project established in 1995 is a plant conservation partnership between RBG Kew and organizations both within the UK and across the globe. It is the largest ex situ plant seed conservation program for wild species in the world (Smith et al. 1998). Seed banking remains a key part of RBG Kew's Science Strategy (RBG Kew 2015), while also contributing to Target 8 (at least 75% of threatened plant species conserved in ex situ collections, preferably in the country of origin, and at least 20% available for recovery and restoration programs) and Target 9 (70% of the genetic diversity of crops including their wild relatives and other socio-economically valuable plant species conserved, while respecting, preserving and maintaining associated indigenous and local knowledge) of the Global Strategy for Plant Conservation (CBD 2012).

The MSB Project's International Program successfully conserved 10% of the world's wild orthodox seed-bearing flora between 2000 and 2010. The MSB Partnership (MSBP, 2011 to present) currently involves 96 countries and territories, with active projects in 54 countries. The main aim of MSBP is to continue safeguarding plant diversity worldwide with a focus on plants most at risk and most useful for the future, while addressing global challenges for food security, sustainable energy, loss of biodiversity and climate change. With one in five vascular plant species currently threatened with extinction (RBG Kew 2016), the need for such conservation measures has never been greater.

Traditional seed banking focuses on the storage of orthodox seeds, those that can be dried (to 15% equilibrium relative humidity) and stored at low temperatures (− 20 °C), in air-tight containers to maximise their longevity. For every 10% decrease in equilibrium relative humidity and 5 °C drop in temperature the lifespan of orthodox seeds is doubled (Harrington 1960). Thus, under conventional storage conditions seeds can be expected to live for 10–100 s, if not 1000 s, of years, the longevity being dependent on the species and seed traits. The ability to store a large diversity of germplasm in a small space at relatively low cost makes seed banking a practical and attractive tool for plant conservation.

There are more than 1750 seed banks in the world, the majority of which conserve crop diversity (Hay and Probert 2013). Since the Global Strategy for Plant Conservation (GSPC) was adopted by the Convention on Biological Diversity (CBD) in 2002, the number of

ex situ conservation facilities for wild species has grown dramatically, but little information is available about the extent to which plant species are appropriately represented in ex situ collections (Godefroid et al. 2011; Cibrian-Jaramillo et al. 2013; Rivière and Müller 2017; Teixido et al. 2017). This study assesses the conservation value, both qualitative and quantitative, of germplasm conserved in MSB in terms of its: (1) biological status; (2) taxonomic diversity (Ojeda et al. 1996); (3) geographic representativeness (Godefroid et al. 2011; Cibrian-Jaramillo et al. 2013; Kricsfalusy and Trevisan 2014); (4) vulnerability to extinction (Rivière and Müller 2017); (5) uniqueness and irreplaceability (Vane-Wright et al. 1991; Ojeda et al. 1995; Isaac et al. 2007); (6) natural capital value; (7) population value; (8) germplasm quality; and (9) use for conservation, research, education and display. Our analysis excludes germplasm conserved under the MSBP in the country of origin and not duplicated in MSB.

Materials and methods

Data for collections were extracted from the MSB Seed Bank Database (SBD) on 31 March 2017. SBD contains in-depth information on seed collections including their heredity, taxonomic identification, geographic origin (bio-geographic region, country, major and minor administrative divisions, locality, geographic coordinates and altitude), habitat (type, associated species, threats, land use, geology, slope, aspect and soil texture), sampled population (abundance, vulnerability, number of plants found, and sampled and area sampled), ethnobotanical uses, quantity, viability, germination, regeneration, propagation and use of collections for conservation, research, education and display.

Biological status

In determining the biological status of a collection (wild or cultivated origin), heredity, geographic origin, habitat, taxonomy and regeneration data were used. Collections originating from natural or semi-natural habitats were considered as ‘wild’ and those originating from cultivated habitats (e.g. orchards, home gardens and botanic gardens) and propagation or regeneration activities were considered as ‘cultivated’ (Alercia et al. 2012).

Taxonomic diversity

Species and taxon abundance and taxonomic composition in terms of families and genera were estimated using the current plant identification status of collections. The representativeness of angiosperm (flowering plant) families was estimated as the percentage of genera and species conserved at the MSB within the family against the total number of naturally occurring genera and species described for the family. Likewise, the representativeness of angiosperm genera was estimated as the percentage of species conserved in the MSB within each genus against the total number of naturally occurring species described for each genus. The total number of accepted genera and species described per family and accepted species described per genus were taken from The Plant List (2013) considering the confidence level and review status of names. If data was not available in The Plant List (2013), or the taxonomy of SBD data was in doubt, problematic taxa were excluded from the analysis. Therefore, angiosperm analysis was restricted to 324 families and 5341 genera.

Geographic representativeness

Geographic origin data were used to yield digital maps and total numbers of taxa and collections originating from bio-geographic regions and countries using ArcInfo and ArcView software. If coordinates were wrong or missing, new coordinates were assigned based on the locality description for the collection where possible. Cultivated collections inherited the geographic origin of the wild plant population from which they were propagated or regenerated. Geographic data were analysed to describe geographic coverage and gaps in geographic representation including representation of MSB collections from the 35 Biodiversity Hotspots described by Conservation International (Mittermeier et al. 2011, http://www.conservation.org).

Vulnerability to extinction

In order to verify whether MSB taxa are extinct, rare or vulnerable to extinction in the wild, SBD taxonomic identifications were cross checked with (1) IUCN (2016) for the global scale; and (2) the National Red List (2016) for the national scale. In addition, Walter and Gillett (1998) was used to identify globally rare taxa and IUCN (2016) was used to identify taxa evaluated with lower risk of extinction at global scale.

The representativeness of taxa that are extinct, rare or vulnerable to extinction at global and national scales was assessed using IUCN Red List categories: extinct (EX); regionally extinct (RE); extinct/endangered (EX/EN); extinct/vulnerable (EX/VU); extinct in the wild (EW); critically endangered (CR); endangered (EN); endangered/vulnerable (EN/VU); vulnerable (VU); rare (R or RR); intermediate (I) and relict. The National Red List (2016) uses both IUCN and non-IUCN criteria (e.g. threatened, critical or declining) for categorizing threatened taxa. In our analysis, all non-IUCN categories from the National Red List (2016) were treated separately and presented as 'Other Threatened' taxa. Taxa categorized under lower risk of extinction in IUCN (2016) include a number of subdivisions, namely: near threatened (NT); lower risk/near threatened (LR/nt); lower risk/conservation dependent (LR/cd); lower risk/least concern (LR/lc); and least concern (LC).

For vascular plants, IUCN (2016) lists global assessments for 23,392 taxa, 12,564 (54%) of which are listed as either extinct or vulnerable to extinction, 9028 (38%) with lower risk of extinction and 1800 (8%) with insufficient information for assessment. The National Red List (2016) lists 24,969 taxa as either extinct, rare or vulnerable to extinction in at least one country. Walter and Gillett (1998) list 14,998 vascular plant taxa as globally rare. To be considered as representative of nationally rare or threatened taxa, MSB collections need to originate from the country where the taxon is declared as rare, extinct or vulnerable to extinction.

Uniqueness and irreplaceability

The rarity of a taxon has been described using its geographic distribution (restricted-range endemics) and/or evolutionary distinctiveness (taxonomically distinct) (Ojeda et al. 1995; Isaac et al. 2007; Cibrian-Jaramillo et al. 2013; Kricsfalusy and Trevisan 2014). Rare taxa will have a greater risk of extinction than common ones (Johnson 1998; Matthies et al. 2004) and are considered as unique and irreplaceable in terms of their restricted distribution pattern and/or taxonomic distinctiveness.

The most commonly used measure of uniqueness and irreplaceability in conservation is plant endemism (Stattersfield et al. 1998). Geographic distribution of plant endemism is highly taxon-dependent (Swenson et al. 2012) and describes the ecological state of a taxon being unique to a defined geographic location, such as an island, nation, country or other defined zone, or habitat type. Organisms that are indigenous to a place are not endemic to it if they are also found elsewhere.

It has been argued that maximizing evolutionary distinctiveness and phylogenetic diversity should be key components of conservation effort as the extinction of a species in an old, monotypic or species-poor clade would result in a greater loss of biodiversity than that of a young species with many close relatives (Mace et al. 2003). Building the phylogenetic diversity of ex situ collections will strengthen their capacity for use in response to biodiversity loss (Griffiths et al. 2015). Therefore, we included both rarity criteria in the analysis.

We applied the concept of endemism at the country or territory scale to identify geographic rarity. A list of endemic or near endemic taxa was compiled using a variety of reference lists including IUCN (2015, 2016), Walter and Gillett (1998), National Red List (2016), WCSP (2015) and SBD. As compiling a full list of endemic taxa is not practically feasible, we have utilised taxon distribution data from the MSB Species Prioritisation Tool (unpublished, Liu and Kenney) to identify endemic taxa, but we treated these designations with a low degree of confidence.

Indices of taxonomic diversity (Vane-Wright et al. 1991) are based on phylogenetic trees and provide a means of identifying distinct species (Ojeda et al. 1995). Phylogenetic studies demonstrate that evolutionary distinctiveness (ED) is derived from a few branches near the tips (i.e. those shared with few other species) and that no ED is gained in clades above ~ 180 species (Isaac et al. 2007). Taxonomic diversity was estimated by evaluation of the singularity of species making up communities. The taxonomic singularity of each species is directly related to the number of co-generic species in a given geographic range (Ojeda et al. 1995, 1996) and the inverse of the average number of species per genus in the community has been proposed as a simple index for assessing taxonomic distinctiveness at the community level, and quantifying its conservation value.

Ideally, the index for taxonomic singularity for the MSB taxa should be calculated from the number of species conserved for a given genus as an average of number of species normally found in a given geographic area from where the seed collections were sampled. For the purpose of this study, we used a rapid assessment method to identify the representation of less diversified and evolutionary distinct genera and families in MSB holdings (Ojeda et al. 1995). The number of genera per family and the number of species per genus were obtained from The Plant List (2013), considering the confidence level and review status of names, for angiosperm families to identify less diversified families (with up to 10 genera) and genera (with up to 10 species) and then their representation at the MSB was measured and expressed as taxonomic singularity to identify taxonomic rarity. Phylogenetic diversity of the MSB legume collections has been assessed in a separate study (Griffiths et al. 2015).

Natural capital value

Any taxon that has an identified use to humans (economic, ecological, social, cultural or scientific) was considered as a taxon with a natural capital value. We compiled 31,413 taxa with an identified use to humans from five reference sources and then used this list as a reference to verify the natural capital value of MSB taxa: (1) RBG Kew's SEPASAL—Survey of Economic Plants for Arid and Semi-Arid Lands Database (RBG Kew 1999); (2) RBG Kew's Economy Botany Collection Database (http://apps.kew.org/ecbot/search); (3) RBG

Kew's Medicinal Plants Names Service Portal (https://www.kew.org/science/data-and-resources/tools-and-services/medicinal-plant-names-services); (4) Germplasm Resources Information Network (https://www.ars-grin.gov/npgs/aboutgrin.html); and (5) Harlan and de Wet Crop Wild Relative Inventory (http://www.cwrdiversity.org/checklist/). In addition, the ethnobotanical uses listed by collectors for MSB collections were extracted from SBD and used as a further resource. As these were not supplemented with a reference, a low degree of confidence was assigned.

Crop wild relatives (CWR) are wild plant species that share a common ancestor with cultivated crop plants of socio-economic value, such as human food, animal forage and fodder crops, etc. (Fielder et al. 2015). A broad definition of a CWR is any taxon belonging to the same genus as a crop (Maxted and Kell 2009). A narrower definition of a CWR was described as "a wild plant taxon that has an indirect use derived from its relatively close genetic relationship to a crop" (Maxted et al. 2006). Three concepts are used to identify how close the relationship to a crop is: (1) Harlan and de Wet (1971) describe the genetic relationship of cultivated plants using the Gene Pool concept at three levels, this is the most commonly used concept, it is relatively objective and widely accepted (Vincent et al. 2013); (2) for the majority of crops the genetic relationship among species remains unknown, and in these cases the Taxon Group concept is used that assumes taxonomic classification is strongly linked to genetic relatedness (Maxted et al. 2006); (3) the provisional gene pool is used where there is no known gene pool concept and taxonomic treatments lack sub-generic information, but there is evidence that the crop and related taxa can be crossed (Vincent et al. 2013).

Based on different definitions, we analysed CWR taxa at two levels: (1) broad definition of CWR (sensu lato) based on any taxa in the same genus as the crop, this was verified against the consolidated list of 99 crop genera cited in Maxted et al. (2013) by combining 51 food crop genera and 81 forage crop species published by the International Treaty on Plant Genetic Resources for Food and Agriculture (FAO 2001) and 77 crop genera listed in Groombridge and Jenkins (2002); and (2) narrow definition of CWR (sensu stricto) based on any taxa specifically listed as CWR using the gene pool, taxon group and provisional gene pool concepts and was verified against 4229 taxa compiled from the Harlan and de Wet Crop Wild Relative Inventory and Germplasm Resources Information Network.

Population value

We considered taxa that are identified as endangered globally and/or nationally, endemic or near endemic at country or territory level or with a natural capital value as '3E taxa' (either endemic, endangered or economically important). Taxa that contribute to all three components were considered as significant taxa or sensu stricto 3E taxa with a high population value. Taxa that contribute to any component were considered as sensu lato 3E taxa. We use 3E taxon status to evaluate the population value of MSB holdings.

Germplasm quality

To be a valuable long-term resource, a collection of germplasm needs to: have accurate plant identification; be genetically representative of the species, population and individual sampled; have high viability with acceptable longevity; contain sufficient germplasm to supply intended uses; and be acquired with all consents and data to facilitate intended users (Way 2003). The key principles at the core of gene bank operations are the preservation

of germplasm identity, maintenance of viability and genetic integrity, and the promotion of access (FAO 2014). In our study, the germplasm quality was analysed in terms of: (1) availability of key data in SBD for taxonomy (plant name), geography (bio-geographic region, country, geo-coordinates and altitude), habitat and population (number of plants found, and sampled and area sampled); (2) current status of plant identification (verified or unverified); and (3) seed quality, quantity and viability.

A sample of seeds (usually 10–50) from most MSB collections are either x-rayed or cut-tested to estimate the number of full, empty and infested seeds. These data are then used as a rapid assessment method to estimate the current number of potentially viable seeds in the whole collection. The true viability of collections (germination and viability percentage) is estimated from routine germination tests throughout the life cycle of the collections. In our study, to identify seed quality, quantity and viability, collections were categorized according to their current number of potentially viable seeds (estimated using seed weights and x-ray or cut-test results of dry seeds) and germination (germinated seeds out of seeds sown, discounting empty and infested seeds) and viability (germinated seeds plus non-germinated seeds that appeared fresh and full when cut-tested) percentage results for the most recent round of germination tests carried out at the MSB.

Use of collections

RBG Kew makes use of the seed collections stored at the MSB in research that both furthers our understanding of seed banking, and also seed biology and species conservation. Subject to terms and conditions, MSB collections with sufficient seed quantity are available to bona fide individuals from recognized organisations around the world for use in non-commercial activities in conservation, research, education and display. These collections are publicised through RBG Kew's MSB Seed List (http://apps.kew.org/seedlist/) and the Genesys website (https://www.genesys-pgr.org/wiews/GBR004), from which a small sample of germplasm can be requested by a third party. In addition, a range of seed biological trait data (storage behaviour, germination, weights, oil and protein content, dispersal and morphology) derived from MSB collections are published in RBG Kew's Seed Information Database (RBG Kew 2017, http://data.kew.org/sid/) and Try Plant Trait Database (Kattge et al. 2011, https://www.try-db.org/TryWeb/Home.php) for use in research and conservation. Although the use of physical collections and associated data are not fully documented, we used seed distribution data from January 2000 to March 2017 to assess the use of collections under four broad categories: conservation, research, education and display. Compiling a full list of scientific outputs and publications generated from MSB collections and associated data is beyond the scope of our study.

Data analysis

In order to maintain taxonomic consistency, plant Latin names and their taxonomic status (e.g. accepted, synonym, unassessed, unresolved or illegitimate) were carefully addressed in the analysis by checking plant names with the WCSP (2015), The Plant List (2013), and The International Plant Name Index (2015). If the data were not available in these resources other online sources, such as Taxonomic Name Resolution Service (http://tnrs.iplantcollaborative.org) and Tropicos.org (http://www.tropicos.org), were used.

There is variability in the precision and accuracy of data both in SBD and reference data sources mainly due to variation in the use of plant names and their taxonomic status.

To address these concerns, prior to the main analysis, we employed an evidence-based approach matching MSB plant names with IUCN (2015) to investigate the accuracy level of estimates. Based on the results (not shown), to reduce under- or over-estimating the number of plant taxa conserved in the MSB under the different criteria applied in the analysis, the following calculation strategy was employed. When matching SBD plant names with reference data sources:

(a) The minimum value is used to describe the number of SBD taxa that match directly with reference lists. These values are reported with a high degree of confidence but are treated as an underestimate of overall representativeness.
(b) The maximum value is used to describe the number of taxa that match directly or indirectly with reference lists based on any taxonomic link. These values are reported with a low degree of confidence and are treated as an overestimate of representativeness. There are multiple examples of this: (i) SBD plant name with 'accepted' taxonomic status matches to a 'synonym' on an external list; (ii) SBD plant name with 'accepted' taxonomic status, but a synonym of it matches to a plant name which is 'accepted' on an external list; (iii) SBD plant name with 'accepted' taxonomic status and a synonym of it matches to a synonym on an external list; and (iv) SBD plant name with 'synonym' taxonomic status and the 'accepted' name of it matches to an 'accepted' name on an external list. If a synonym's accepted plant name has already been matched to an accepted plant name then it is not included again to avoid double counting.
(c) The median is the middle value between the minimum and maximum values, and is used as the total figure reported and for calculating percentages. This ensures neither under- nor over- reporting.

Results

Biological status

Of the 82,556 collections conserved in MSB, 75,749 (91.8%) originated from material collected in natural or semi-natural habitats (wild collections) and 6807 (8.2%) originated from either cultivated habitats or propagation and regeneration activities in the UK or elsewhere (cultivated collections).

Taxonomic diversity

The majority of the collections (99.9%) conserved are seeds and only 0.1% are either spores or dormant organs (e.g. bulbils). MSB collections represent 365 families, 5813 genera, 36,975 species and 39,669 taxa (Table 1). Species of angiosperms contained within the Plant List (2013) belong to 405 families and 14,559 genera. About 82% of these families and 39% of these genera are represented in MSB holdings. Fifty percent of MSB collections originated from 355 families, 5558 genera, 30,956 species and 32,747 taxa, whilst the other 50% originated from 10 families, 255 genera, 6019 species and 6922 taxa (Appendix 1; Table 2). Fifty percent of species originated from 5476 genera, whilst the other 50% of species originated from 337 genera (Supplement 1). All the naturally occurring angiosperm genera are conserved for at least 93 families, and all the naturally occurring

Table 1 Overview of taxonomic diversity of MSB collections in terms of number of families, genera, species and taxa conserved under different vascular plant groups

Vascular plants	Collections	Families	Genera	Species	Taxa[a]
Angiosperms—dicots	65,101	267	4598	29,420	31,678
Angiosperms—monocots	16,277	64	1125	7212	7609
Gymnosperms	963	10	52	272	308
Lycophytes	15	3	6	10	11
Pteridophytes	93	20	32	61	63
Unknown	107	1			
Total	82,556	365	5813	36,975	39,669

Both wild and cultivated collections are considered

[a]Includes species, subspecies, varieties, etc.

angiosperm species conserved for at least 20 families (Appendix 1) and 949 genera (Supplement 1). For angiosperm families with one to 50 naturally occurring genera or species, more than 50% of genera are conserved for at least 150 families out of 265 and more than 50% of species conserved for at least 28 families out of 121 (Appendix 1; Table 3). Of the 12 families with many naturally occurring genera (> 200), three of them (Malvaceae, Leguminosae and Lamiaceae) each have more than 50% of genera conserved, and of the five families with a large number of naturally occurring species (> 10,000), three of them (Leguminosae, Poaceae and Compositae/Asteraceae) each have more than 10% of species conserved (Table 3; Appendix 1).

Out of the 5341 angiosperm genera analysed (Supplement 1), only 536 genera have more than 100 naturally occurring species, and these are underrepresented in MSB collections, except for nine genera which have at least 50% of their naturally occurring species conserved: *Acacia* (51%, 717 out of 1393); *Eucalyptus* (88%, 722 out of 822); *Melaleuca* (80%, 212 out of 265); *Plantago* (56%, 89 out of 158); *Lachenalia* (51%, 59 out of 115); *Terminalia* (55%, 60 out of 109); *Gastrolobium* (51%, 54 out of 105); *Pultenaea* (62%, 64 out of 104); and *Protea* (58%, 59 out of 101).

Geographic representativeness

Collections originating from all nine bio-geographic regions and 189 countries and territories are stored at the MSB (Table 4). The greatest number of collections and taxa originated from Africa and the lowest numbers from the Pacific bio-geographic region. The highest number of collections were collected in Australia (11,563), UK (6779), South Africa (5858), USA (4682), Madagascar (3692), Mexico (3332), Israel (3094), Italy (3021), China (2355) and Kenya (2225) (Fig. 1). Gaps in the geographic representation of collections were Belarus, Benin, Republic of Congo, Djibouti, Equatorial Guinea, Eritrea, Iceland, Kosovo, Latvia, Moldova, Mongolia, North Korea, Paraguay, Timor-Leste, Togo, Pacific Islands (American Samoa, Cook Islands, Guam, Kiribati, Marshall Islands, Micronesia, Nauru, Niue, Northern Mariana Islands, Palau, Samoa, Tokelau, Tuvalu), Middle Eastern States (Bahrain, Kuwait, Qatar) and Caribbean Islands (Antigua and Barbuda, Barbados, Saint Kitts and Nevis, Saint Vincent and the Grenadines).

MSB collections originating from the 35 biodiversity hotspots (Mittermeier et al. 2011) are provided in Appendix 2 and illustrated in Fig. 2. They represent 53% of MSB collections

Table 2 The most represented angiosperm families, genera and species in MSB seed collections

Families	Collections	% Collections	Genera	Collections	% Collections	Species[a]	Collections	% Collections
Leguminosae	13,092	15.86	*Vicia*	1950	2.36	*Vicia sativa*	497	0.60
Asteraceae/Compositae	8486	10.28	*Acacia*	1538	1.86	*Lathyrus aphaca*	222	0.27
Poaceae	6184	7.49	*Lathyrus*	1345	1.63	*Pennisetum violaceum*	196	0.24
Lamiaceae	2453	2.97	*Eucalyptus*	1057	1.28	*Vicia hybrida*	165	0.20
Rosaceae	2283	2.77	*Silene*	853	1.03	*Daucus carota*	161	0.20
Cyperaceae	2266	2.74	*Trifolium*	754	0.91	*Lathyrus sativus*	158	0.19
Mytaceae	2110	2.56	*Carex*	687	0.83	*Acacia nilotica*	140	0.17
Orchidaceae	2028	2.46	*Hieracium*	608	0.74	*Vicia narbonensis*	130	0.16
Malvaceae	1899	2.30	*Medicago*	564	0.68	*Vicia peregrina*	130	0.16
Caryophyllaceae	1886	2.28	*Cyperus*	499	0.60	*Neonotonia wightii*	125	0.15

[a]Includes subspecies, varieties, etc.

Table 3 The representativeness of angiosperm families and genera conserved at MSB

Naturally occurring genera and species			Total number of families or genera for which the naturally occurring genera or species have been conserved at the MSB in percentage classes				
Plant List (2013)		MSB holdings	% Conserved at MSB	1–25%	26–50%	51–75%	> 75%
Number of genera per family							
1–50	344	265	Families	33	82	49	*101*
51–100	29	28		5	15	8	0
101–150	12	12		2	4	6	0
151–200	7	7		3	3	1	0
> 200	12	12		2	7	3	0
Total	404	324		45	111	67	101
Number of species per family							
1–50	191	121	Families	46	47	6	22
51–100	40	34		26	6	2	0
101–500	69	66		58	7	0	0
501–1000	34	34		32	2	0	0
1001–5000	57	57		53	2	1	0
5001–10,000	6	5		5	0	0	0
> 10,000	5	5		5	0	0	0
Total	402	322		225	64	9	22
Number of species per genera							
1–50	13,348	4394	Genera	1686	1384	323	1001
51–100	552	411		322	71	13	5
101–500	560	475		393	63	6	1
501–1000	45	42		36	2	0	1
> 1000	20	19		17	0	1	0
Total	14,525	5341		2454	1520	343	1008

The representativeness of each angiosperm family was estimated as the percentage of genera and species conserved within the family against the total number of naturally occurring genera and species described for the family. Likewise, the representativeness of each angiosperm genus was estimated as the percentage of species conserved at MSB within the genus against the total number of naturally occurring species described for the genus. The total number of accepted genera and species described per family and accepted species described per genus were taken from The Plant List (2013) considering the confidence level and review status of names. Doubtful data were excluded from the analysis. Full dataset available in Appendix 1 and Supplement 1. Example: *101* (highlighted in italic) shows that 101 families of plants containing 1–50 genera have > 75% of their naturally occurring genera conserved at the MSB

(44,130). The hotspots with the greatest number of MSB collections are the Mediterranean Basin (13,020), Madagascar and the Indian Ocean Islands (4027), the Cape Floristic Region (3016), Southwest Australia (2935) and the Caucasus (2753). Hotspots with the lowest number of collections are Polynesia-Micronesia (9), the Western Ghats and Sri Lanka (12), the East Melanesian Islands (26), New Caledonia (33) and the Philippines (62).

Vulnerability to extinction

The representativeness of taxa in the MSB that are extinct, rare or vulnerable to extinction and taxa with a lower risk of extinction at global and/or national scales are given in

Table 4 Overview of seed collections conserved at MSB from each bio-geographic region

Bio-geographic region	Collections	Families	Genera	Species	Taxa[a]
Africa	21,109	235	2256	9998	10,677
Antarctic	388	40	104	177	178
Asia—temperate	15,868	178	1653	6649	6938
Asia—tropical	1145	104	352	575	589
Australasia	12,107	183	1281	8582	9068
Europe	17,155	143	1028	5390	6066
Northern America	8662	194	1500	5476	5738
Southern America	3908	172	974	2342	2435
Pacific	63	27	51	56	56
Unknown	2151	134	581	1283	1335

Cultivated collections inherited the geographic origin of the wild plant population from which they were propagated or regenerated. The total number of collections, families, genera, species and taxa originated from each bio-geographic region is presented based on geographic origin data

[a]Includes species, sub species, varieties, etc.

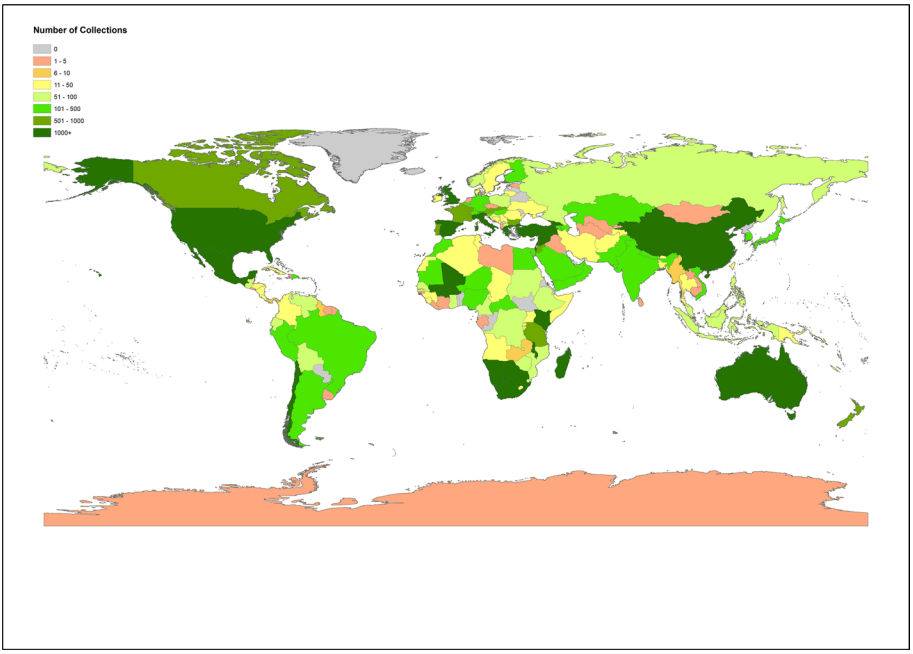

Fig. 1 Geographic origin of MSB collections. Cultivated collections inherited the geographic origin of the wild plant population from which they were propagated or regenerated. Total number of collections are shown according to different size classes. (Color figure online)

Tables 5 and 6. When compared with IUCN (2016) global assessments, MSB conserved at least 5.31% of taxa, representing nearly 1600 collections, which are declared as extinct or vulnerable to extinction (EW, CR, EN and VU). This includes seven of 36 taxa which

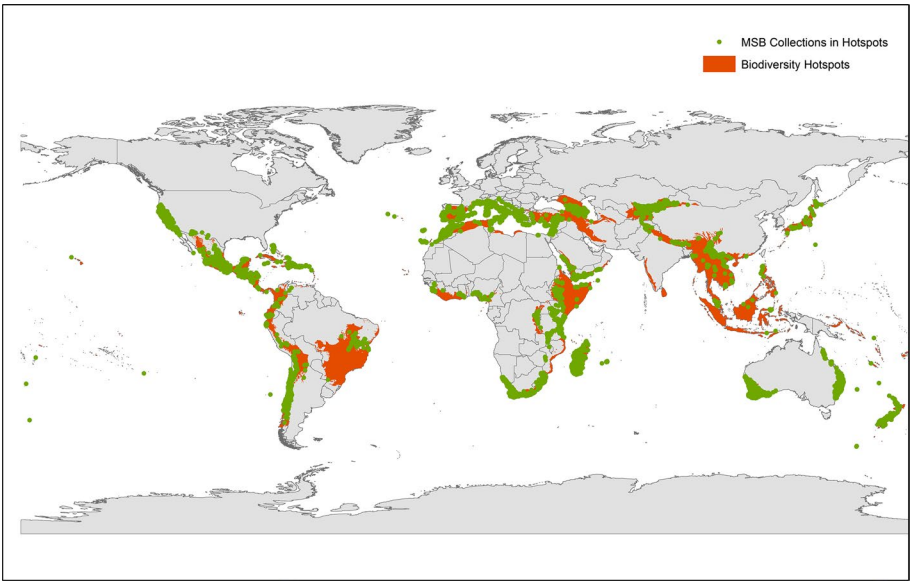

Fig. 2 MSB collections originating from 35 biodiversity hotspots. Cultivated collections inherited the geographic origin of the wild plant population from which they were propagated or regenerated. Collections with missing coordinates but originating from biodiversity hotspots according to locality data are not shown. The complete list of collections is given in Appendix 2. (Color figure online)

are declared as extinct in the wild (Table 7). For those taxa declared as having a lower risk of extinction, MSB collections include 23.19% taxa, represented by over 7400 collections. Nearly 10% of taxa (1487 out of 14,998) that are declared as globally rare in Walter and Gillett (1998) are conserved at the MSB, representing over 2700 collections. At the national scale, the MSB conserved at least 6.93% of taxa (over 3175 collections) that are declared as extinct, rare or vulnerable to extinction nationally in at least one country according to the National Red Lists (2016). In total, at least 10% of taxa (3801 out of 39,669) conserved at MSB, represented by over 6703 collections (> 8% of total holdings), are either extinct, rare or vulnerable to extinction at the global and/or national scale.

Uniqueness and irreplaceability

Twenty percent of MSB taxa (7764 out of 39,669), representing 13% of collections (11,064 out of 82,556), are endemic or near endemic at the country or territory scale.

At least 21% of angiosperm families conserved (70 out of 331) are taxonomically distinct at the species level, and 58% of angiosperm families conserved (191 out of 331) are taxonomically distinct at the genus level, with 11 of the families (Biebersteiniaceae, Cephalotaceae, Drosophyllaceae, Eucommiaceae, Gomortegaceae, Lanariaceae, Plocospermataceae, Quillajaceae, Scheuchzeriaceae, Setchellanthaceae and Strasburgeriaceae) with only one genera and species described in nature (Table 8; Appendix 1). At least 49.6% of angiosperm genera conserved at MSB (2843 out of 5723) are taxonomically distinct with only a few species (up to 10) occurring in nature (Supplement 1).

Table 5 The representativeness of taxa at MSB that are extinct or vulnerable to extinction and taxa with lower risk of extinction at the global scale

Category	Extinct or vulnerable to extinction						Lower risk of extinction						Data deficient	Grand total
	EX	EW	CR	EN	VU	Total	NT	LR/nt	LR/cd	LR/lc	LC	Total	DD	
IUCN (2016)	121	36	2640	3865	5902	12,564	1102	666	221	646	6393	9028	1800	23,392
MSB														
Minimum	0	7	144	210	252	613	115	61	10	37	1781	2004	107	2724
Median	0	7	156	225	279	667	124	63	12	38	1857	2094	125	2885
Maximum	0	7	168	240	306	721	133	65	13	39	1932	2182	142	3045
% Conserved	0.00	19.44	5.91	5.82	4.73	5.31	11.25	9.46	5.20	5.88	29.04	23.19	6.92	12.33

To identify these taxa, SBD taxonomic identifications were cross checked with IUCN (2016). Minimum value refers to the total number of SBD taxa that matched directly with IUCN (2016) but is considered an under-estimate. Maximum value refers to both direct and indirect matches of SBD taxa with IUCN (2016) but is considered an over-estimate. The median is the middle value between the minimum and maximum and is considered as neither under- nor over-estimate of overall representativeness

Table 6 The representativeness of taxa in MSB that are extinct, rare or vulnerable to extinction at the national scale

Category	Extinct, rare or vulnerable to extinction													
	EX	EX/EN	EX/VU	EW	CR	EN	EN/VU	VU	R or RR	I	RE	Relict	Other threatened	Total taxa[a]
National Red Lists (2016)	1284	20	2	32	4741	7850	2	8424	2839	313	222	7	1760	24,969
MSB														
Minimum	7	0	0	7	226	556	0	734	55	0	9	1	118	1681
Median	8	0	0	7	234	575	0	754	57	0	9	1	129	1730
Maximum	8	0	0	7	241	593	0	773	58	0	9	1	139	1778
% Conserved	0.58	0.00	0.00	21.88	4.93	7.32	0.00	8.94	1.99	0.00	4.05	14.29	7.30	6.93

To identify these taxa, SBD taxonomic identifications and their geographic origin were cross checked with National Red List (2016). Minimum value refers to the total number of SBD taxa that matched directly with National Red List (2016) but is considered an under-estimate. Maximum value refers to both direct and indirect matches of SBD taxa with National Red List (2016) but is considered an over-estimate. The median is the middle value between the minimum and maximum and is considered as neither under- nor over-estimate of overall representativeness

[a]IUCN or non-IUCN status category declared for a taxon may differ among countries. Therefore, the total taxa figure is calculated separately by discounting the duplication of taxa by countries across the categories

Table 7 Taxa conserved at MSB that are declared as extinct in the wild (EW) by IUCN (2016)

Family	Taxa
Nymphaeaceae	*Nymphaea thermarum*
Poaceae/Gramineae	*Bromus bromoideus*
Poaceae/Gramineae	*Bromus interruptus*
Primulaceae	*Lysimachia minoricensis*
Solanaceae	*Brugmansia arborea*
Malvaceae	*Trochetiopsis erythroxylon*
Thecaceae	*Franklinia alatamaha*

Table 8 Taxonomic singularity and rarity of MSB seed collections at family level

Taxonomic singularity	Number of angiosperm families conserved at MSB	
Number of genera or species described from The Plant List (2013)	For genera	For species
1	70	11
2	25	19
3	27	8
4	22	6
5	11	5
6	9	5
7	7	7
8	8	6
9	6	0
10	6	3
Total	191	70

The number of genera and species for angiosperm families was obtained from The Plant List (2013) considering the confidence level and review status of names to identify less diversified families (with up to 10 genera or 10 species), then their representation at MSB was measured and expressed as taxonomic singularity to identify taxonomic rarity. Doubtful data were excluded from the analysis (see Appendix 1)

Natural capital value

The majority of MSB collections (49%, 40,430 out of 82,556) have at least one natural capital value and these represent 32% of MSB taxa (12,643 out of 39,669). From a total of 99 CWR genera, 81% (80 genera) are conserved at the MSB. At least 11.3% (9294 out of 82,556) of MSB collections can be considered as CWR sensu lato, represented by 1933 taxa. Of the 4229 CWR sensu stricto taxa compiled from reference sources, 22% (953) are stored at the MSB, representing 8.7% of MSB collections (7160 out of 82,556).

Population value

The majority of taxa (74%, 29,326 out of 39,669) and collections (78%, 64,501 out of 82,556) stored at the MSB are represented by sensu lato 3E taxa. Two-3E taxa are those

that meet two of the three criteria of endemic, endangered or economic, we found that 16% of MSB taxa (6322 out of 39,669), represented by over 12,220 collections (15%), fall under this category. Sensu stricto 3E taxa are those that are endemic, endangered and of economic value i.e. meet all 3Es; 315 MSB taxa represented by 566 collections fall under this category.

Germplasm quality

Figure 3 illustrates the availability of data for MSB collections. A high percentage of collections (> 95%) have taxonomy (Latin plant name) and geographic data at least at country level, followed by geo-coordinates (88%), habitat (87%), altitude (79%) and population data (49–71%). Plant taxonomy of 89% of collections (73,600) are verified either by a field expert, RBG Kew herbarium or another institute in the UK or elsewhere.

Seed quantity data are available for 95% of collections (78,768 collections out of 82,556), with 34% of them represented by more than 5000 potentially viable seeds per collection, whereas 25% are represented by < 501 potentially viable seeds (Fig. 4). About 56,595 collections processed at the MSB are also duplicated in another seed bank in the UK and/or in the country of origin.

For the last round of viability tests carried out for seed collections, germination data are available for 70% of collections (57,658 out of 82,556) and seed viability data are available for 69% collections (57,141 out of 82,556). Of these collections, 47% and 65% respectively showed more than 90% germination and viability and 21% and 11% respectively showed less than 50% germination and viability (Fig. 5). Of the tested collections, 7.7% (4457 out of 57,658) showed 0% germination and 3.2% (1861 out of 57,141) showed 0% viability. As viability percentages were estimates calculated using cut-tested results of non-germinated

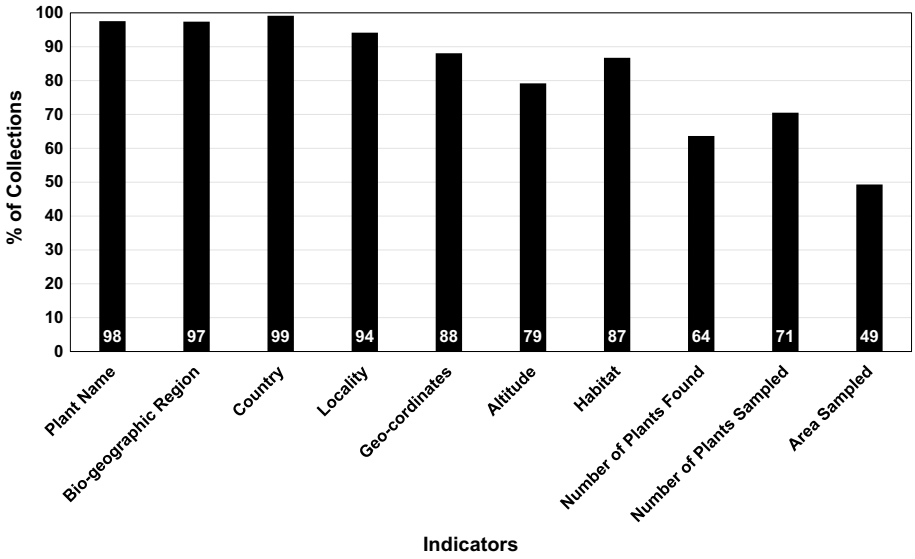

Fig. 3 Availability of key data for MSB collections. These includes taxonomy (plant name), geography (bio-geographic region, country, locality, geo-coordinates and altitude), habitat and population (number of plants found and sampled, and area sampled)

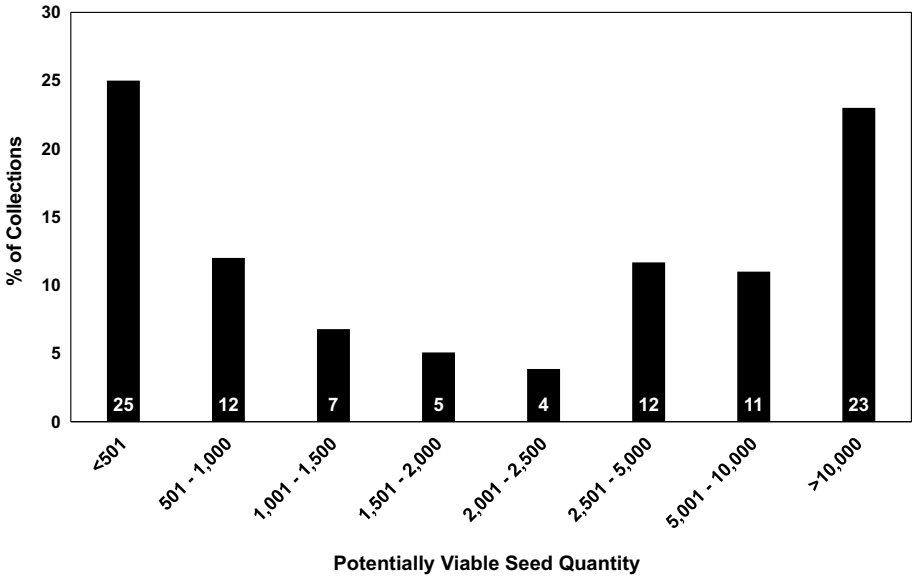

Fig. 4 Potentially viable seed quantity. Percentage of MSB collections in different size classes of estimated number of viable seeds are illustrated taking into account the results of X-rayed or cut-tested samples of dry seeds

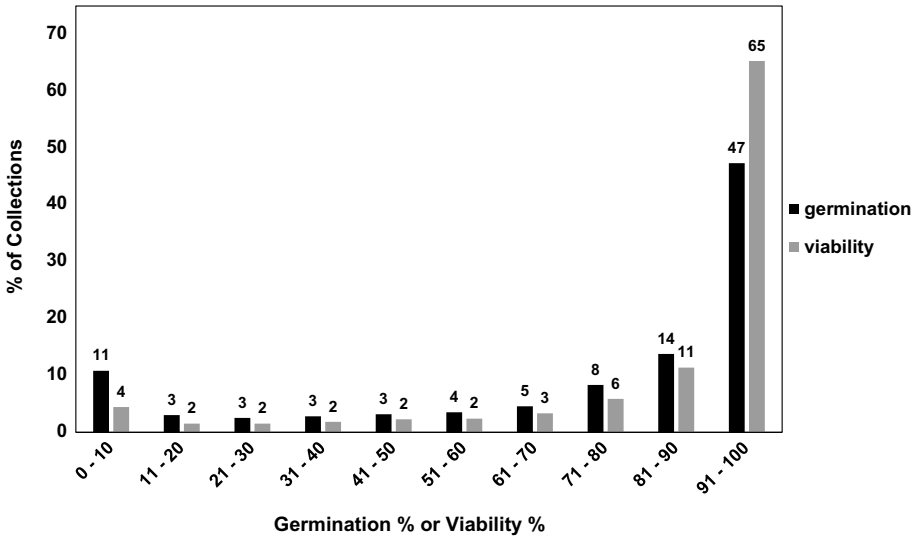

Fig. 5 Seed germination and viability. Percentage of MSB collections in different size classes of germination and viability percentage are illustrated for the last round of germination tests carried out at the MSB. The highest percentages achieved for germination and viability are considered

seeds, collections with 0% germination and viability will be further assessed using experimental controls or Tetrazolium Chloride stain.

Use of collections

During January 2000–March 2017, a total number of 11,182 seed samples representing 6759 collections, 4811 taxa (including subspecies and varieties) and 200 families were supplied to over 410 organizations across 57 countries (includes RBG Kew and UK). At least 8.2% of collections and 12.1% of taxa were distributed globally for a diverse range of uses in research (75%), conservation (13%), education (2%) and display (1%). Details of use for 9% of samples are unknown. Seed samples supplied for research were used across at least 80 pure and applied science disciplines including agriculture, horticulture and archaeology. Those supplied for conservation were used in a range of programs including: habitat restoration and enrichment; species re-introduction or recovery; developing or improving botanic garden, arboretum or nursery collections; regeneration; propagation; breeding; biological control; livelihood programmes; and ex-situ conservation. Samples supplied for education were used during course work and workshops at educational institutes including schools and universities and those supplied for national and international displays were used to raise public awareness of plant conservation. The most distributed taxa were (number of samples distributed in brackets): *Lotus corniculatus* (71); *Brassica oleracea* (60); *Trifolium repens* (48); *Lolium perenne* (36); *Sorghum arundinaceum* (33); *Beta vulgaris* (33); *Daucus carota* (32); *Dactyloctenium aegyptium* (31); *Trifolium pratense* (30); and *Chenopodium album* (30).

Discussion

Plant conservation by seed banking orthodox species at RBG Kew contributes towards RBG Kew's Science Strategy (RBG Kew 2015), Targets 8 and 9 of the GSPC and tackling the challenges of food security, sustainable energy, loss of biodiversity and climate change. This study highlights the following strengths of MSB holdings: a rich biological resource; substantial taxonomic diversity; wide geographic coverage; notable uniqueness and irreplaceability; significant natural capital and population value; and high quality germplasm. Seventy four percent of MSB taxa, representing 78% of collections, fall into one or more of three important categories: (1) endemic; (2) endangered at the national or global scale; and (3) of economic, ecological, social, cultural or scientific value. In addition, MSB holdings represent 81% of CWR genera.

The importance of these collections in the face of threats to global plant diversity cannot be overstressed. An estimated 369,434 species of angiosperms are known to science, and 21% of global plant species are currently threatened with extinction (Nic Lughadha et al. 2016; RBG Kew 2016). By focusing seed conservation efforts on endangered species RBG Kew's MSB is helping to conserve plant diversity that is most at risk of extinction. As many as 44% of all species of vascular plants are confined to 25 hotspots comprising only 1.4% of the land surface of the earth (Myers et al. 2000). Using these regions as collection foci has enabled RBG Kew to maximize its impact with limited resources, and effort continues to be directed to those hotspots currently under-represented in RBG Kew's collections.

Despite the importance of the collections and the targeted approach, limitations to seed conservation remain. Challenges include: the absence of a red list or up to date, accurate information on threatened species for some partner countries; accessible data on the locality, phenology and identification of taxa; infrastructure for processing and storage of seeds; and skills enabling collection, handling and storage of seed (Smith 2007). The level of emphasis on which taxa are to be conserved is governed by the Access and Benefit Sharing Agreements (ABSA), the needs of project partners, the political, legal and administrative requirements of the regional, national or local government in question and the level of implementation of the CBD within a particular country or region (Cheyne 2003). Due to the current global financial crisis, the availability of monetary funds is another bottle neck for initiating conservation programmes. Lack of political will and funding are the biggest constraints preventing the achievement of Target 8 of GSPC (Smith 2007). An estimated £50 million was spent to secure 10% of world's flora during the first phase of ex situ conservation in RBG Kew's international programme (Smith 2007), representing about £2100 per plant species to guarantee its long-term ex situ seed conservation (Li and Pritchard 2009).

Most importantly, only orthodox species whose seeds can survive considerable desiccation and freezing during ex situ conservation are bankable in traditional seed banks. Species adapted to hot, dry environments may have evolved longer lifespans in the dry state and are suitable to conserve in seed banks (Li and Pritchard 2009). The world's drylands are home to an immense variety of plant life, with a high proportion of species producing orthodox seeds, and supporting approximately one-fifth of the world's population (far more than the tropical rain forests), as well as 50% of the world's livestock, and provide forage for both domestic animals and wildlife (van Slageren 2003). However, drylands are among the most threatened environments on Earth, with large areas being lost due to desertification each year (van Slageren 2003). As a result, the MSB initially focused its seed conservation efforts in drylands.

Representation of plant taxa in gene banks is also subjected to species, species-area, hotspot and infrastructure biases that result in over- or under- representing certain taxa (Hijmans et al. 2000). Under representation of threatened taxa may be linked to geographic rarity or recalcitrant seeds. Families with a high incidence of recalcitrant species which are less likely to be conserved in conventional seed banks are Fagaceae, Lauraceae, Sapotaceae, Moraceae, Clusiaceae, Sapindaceae (including Aceraceae), Arecaceae (= Palmae), Myrtaceae, Annonaceae, Rurtaceae, Anarcardiaceae, Dipterocarpaceae, Meliaceae and Rhizophoraceae (Dickie and Pritchard 2002). Some wild species have been found to produce seeds that are extremely short lived in traditional seed bank storage and cryopreservation may be the only resource to ensure the effective ex situ seed conservation of such species (Li and Pritchard 2009; Hay and Probert 2013). Li and Pritchard (2009) highlighted the need to increase our effort at developing ex situ conservation approaches for plants, particularly those from biodiversity hotspots with recalcitrant seeds.

Placing a monetary value on conservation collections is difficult. A recent estimate suggests that the potential value of benefits from CWR traits of the MSB's 29 current priority crops alone amounts to approximately $120 billion (PwC 2013). Economic assessment is extraordinarily difficult beyond the main crops, particularly for species not yet fully characterised for traits of societal value (Li and Pritchard 2009). If CWR of 29 crops alone are worth $120 billion, the entire MSB holdings of 39,669 taxa can be considered as a very successful output of a global ex situ conservation program, and an extremely valuable biological resource.

MSB collections and associated data are already being used by partners to restore populations of wild plants (South Africa), to manage damaged ecosystems (Madagascar), to rehabilitate degraded lands (Burkina Faso and Australia), to develop opportunities for the sustainable use of plants by local communities (Kenya) and for scientific research (unpublished, RBG Kew). In addition, small quantities of MSB collections (usually 50 seeds) and associated data are also being supplied for use in conservation, research, education and display across the world. While the sample size is too small to support large-scale restoration projects, or research programmes requiring a large number of seeds, the importance of the collections for these areas cannot be overstated. The depth and breadth of information, knowledge and research outputs underpinning plant conservation being generated by MSB collections and associated data across the world is unparalleled. It is shared with policy makers, disseminated to the wider scientific community and general public through mainstream scientific literature, articles, conferences, social media and the RBG Kew website and those of our partners. Although compilation of a full list of research publications is beyond the scope of this paper, recent research studies that used MSB collections or their associated data include Colville et al. (2015), Díaz et al. (2016), Fernández-Marín et al. (2017), Mattana et al. (2017), Rodríguez-Arévalo et al. (2017), Seal et al. (2017), Ulian et al. (2017) and Wyse and Dickie (2017).

Our analysis was focused only on taxa conserved at the MSB and not on those stored only in seed banks of partner countries within the MSBP network. At present, it is difficult to assess the conservation value of taxa conserved within the whole network due to various constraints including lack of access to databases in partner countries. This is a problem that is being overcome by the MSBP Data Warehouse (http://brahmsonline.kew.org/msbp), an online BRAHMS database holding collection data from across the MSBP, and crucially, including data on collections not duplicated to the MSB. Further data analysis is essential across the MSBP network to underpin future collection activities to maximize the usefulness of collections while concentrating on gaps in threatened taxa, geographic representation and taxonomic diversity highlighted in our study, while identifying their suitability for conserving in conventional seed banks including cryopreservation.

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Appendix 1

See Table 9.

Table 9 The representativeness of angiosperm families at the MSB

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Acanthaceae	316	51	244	21	213	4016	5
Achariaceae	13	7	27	26	7	98	7
Acoraceae	1	1	1	100	1	4	25
Actinidiaceae	14	2	3	67	10	193	5
Adoxaceae	159	3	5	60	51	230	22
Aizoaceae	732	93	146	64	447	2330	19
Alismataceae	79	10	17	59	28	130	22
Alseuosmiaceae	2	2	2	100	2	7	29
Alstroemeriaceae	79	3	5	60	32	271	12
Altingiaceae	5	2	2	100	4	16	25
Amaranthaceae	1410	99	179	55	572	2211	26
Amaryllidaceae	611	27	81	33	276	2376	12
Anacampserotaceae	5	1	3	33	3	137	2
Anacardiaceae	322	44	75	59	139	750	19
Anarthriaceae	2	2	3	67	2	11	18
Anisophylleaceae	2	1	4	25	2	39	5
Annonaceae	81	22	128	17	49	2135	2
Aphanopetalaceae	1	1	1	100	1	4	25
Aphloiaceae	8	1	1	100	1	2	50
Apiaceae	1792	206	416	50	744	3492	21
Apocynaceae	585	117	318	37	321	3400	9
Aponogetonaceae	1	1	2	50	1	58	2
Aquifoliaceae	114	1	4	25	40	519	8
Araceae	74	14	117	12	41	3459	1
Araliaceae	151	23	44	52	112	1609	7

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Arecaceae	115	36	185	19	76	2625	3
Argophyllaceae	5	2	2	100	4	[a]	[a]
Aristolochiaceae	39	4	8	50	24	643	4
Asparagaceae	1165	88	126	70	549	3087	18
Asteliaceae	17	4	4	100	11	39	28
Asteropeiaceae	1	1	1	100	1	8	12
Atherospermataceae	9	4	1	[a]	6	1	[a]
Balanophoraceae	1	1	16	6	1	52	2
Balsaminaceae	46	1	2	50	34	506	7
Basellaceae	3	2	4	50	3	19	16
Bataceae	3	1	1	100	1	2	50
Begoniaceae	45	1	2	50	34	1618	2
Berberidaceae	92	10	20	50	60	780	8
Berberidopsidaceae	2	1	1	100	1	2	50
Betulaceae	259	5	6	83	67	282	24
Biebersteiniaceae	2	1	1	100	1	1	100
Bignoniaceae	158	37	85	44	78	865	9
Bixaceae	14	3	4	75	8	26	31
Blandfordiaceae	6	1	1	100	4	4	100
Boraginaceae	799	78	155	50	456	2983	15
Boryaceae	7	2	2	100	6	12	50
Brassicaceae	1577	162	372	44	653	4507	14
Bromeliaceae	200	23	52	44	112	3404	3
Brunelliaceae	3	1	1	100	2	65	3

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Bruniaceae	20	6	11	55	14	92	15
Burmanniaceae	3	3	17	18	3	165	2
Burseraceae	95	6	19	32	56	682	8
Butomaceae	8	1	1	100	1	2	50
Buxaceae	21	2	5	40	10	132	8
Byblidaceae	3	1	1	100	3	7	43
Cactaceae	330	45	176	26	176	2715	6
Calceolariaceae	83	2	4	50	47	314	15
Calophyllaceae	4	2	7	29	4	57	7
Calycanthaceae	4	2	3	67	4	12	33
Calyceraceae	3	2	6	33	3	56	5
Campanulaceae	697	37	87	43	354	2575	14
Canellaceae	1	1	5	20	1	21	5
Cannabaceae	78	7	8	88	31	115	27
Cannaceae	6	1	1	100	1	12	8
Capparaceae	178	11	35	31	75	492	15
Caprifoliaceae	453	26	55	47	205	954	21
Caricaceae	3	3	6	50	3	51	6
Caryophyllaceae	1887	55	93	59	590	2865	21
Casuarinaceae	98	3	4	75	70	94	74
Celastraceae	283	39	86	45	138	1217	11
Centrolepidaceae	19	3	3	100	12	39	31
Cephalotaceae	1	1	1	100	1	1	100
Ceratophyllaceae	1	1	1	100	1	10	10

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Cercidiphyllaceae	3	1	1	100	2	2	100
Chenopodiaceae	1	1	[a]	[a]	1	[a]	[a]
Chloranthaceae	1	1	5	20	1	74	1
Chrysobalanaceae	23	7	18	39	13	564	2
Cistaceae	300	7	9	78	78	258	30
Cleomaceae	136	6	11	55	56	269	21
Clethraceae	8	1	2	50	4	89	4
Clusiaceae	19	4	31	13	14	1142	1
Colchicaceae	114	10	18	56	61	300	20
Columelliaceae	1	1	2	50	1	10	10
Combretaceae	298	13	17	76	123	519	24
Commelinaceae	91	14	41	34	55	766	7
Compositae	8486	774	1923	40	4386	36,722	12
Connaraceae	9	4	19	21	7	284	2
Convolvulaceae	496	25	67	37	221	1409	16
Coriariaceae	12	1	1	100	6	18	33
Cornaceae	83	3	12	25	33	169	20
Corsiaceae	3	1	3	33	1	27	4
Corynocarpaceae	1	1	1	100	1	5	20
Costaceae	4	2	7	29	3	145	2
Crassulaceae	535	22	50	44	295	1670	18
Crossosomataceae	1	1	3	33	1	12	8
Cucurbitaceae	437	46	134	34	143	1007	14
Cunoniaceae	28	10	20	50	18	255	7

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Cynomoriaceae	4	1	1	100	1	2	50
Cyperaceae	2266	58	110	53	971	6265	15
Cyrillaceae	1	1	3	33	1	11	9
Daphniphyllaceae	4	1	1	100	2	35	6
Dasypogonaceae	4	3	4	75	4	16	25
Datiscaceae	2	1	1	100	1	2	50
Diapensiaceae	1	1	7	14	1	15	7
Dichapetalaceae	6	2	3	67	5	208	2
Didiereaceae	7	4	7	57	6	22	27
Dilleniaceae	64	3	11	27	52	232	22
Dioscoreaceae	103	2	6	33	50	665	8
Dipentodontaceae	1	1	2	50	1	23	4
Dipterocarpaceae	5	1	16	6	3	155	2
Doryanthaceae	3	1	1	100	2	2	100
Droseraceae	78	2	3	67	36	210	17
Drosophyllaceae	2	1	1	100	1	1	100
Ebenaceae	184	2	3	67	77	777	10
Ecdeiocoleaceae	1	1	2	50	1	3	33
Elaeagnaceae	23	3	4	75	11	111	10
Elaeocarpaceae	37	4	9	44	26	677	4
Elatinaceae	24	2	2	100	17	57	30
Ericaceae	979	67	130	52	574	3682	16
Eriocaulaceae	58	2	11	18	35	1254	3
Erythroxylaceae	35	1	4	25	17	269	6

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Escalloniaceae	20	3	10	30	11	115	10
Eucommiaceae	1	1	1	100	1	1	100
Euphorbiaceae	605	55	227	24	351	6834	5
Eupomatiaceae	1	1	1	100	1	3	33
Eupteleaceae	3	1	1	100	2	2	100
Fagaceae	12	1	9	11	4	1151	0
Flagellariaceae	9	1	1	100	2	4	50
Fouquieriaceae	15	1	2	50	6	13	46
Frankeniaceae	27	1	1	100	13	80	16
Garryaceae	11	2	2	100	6	33	18
Gelsemiaceae	1	1	2	50	1	8	12
Gentianaceae	467	33	95	35	176	1874	9
Geraniaceae	250	5	10	50	130	882	15
Gesneriaceae	103	27	161	17	61	3191	2
Gisekiaceae	11	1	1	100	3	7	43
Gomortegaceae	1	1	1	100	1	1	100
Goodeniaceae	172	9	13	69	120	333	36
Griseliniaceae	2	1	1	100	2	[a]	[a]
Grossulariaceae	52	1	2	50	36	229	16
Gunneraceae	16	1	1	100	6	72	8
Gyrostemonaceae	16	3	3	100	13	17	76
Haemodoraceae	74	10	14	71	53	108	49
Haloragaceae	58	8	9	89	45	99	45
Hamamelidaceae	16	8	23	35	11	108	10

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Heliconiaceae	1	1	1	100	1	208	0
Hernandiaceae	15	4	4	100	6	57	11
Huaceae	1	1	1	100	1	3	33
Hydatellaceae	3	1	1	100	3	12	25
Hydrangeaceae	64	9	17	53	42	283	15
Hydrocharitaceae	15	4	16	25	7	147	5
Hydroleaceae	4	1	1	100	4	4	100
Hydrostachyaceae	1	1	1	100	1	19	5
Hypericaceae	267	5	12	42	111	660	17
Hypoxidaceae	41	4	10	40	21	169	12
Icacinaceae	8	3	34	9	5	197	3
Iridaceae	929	46	80	57	508	2456	21
Irvingiaceae	1	1	3	33	1	11	9
Iteaceae	4	3	3	100	4	20	20
Ixioliriaceae	5	1	1	100	1	5	20
Joinvilleaceae	1	1	1	100	1	4	25
Juglandaceae	16	5	12	42	10	98	10
Juncaceae	431	6	8	75	175	581	30
Juncaginaceae	44	3	4	75	18	36	50
Kewaceae	2	1	1	100	1	8	12
Kirkiaceae	5	1	1	100	3	3	100
Koeberliniaceae	1	1	1	100	1	3	33
Krameriaceae	7	1	1	100	5	25	20
Lamiaceae	2453	127	246	52	1044	8608	12

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Lanariaceae	1	1	1	100	1	1	100
Lardizabalaceae	11	5	9	56	7	41	17
Lauraceae	47	9	69	13	29	3029	1
Lecythidaceae	8	2	24	8	4	345	1
Leguminosae	13,092	392	732	54	3917	21,664	18
Lentibulariaceae	46	2	4	50	30	321	9
Liliaceae	246	13	19	68	142	807	18
Limeaceae	13	2	1	[a]	10	13	77
Limnanthaceae	2	1	2	50	1	18	6
Linaceae	155	5	16	31	57	241	24
Linderniaceae	26	5	16	31	25	167	15
Loasaceae	78	10	22	45	50	338	15
Loganiaceae	81	9	18	50	47	393	12
Lophiocarpaceae	7	2	1	[a]	5	3	[a]
Loranthaceae	9	7	84	8	7	1170	1
Lythraceae	140	16	32	50	54	625	9
Magnoliaceae	28	2	6	33	23	275	8
Malpighiaceae	70	25	81	31	47	1327	4
Malvaceae	1899	132	216	61	821	3503	23
Marantaceae	6	2	28	7	3	571	1
Marcgraviaceae	2	1	8	12	2	140	1
Martyniaceae	12	2	5	40	5	24	21
Melanthiaceae	40	9	17	53	23	207	11
Melastomataceae	148	38	169	22	96	4131	2

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Meliaceae	84	20	51	39	47	693	7
Melianthaceae	18	4	4	100	9	22	41
Menispermaceae	49	14	68	21	28	460	6
Menyanthaceae	23	6	7	86	18	55	33
Molluginaceae	82	8	14	57	35	125	28
Monimiaceae	14	3	23	13	8	155	5
Montiaceae	133	8	20	40	71	283	25
Montiniaceae	5	3	2	[a]	3	6	50
Moraceae	154	12	40	30	88	1346	7
Moringaceae	9	1	1	100	6	13	46
Muntingiaceae	2	1	3	33	1	3	33
Musaceae	68	2	2	100	8	98	8
Myricaceae	28	2	5	40	12	57	21
Myristicaceae	1	1	17	6	1	154	1
Myrothamnaceae	4	1	1	100	1	2	50
Myrtaceae	2110	76	145	52	1501	6133	24
Nartheciaceae	15	3	5	60	6	38	16
Nelumbonaceae	1	1	1	100	1	2	50
Nepenthaceae	6	1	1	100	5	8	62
Neuradaceae	6	3	3	100	5	7	71
Nitrariaceae	39	3	2	[a]	9	8	[a]
Nothofagaceae	24	1	1	100	7	40	18
Nyctaginaceae	79	10	34	29	48	482	10
Nymphaeaceae	13	2	8	25	9	83	11

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Ochnaceae	21	5	38	13	16	515	3
Olacaceae	44	5	32	16	12	154	8
Oleaceae	277	14	25	56	82	767	11
Onagraceae	484	16	43	37	190	985	19
Opiliaceae	12	4	10	40	6	39	15
Orchidaceae	2028	256	897	29	1294	28,347	5
Orobanchaceae	640	43	77	56	313	1727	18
Oxalidaceae	25	2	8	25	14	643	2
Paeoniaceae	36	1	1	100	17	52	33
Pandanaceae	12	3	4	75	12	1069	1
Papaveraceae	355	24	41	59	131	1028	13
Paracryphiaceae	2	1	1	100	2	1	[a]
Passifloraceae	94	8	34	24	55	880	6
Paulowniaceae	8	1	3	33	5	20	25
Pedaliaceae	88	12	14	86	37	72	51
Penaeaceae	13	5	13	38	7	48	15
Pennantiaceae	1	1	[a]	[a]	1	5	20
Pentaphylacaceae	15	3	8	38	15	99	15
Penthoraceae	2	1	1	100	2	2	100
Peraceae	12	3	5	60	10	125	8
Philesiaceae	2	1	2	50	1	2	50
Philydraceae	3	2	3	67	2	6	33
Phrymaceae	58	7	11	64	33	215	15
Phyllanthaceae	198	19	58	33	87	2204	4

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Phyllonomaceae	1	1	1	100	1	5	20
Physenaceae	5	1	1	100	1	2	50
Phytolaccaceae	31	6	13	46	9	64	14
Picramniaceae	3	2	4	50	2	52	4
Picrodendraceae	24	9	25	36	18	97	19
Piperaceae	45	3	13	23	30	2717	1
Pittosporaceae	88	10	10	100	57	192	30
Plantaginaceae	1071	50	91	55	472	1662	28
Platanaceae	13	1	1	100	5	10	50
Plocospermataceae	4	1	1	100	1	1	100
Plumbaginaceae	163	10	24	42	91	698	13
Poaceae	6184	349	757	46	1980	11,881	17
Polemoniaceae	76	13	30	43	58	562	10
Polygalaceae	96	8	27	30	59	1200	5
Polygonaceae	602	29	59	49	275	1584	17
Pontederiaceae	7	4	6	67	7	34	21
Portulacaceae	57	3	7	43	21	186	11
Potamogetonaceae	53	5	6	83	25	195	13
Primulaceae	450	31	51	61	217	1510	14
Proteaceae	1114	35	68	51	730	1360	54
Putranjivaceae	7	1	3	33	6	217	3
Quillajaceae	1	1	1	100	1	1	100
Ranunculaceae	1022	33	66	50	446	2767	16
Resedaceae	93	5	7	71	26	61	43

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Restionaceae	184	29	47	62	118	485	24
Rhamnaceae	595	35	53	66	265	915	29
Rhipogonaceae	2	1	1	100	2	6	33
Rhizophoraceae	1	1	18	6	1	142	1
Roridulaceae	3	1	1	100	2	2	100
Rosaceae	2283	67	104	64	893	5317	17
Rousseaceae	4	3	3	100	3	14	21
Rubiaceae	1410	193	608	32	736	14,268	5
Ruppiaceae	4	1	1	100	3	8	38
Rutaceae	553	58	158	37	331	1848	18
Sabiaceae	6	1	4	25	3	127	2
Salicaceae	88	18	39	46	49	995	5
Salvadoraceae	12	3	3	100	4	8	50
Santalaceae	76	14	42	33	48	762	6
Sapindaceae	438	45	139	32	194	1857	10
Sapotaceae	79	13	59	22	46	1334	3
Sarcobataceae	1	1	1	100	1	2	50
Sarcolaenaceae	22	6	10	60	12	69	17
Sarraceniaceae	12	1	3	33	5	38	13
Saururaceae	5	3	4	75	3	7	43
Saxifragaceae	241	17	50	34	122	932	13
Scheuchzeriaceae	2	1	1	100	1	1	100
Schisandraceae	11	3	3	100	6	78	8
Schoepfiaceae	1	1	3	33	1	52	2

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Scrophulariaceae	605	42	139	30	364	1680	22
Setchellanthaceae	1	1	1	100	1	1	100
Simaroubaceae	19	5	21	24	11	127	9
Simmondsiaceae	1	1	1	100	1	2	50
Smilacaceae	45	1	2	50	24	270	9
Solanaceae	1008	47	114	41	388	2767	14
Sphaerosepalaceae	7	1	2	50	3	20	15
Sphenocleaceae	6	1	1	100	1	2	50
Stachyuraceae	4	1	1	100	4	7	57
Staphyleaceae	13	3	4	75	9	31	29
Stegnospermataceae	2	1	1	100	1	5	20
Stilbaceae	14	5	12	42	11	38	29
Strasburgeriaceae	1	1	1	100	1	1	100
Strelitziaceae	5	2	3	67	4	7	57
Stylidiaceae	134	5	5	100	115	30	[a]
Styracaceae	35	5	12	42	14	145	10
Surianaceae	11	3	4	75	4	6	67
Symplocaceae	16	1	3	33	8	219	4
Talinaceae	25	2	2	100	10	23	43
Tamaricaceae	24	3	3	100	13	83	16
Tecophilaeaceae	33	6	9	67	16	26	62
Tetracarpaeaceae	1	1	1	100	1	[a]	[a]
Tetrachondraceae	1	1	2	50	1	3	33
Theaceae	17	6	25	24	12	783	2

Table 9 (continued)

Family	Collections conserved at MSB	Genera conserved at MSB	Genera in The Plant List (2013)	% Genera conserved at MSB	Species conserved at MSB	Species in The Plant List (2013)	% Species conserved at MSB
Thymelaeaceae	137	16	55	29	95	1008	9
Tofieldiaceae	19	3	4	75	8	33	24
Trochodendraceae	4	2	2	100	2	2	100
Typhaceae	65	2	2	100	15	69	22
Ulmaceae	65	5	8	62	20	67	30
Urticaceae	126	22	55	40	66	1541	4
Vahliaceae	3	1	1	100	3	6	50
Velloziaceae	12	3	6	50	7	286	2
Verbenaceae	217	19	34	56	102	1068	10
Violaceae	167	5	25	20	89	885	10
Vitaceae	183	10	16	62	99	1045	9
Vivianiaceae	7	2	4	50	4	16	25
Winteraceae	30	3	6	50	11	38	29
Xanthorrhoeaceae	568	25	34	74	330	1404	24
Xeronemataceae	2	1	1	100	1	3	33
Xyridaceae	42	1	5	20	28	400	7
Zingiberaceae	39	8	52	15	25	1611	2
Zygophyllaceae	207	17	29	59	104	230	45

The representativeness of each angiosperm family was estimated as the percentage of genera and species conserved within the family against the total number of naturally occurring genera and species described for the family. The total number of accepted genera and species described per family were taken from The Plant List (2013) considering the confidence level and review status of names

[a]Doubtful data were excluded from the analysis

Appendix 2

See Table 10.

Table 10 Number of MSB collections originated from 35 biodiversity hotspots

Biodiversity hotspot	Main countries	Number of MSB collections originated from hotspots
Mediterranean Basin	Portugal, Spain, France, Italy, Greece, Croatia, Montenegro, Turkey, Israel, Morocco, Tunisia, Algeria, Libya, Cyprus, Albania, Lebanon, Malta, Syria	13,020
Madagascar and the Indian Ocean Islands	Madagascar, Reunion, Mauritius, Comoros, Seychelles, Mayotte	4027
Cape Floristic Region	South Africa: South, East and Western Capes	3016
Southwest Australia	Australia: Western	2935
Caucasus	Armenia, Azerbaijan, Georgia, Iran, Russia, Turkey	2753
Forests of East Australia	Australia: New South Wales, Queensland	1745
Eastern Afromontane	Burundi, DRC, Eritrea, Ethiopia, Kenya, Malawi, Mozambique, Rwanda, Saudi Arabia, Somalia, Sudan, South Sudan, Tanzania, Uganda, Yemen, Zambia, Zimbabwe	1571
Mountains of Central Asia	Kyrgyzstan, Kazakhstan, Tajikistan, Uzbekistan, China: Xinjiang	1457
Chilean Winter Rainfall and Valdivian Forests	Chile, Argentina	1416
Mesoamerica	Mexico, Guatemala, Belize, Costa Rica, El Salvador, Honduras, Nicaragua, Panama	1345
Maputaland-Pondoland-Albany	South Africa, Swaziland, Mozambique	1242
Succulent Karoo	South Africa, Namibia	1103
Caribbean Islands	All Caribbean Islands	1100
Indo-Burma	Myanmar, Bangladesh, China: Yunnan, Laos, Cambodia, Vietnam, Thailand, Malaysia	1035
California Floristic Province	Mexico: Baja California, USA: Oregon, Nevada, California	1011
Madrean Pine-Oak Woodlands	Mexico, USA: Texas, New Mexico, Arizona	847
Horn of Africa	Djibouti, Eritrea, Ethiopia, Kenya, Oman, Saudi Arabia, Somalia, Sudan, South Sudan, Tanzania, Yemen	671
New Zealand	New Zealand	592
Tropical Andes	Argentina, Bolivia, Chile, Colombia, Ecuador, Peru, Venezuela	540
Coastal Forests of Eastern Africa	Kenya, Tanzania, Mozambique, Somalia	485
Irano-Anatolian	Armenia, Azerbaijan, Georgia, Iraq, Iran, Turkmenistan	465
Himalaya	Nepal, Pakistan, Bhutan, India, China: Xizang	392
Mountains of Southwest China	China: Sichuan, Xizang, Yunnan	366
Cerrado	Brazil, Bolivia, Paraguay	188

Table 10 (continued)

Biodiversity hotspot	Main countries	Number of MSB collections originated from hotspots
Guinean Forests of West Africa	Guinea, Sierra Leone, Cameroon, Benin, Ivory Coast, Equatorial Guinea, Ghana, Liberia, Nigeria, Sao Tome & Principe, Togo	183
Japan	Japan	157
Atlantic Forest	Brazil, Argentina, Paraguay	98
Tumbes-Choco-Magdalena	Colombia, Ecuador, Panama, Peru	85
Sundaland	Malaysia, Indonesia, Brunei, Singapore	76
Wallacea	Indonesia, Timor-Leste	67
Philippines	Philippines	62
New Caledonia	New Caledonia	33
East Melanesian Islands	Papua New Guinea, Vanuatu, Solomon Islands	26
Western Ghats and Sri Lanka	India, Sri Lanka	12
Polynesia-Micronesia	Cook Islands, Fiji, French Polynesia, Guam, Kiribati, Marshall Islands, Micronesia, Nauru, Niue, Northern Mariana Islands, Palau, Pitcairn Islands, Samao, Tokelau, Tonga, Tuvalu, Wallis and Futana	9
Total		44,130

Cultivated collections inherited the geographic origin of the wild plant population from which they were propagated or regenerated

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