

Collecting Genetic Variation on a Small Island¹

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Abstract

Genetic variation is the most powerful factor in ensuring the long term success of trees and forests in times of change. In order to protect against loss of genetic variation from threats, including pests and diseases and climate change, the Royal Botanic Gardens, Kew, is developing a national tree seed collection for the United Kingdom.

This paper describes the authors methodologies and experiences of developing the national collection: firstly, forming species target lists using plant health risk and conservation assessments; then, developing national sampling strategies based on the distribution of target species in biogeographic zones; and, finally, how collecting strategies were designed to capture and preserve genetic variation within populations.

Additionally, we discuss social and ecological factors taken into account when developing and implementing the sampling program. These include phylogeography; the history of woodland management, ownership and fragmentation; and introgression risks from plantation and garden escapees. Finally citizen participation is discussed.

Introduction

The Royal Botanic Gardens, Kew launched the United Kingdom National Tree Seed Project (UKNTSP) in 2013 as an *ex situ* seed conservation initiative. The UKNTSP's stated aim is: 'To provide a national repository of plant material and associated knowledge, for the purposes of long term conservation, and to make these resources available to users, in order to better understand and manage tree and shrub species in the United Kingdom landscape.'

In this paper we set out our key considerations and experiences in targeting, sampling and collecting seed for the project.

Developing Target Lists

Native Species

The United Kingdom woody flora consists of 139 native species, 11 archeophytes, (naturalized non-native species introduced prior to 1500), and 57 neophytes (naturalized and introduced after 1500). Native species are considered to be those that recolonized the British Isles between the last ice age and the separation of the British Isles from mainland Europe, around 8 to 10,000 years ago; or those that speciated in the United Kingdom after this period. Re-colonization occurred from glacial refugial populations from Spain, Italy and the Balkans, in the main (Newton et al. 1999).

Prioritization

To prioritize native species for seed banking in the project timeframe, a target list was developed, using a scoring and ranking system. Impacts from invasive species, including pests and diseases, is one of the key drivers for change affecting forest genetic resources (FAO 2014). Plant health, therefore, was a primary factor in our prioritization. We used the United Kingdom Plant Health Risk Register (DEFRA 2014) to prioritize species most at risk. This tool adopts a risk assessment approach to potential plant health threats to hosts, by assessing impact and likelihood and setting out mitigation approaches. 'Unmitigated risk'

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assessments for plant health risks were totalled for each host, which were then ranked accordingly. Wider threats to species were scored using Great Britain Red List status (Cheffings et al. 2005). Species were again ranked, those with a higher 'Extinction Risk' threat status received greater priority. Finally, as a measure of potential ecological impact at a landscape-scale, 'prevalence in the landscape' was also used to rank targets, using the PLANTATT database (Hill et al. 2004); species that had wider distributions received greater priority. For each ranking 'Plant health risk' was used as a secondary factor in ranking if species received the same score, for example both *Pinus sylvestris* and *Acer campestre* have a Red List status of 'Least Concern', but *P. sylvestris* received a higher rank in this category because there is a greater 'Plant health risk' for this species. See table 1 for a list of target taxa and prioritisation data.

Table 1—Prioritization list of target taxa

Taxon name	Unmitigated risk total score	Risk rank	Red list GB ^a	Red list rank	No. 10 km ² in GB	Distribution rank	Native status ^b	Native rank	Total rank
<i>Acer campestre</i>	745	25	LC	68	1389	37	N	62	192
<i>Alnus glutinosa</i>	420	46	LC	85	2478	8	N	80	219
<i>Betula nana</i>	591	45	LC	84	125	77	N	79	285
<i>Betula pendula</i>	591	44	LC	83	2293	21	N	78	226
<i>Betula pubescens</i>	651	30	LC	72	2399	15	N	66	183
<i>Buxus sempervirens</i>	54	111	DD	13	2	112	N	108	344
<i>Carpinus betulus</i>	281	51	LC	90	1488	35	N	84	260
<i>Cornus sanguinea</i>	182	56	LC	95	1179	41	N	89	281
<i>Crataegus laevigata</i>	153	58	LC	97	597	59	N	91	305
<i>Crataegus monogyna</i>	233	54	LC	93	2496	7	N	87	241
<i>Erica vagans</i>	16	115	LC	116	6	99	N	110	440
<i>Fagus sylvatica</i>	280	52	LC	91	2397	16	N	85	244
<i>Fraxinus excelsior</i>	407	47	LC	86	2459	11	N	81	225
<i>Ilex aquifolium</i>	183	55	LC	94	2353	17	N	88	254
<i>Juniperus communis</i>	147	59	LC	98	1020	46	N	92	295
<i>Juniperus communis</i> subsp. <i>hemisphaerica</i>	87	65	CR	1	2	107	N	98	271
<i>Lonicera periclymenum</i>	0	120	LC	119	2622	1	N	115	355
<i>Malus sylvestris</i>	1175	19	LC	62	2023	26	N	56	163
<i>Pinus sylvestris</i>	2114	2	LC	46	65	83	N	39	170
<i>Prunus avium</i>	2225	1	LC	45	2136	24	N	38	108
<i>Prunus padus</i>	1901	4	LC	48	1089	44	N	41	137
<i>Prunus spinosa</i>	1901	3	LC	47	2308	20	N	40	110
<i>Pyrus cordata</i>	642	31	VU	32	9	94	NA	128	285
<i>Rubus idaeus</i>	914	22	LC	65	2425	13	N	59	159
<i>Sambucus nigra</i>	4	118	LC	118	2457	12	N	113	361
<i>Sorbus admonitor</i>	72	87	EN	24	1	122	NE	22	255

<i>Sorbus anglica</i>	72	99	NT	41	12	93	NE	34	267
<i>Sorbus aria</i>	72	106	LC	109	341	67	N	103	385
<i>Sorbus arranensis</i>	72	96	VU	36	1	130	NE	31	293
<i>Sorbus arvonicola</i>	72	67	CR	3	1	113	NE	3	186
<i>Sorbus aucuparia</i>	72	103	LC	106	2472	9	N	100	318
<i>Sorbus bristoliensis</i>	72	88	EN	25	1	123	NE	23	259
<i>Sorbus cambrensis</i>	72	84	EN	21	2	109	NE	19	233
<i>Sorbus cheddarensis</i>	72	68	CR	4	1	114	NE	4	190
<i>Sorbus cuneifolia</i>	72	89	EN	26	1	124	NE	24	263
<i>Sorbus devoniensis</i>	72	101	LC	104	32	89	NE	36	330
<i>Sorbus domestica</i>	72	76	CR	12	4	101	N	99	288
<i>Sorbus eminens</i>	72	77	EN	14	9	95	NE	12	198
<i>Sorbus eminentiformis</i>	72	85	EN	22	2	110	NE	20	237
<i>Sorbus eminentoides</i>	72	69	CR	5	1	115	NE	5	194
<i>Sorbus evansii</i>	72	90	EN	27	1	125	NE	25	267
<i>Sorbus greenii</i>	72	91	EN	28	1	126	NE	26	271
<i>Sorbus herefordensis</i>	72	92	EN	29	1	127	NE	27	275
<i>Sorbus lancastricensis</i>	72	100	NT	42	9	96	NE	35	273
<i>Sorbus leighensis</i>	72	93	EN	30	1	128	NE	28	279
<i>Sorbus leptophylla</i>	72	81	EN	18	3	104	NE	16	219
<i>Sorbus leyana</i>	72	66	CR	2	2	108	NE	2	178
<i>Sorbus margaretæ</i>	72	79	EN	16	5	100	NE	14	209
<i>Sorbus minima</i>	72	97	VU	37	1	131	NE	32	297
<i>Sorbus parviloba</i>	72	70	CR	6	1	116	NE	6	198
<i>Sorbus porrigentiformis</i>	72	102	LC	105	30	90	NE	37	334
<i>Sorbus pseudofennica</i>	72	98	VU	38	1	132	NE	33	301
<i>Sorbus pseudomeinchii</i>	72	71	CR	7	1	117	NE	7	202
<i>Sorbus richii</i>	72	86	EN	23	2	111	NE	21	241
<i>Sorbus rupicola</i>	72	107	LC	110	98	81	N	104	402
<i>Sorbus rupicoloides</i>	72	72	CR	8	1	118	NE	8	206
<i>Sorbus saxicola</i>	72	73	CR	9	1	119	NE	9	210
<i>Sorbus spectans</i>	72	94	EN	31	1	129	NE	29	283
<i>Sorbus</i>	72	82	EN	19	3	105	NE	17	223

stenophylla

<i>Sorbus stirtoniana</i>	72	74	CR	10	1	120	NE	10	214
<i>Sorbus subcuneata</i>	72	95	VU	35	4	103	NE	30	263
<i>Sorbus torminalis</i>	72	105	LC	108	573	60	N	102	375
<i>Sorbus vexans</i>	72	80	EN	17	4	102	NE	15	214
<i>Sorbus whiteana</i>	72	83	EN	20	3	106	NE	18	227
<i>Sorbus wilmottiana</i>	72	75	CR	11	1	121	NE	11	218
<i>Taxus baccata</i>	57	110	LC	113	1881	28	N	107	358
<i>Tilia cordata</i>	72	104	LC	107	896	51	N	101	363
<i>Tilia platyphyllos</i>	72	108	LC	111	84	82	N	105	406
<i>Ulmus glabra</i>	659	29	LC	71	2338	18	N	65	183

^a Red list Great Britain (GB): Not Evaluated (NE), Data Deficient (DD), Least Concern (LC), Near Threatened (NT), Vulnerable (VU), Endangered (EN), Critically endangered (CR).

^b Native status: Native (N), Native endemic (NE), Native or alien; Native status doubtful (NA).

Practical and Technical Considerations

The Millennium Seed Bank Seed Information Database (Royal Botanic Gardens, Kew 2008) was used to identify the seed storage behavior of target species. Recalcitrant *Quercus* species, and short lived Salicaceae were removed from the target list, as these cannot be dried and stored under conventional seed bank conditions or require careful post-harvest handling, not considered possible when using volunteer seed collectors for much of the collecting. Similarly, taxonomically complex woody taxa including most *Rubus* and *Rosa* species were removed to reduce problems of identification. The *Sorbus* micro-species were agreed as targets as, while taxonomically complex, they are well-recorded in the United Kingdom, have limited distribution, and there was an expert available to collect them. All *Ulmus*, apart from *U. glabra*, were removed because of the on-going effects of the Dutch elm disease (caused by species of *Ophiostoma*) outbreak which limits the ability of elm trees to reach maturity. *Ulmus glabra* is less susceptible than other *Ulmus* to Dutch elm disease, and has a greater propensity for sexual reproduction and therefore seed production. The resulting ranked target list was used as the basis for a project target list of 70 taxa which was agreed upon following further discussions by the project advisory group, a group of professionals from both the forestry and conservation sector brought together for oversight and technical advice.

Sampling Strategy

National Level

Unfortunately, most published research dealing with the genetic diversity and structure of United Kingdom woody species is only at the European scale, providing an insufficient basis for a national sampling strategy. The Forestry Commission (the United Kingdom government department responsible for forestry), has developed a seed zone map to guide seed transfer under the Forest Reproductive Material (Great Britain) Regulations 2002 legislation implemented under the EC Directive 1999/105. The map divides Britain into four broad regions of provenance and 24 smaller native seed zones; each zone is further divided by elevation above and below 300 m (Herbert et al. 1999) (fig. 1). Zones are based on biogeographic factors, such as watersheds and climate as well as man-made barriers which may influence dispersal and gene-flow, such as major roads. We decided to use the seed zone map as a framework for our sampling strategy, aiming for a seed collection in each seed zone where there is a native population for each of our target species, and an additional collection over 300 m, where this is possible. This standardised approach was adopted across all target taxa.

The native distribution of each target taxa was mapped against each seed zone using the Atlas of the British and Irish Flora (Preston et al. 2002), excluding the introduced distribution, based on historical pollen records and archaeological findings. Unfortunately, elevation analysis is not provided by the Atlas. The open access data from the Shuttle Radar Topography Mission (Jarvis et al. 2008), and records from the Botanical Society of Britain and Ireland database (BSBI 2016) were used to estimate the elevation of specific records. Estimates of occurrence over 300 m in seed zones were then confirmed on the ground by partners in seed zones 105, 108, 109 and 204, to validate the desk-based analysis, in order to produce a target list which is both vertically and horizontally distributed. This resulted in a total of 680 target seed collections.

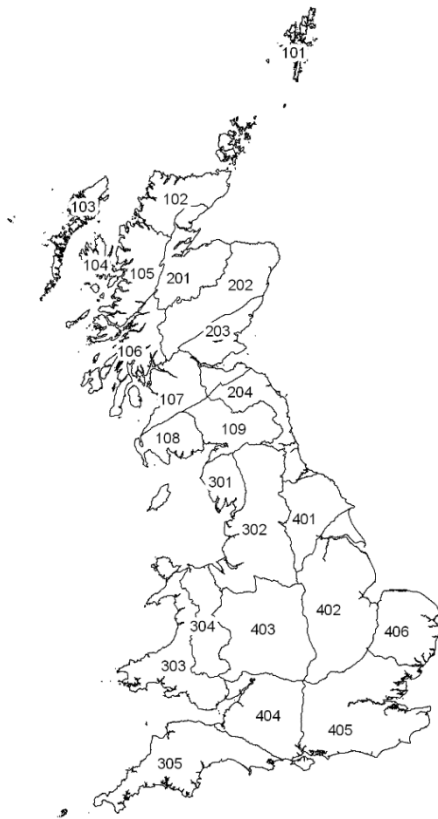


Figure 1—Native seed zones in Great Britain (Herbert et al. 1999).

This sampling strategy is a highly-effective way of sampling both common and rare locally adapted alleles (Hoban and Schlarbaum 2014), which are ecologically and economically important for conservation. Additionally, this form of sampling allows for conservation of the spatial structure of populations, which provides important data for research and restoration use of the collections. However, as we are using a standard strategy for all species which is based upon assumptions of genetic diversity and structure, rather than actual data, the strategy risks missing important species-specific factors and potentially over or under sampling.

One further implication of our sampling approach is that some of the populations collected from are at the northwestern edge of their ranges. Range-edges have profound implications on seed collecting programs, for example, marginal population may have decreased population density or reduced seed production (Garcia et al. 2000, Rasmussen and Kollmann 2004). Anecdotally, both the number of trees and number of seeds collected for the project have been less at range edges. Such peripheral populations may be adapted to the extremes possible for the species, and are, therefore, important sources of evolutionary potential, especially in the context of climate change (Fady et al. 2016, Havens et al. 2015).

The north and west are also the most deforested regions of the United Kingdom; therefore, also reducing seed sample sizes. The project is therefore re-assessing how to maximize sampling in these regions.

Sampling From Appropriate Sites

To meet our project aim, it is important that seeds are sampled from autochthonous, locally-adapted populations. As the United Kingdom is one of the least forested countries in Europe, with around 12 percent forest cover in contrast to averages in Europe of 45 percent and the United States of 40 percent (World Bank 2016a), this can be a challenge. Average woodland size (including exotic plantations), is only 7.9 ha in the United Kingdom overall, and only 4.9 ha in England (Watts 2006). The forest which once covered Britain was cleared for agriculture, mainly in the Bronze and Iron Age (Rackham 2001). The remaining ‘ancient semi-natural woodland’ (defined as having a continuous woodland cover since the earliest maps of 1600), is highly fragmented and now accounts for only 2 percent of land cover (Atkinson and Townsend 2011). The remaining woodlands also have a long history of management, usually by coppicing and favouring ‘useful’ species (Rackham 2001). There is, therefore, a high probability that any sampled population is not strictly autochthonous. There is an inevitable risk, therefore, that seed collected for the project may result from planted or naturalized trees originating from other parts of the United Kingdom or abroad, and in some cases form hybrid progeny. Indeed, the challenge is so great that even when we hand-pollinated native black poplars (*Populus nigra* subsp. *betulifolia*) and excluded pollen using bags, DNA analysis showed several resulting progeny had been pollinated by exotic, but abundant hybrid poplar (*Populus x euramericana*) (Gargiulo, personal communication, 2016). Resources such as the Native Woodland Survey of Scotland (Forestry Commission Scotland 2016), the Ancient Woodland Inventory (Natural England 2015), and sites protected under the Wildlife and Countryside Act (1981), and the EC Habitats Directive have been invaluable in guiding seed collectors to autochthonous populations. While every effort has been made to collect from such populations, we recognize that sampling is of the current gene pool, which includes introduced material.

Sampling Strategy at the Seed Zone

As a citizen science project working with a wide range of, mostly voluntary sector, partner organizations, we had to be realistic and pragmatic when setting an achievable sample size. Standard population sampling guidelines advise collection from either 30 randomly chosen individuals in a fully out-crossing species, or 59 random individuals in a self-fertilizing species (Brown and Marshall 1995) in order to collect 95 percent of the non-rare alleles present in a population. For many of our targets, collecting from this number of individuals in a population is not possible, because of the limited numbers of accessible seed-bearing plants. Even when populations were large enough, practical experience in the first year of the project demonstrated that collecting from 15 individuals in a day represented good effort. We therefore agreed to aim for sampling 10,000 seeds from at least this many individuals, where possible. This number of seeds is enough for some to be used for standard MSB germination tests, and allows for others to be available for distribution to researchers and conservation projects, while maintaining a sizeable sample in the seed bank.

In order to maximize the genetic diversity of the sample, collections were taken throughout the population, and over 50 m apart where possible; and seeds were collected from across the canopy at various places, to maximize the number of pollen sources. Individual mother trees were georeferenced using a GPS and were also physically tagged with a project aluminium tree tag. The seed from each mother tree was collected and stored separately in order to maximize the potential use of the collections for research and breeding. On every occasion, herbarium specimens accompanied a collection, for identification. In some cases tissue samples were collected for dry storage in silica gel, these can be used for DNA extraction and genomic analysis.

Building a Network of Seed Collectors

Although the United Kingdom is one of the least forested and most densely populated countries in Europe, with 267 people per km² compared to an European Union average of 116, and United States of 35

people per km² (World Bank 2016b), many people have easy access to local woodlands. In fact, 65 percent of the population live within 4 km of a 20 ha woodland (The Woodland Trust 2010). The link between health and wellbeing and access to natural environments, such as woodlands, is increasingly being valued (DEFRA 2012). It was important for the project, therefore, to develop an approach which capitalised on local community engagement in woodlands and forests, by working with conservation non-governmental organizations (NGOs). Working with land-owning NGOs also increased access to collecting sites, particularly useful as state ownership of woodlands and forests in the United Kingdom is also relatively low, at around 28 percent (DEFRA 2012).

Since the project began in 2013, seed collecting partnerships have been developed with over 30 organizations. Small local groups, for example Cree Valley Community Woodland Trust in Dumfries and Galloway, Scotland, or Suffolk Wildlife Trust, England, have collected the full target list in their respective seed zones. Locally-based NGOs, with access to woodland and existing experience in seed collecting with a team of engaged volunteers, are an incredible resource for the UKNTSP. Partners like this are able to monitor the availability and maturity of seed for collection without having to travel long distances. We also worked with the Forestry Commission, which led on collecting ash (*Fraxinus excelsior*), yew (*Taxus baccata*) and juniper (*Juniperus communis*) right across the United Kingdom.

In order to communicate the sampling strategy and collecting methodology to NGOs and volunteers, we produced a seed collecting manual (Kallow 2013), and ran training sessions across the United Kingdom each year. We developed a standard memorandum of collaboration and grant agreement in order to provide funding to partners and to agree the terms of ownership and transfer of seed collections, herbarium vouchers and data. We devised a new landowner consent form for use across the project partnership and made consent agreements with statutory agencies for collecting from protected sites. Working with these NGOs has also provided an opportunity to raise public awareness of forest conservation and *ex situ* conservation.

To conclude, the UKNTSP took into account scientific knowledge, existing national infrastructures, and engaged with local communities to build a seed repository which provides an important resource of genetic material and knowledge to help the conservation and management of trees and forests for the future.

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