

Collaborative action for updating the documentation on berry genetic resources in Europe

Monika Höfer

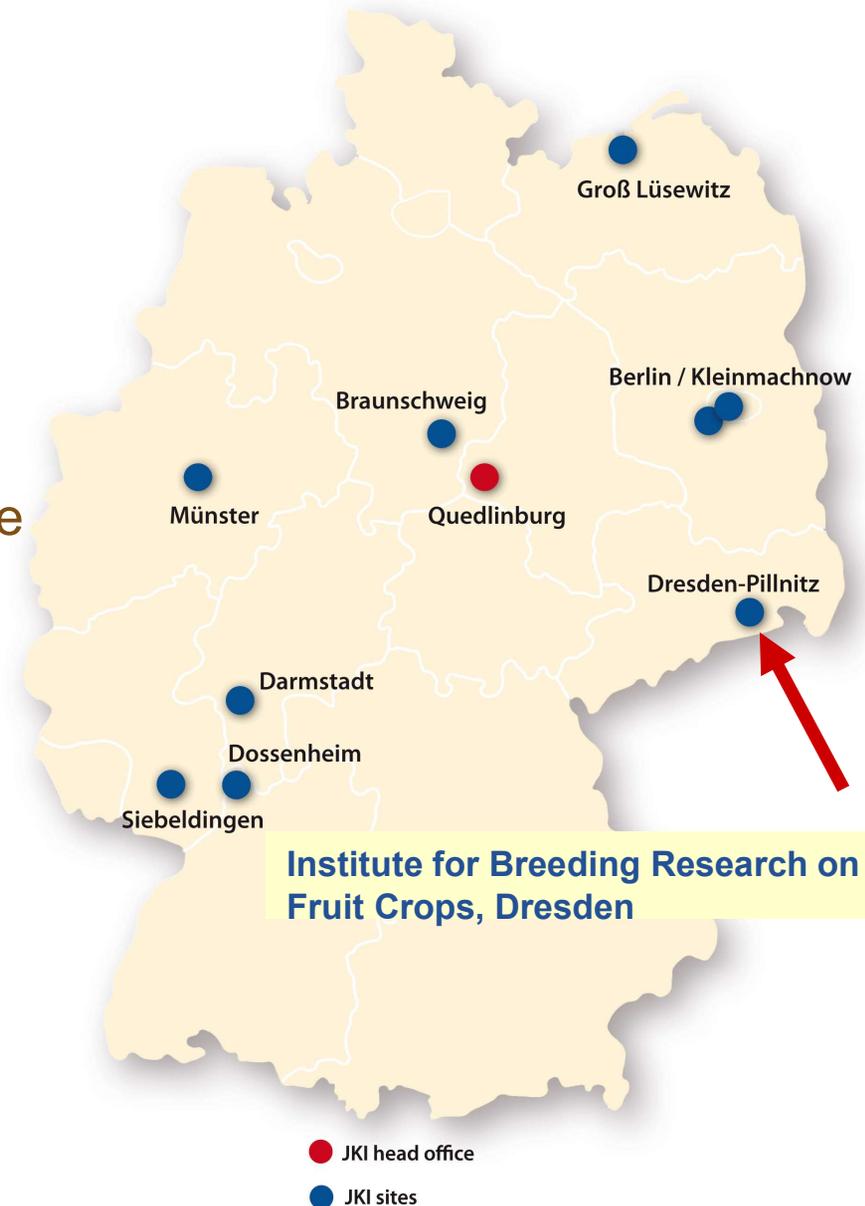
Julius Kühn Institute (JKI)

Institute of Breeding Research on Fruit Crops,
Dresden-Pillnitz, Germany



Julius Kühn-Institute, Federal Research Centre for Cultivated Plants

- ➔ **18 Research Institutes**
- ➔ **Total Staff: 1,180**
- ➔ **Head office in Quedlinburg**
- ➔ **Research field area:**
 - plant genetics, breeding research
 - plant nutrition, agronomy, soil science
 - plant protection and plant health



UNTIL 2019, THERE IS NO ECPGR WG FOR BERRIES. BUT EUROPEAN PROJECTS HAVE BEEN REALIZED SO FAR

Problems

1. only a limited number of partners/countries could be involved
2. the time frame was limited
3. the topic “genetic resources” covers only a very limited part

- Strawberry is on the Annex 1 list of the International Treaty on Plant Genetic Resources for Food and Agriculture
- ECPGR WGs for *Malus/Pyrus* (1995) and *Prunus* (1983 as one of the original six WGs) with a long active work

December 2018	Proposal to the ECPGR Executive/Steering Committee for the establishment of a Berries Working Group
January 2019	Expression of Interest to serve as ECPGR Working Group Chair during Phase X
March 2019	Nomination of the WG Chair
From April 2019	Nomination of the members of the WG by respective National Coordinators
January 2020	1. Meeting of the ECPGR WG Berries in Dresden



<http://www.ecpgr.cgiar.org/working-groups/berries/>

ECPGR Berries Working Group

ECPGR Homepage / ECPGR Working Groups / ECPGR Berries Working Group

51 ECPGR Berries Working Group Members

[✉ Email Distribution List](#)

Chair: Monika Höfer ✉ monika.hoefer@julius-kuehn.de

(22 countries)

[☰ Full list](#)

Genebank Curator (14)

Crop specialist (23)

Information/Documentation (8)

Plant breeder (19)

Policy and law (3)

Other expertise (8)

Contact Persons (11)

▾ LIST OF COUNTRIES

Albania

Denmark

France

Ireland

Lithuania

Romania

Sweden

United Kingdom

Bosnia and Herzegovina

Estonia

Germany

Italy

Norway

Serbia

Switzerland

Czech Republic

Finland

Greece

Latvia

Portugal

Slovenia

Turkey



18 members from 17 countries present

1. Meeting of the ECPGR WG Berries, 14-15 January 2020, Dresden, Germany

It was organized in collaboration with the Julius Kühn-Institut, Institute for Breeding Research on Fruit Crops, Dresden, Germany and financially supported by the German Federal Ministry of Food and Agriculture.



Proposal for a project for the Fourth Call for proposals under the ECPGR Activity Grant Scheme (Phase X).

‘Collaborative action for updating the documentation about berries genetic resources in Europe’

Expected impact.

1. With the implementation of the project, the varieties/ cultivars of the **berries genetic resources will be recorded** in the respective countries. The data will be harmonized, qualitatively evaluated and finally **will be available for inclusion to the respective National Focal Points for EURISCO.**
2. **A first draft of crop-specific technical guidelines for genebank management (Manuals) of berries genetic resources** will be elaborated and a related article will possibly be published.
3. Both results provide the basis for the future work of the Berries Working group. With the inventory of which berries genetic resources are conserved under which conditions in the gene banks of the European countries, **further projects on characterisation (phenotypic/molecular) will be derived in order to identify unique accessions for inclusion in AEGIS in the future.**



What is included in the genetic resources in berries?

In **botanical terminology**, a berry is a simple fruit with seeds and pulp produced from the ovary of a single flower, in which the complete pericarp is still juicy or at least fleshy when ripe (Kiger and Porter, 2001).

Traditionally and in everyday language, the term has different meanings and is usually used for small, sweet fruits, and the berry fruit.

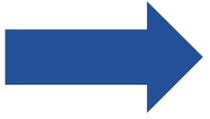
Berries under both definitions include blueberries, cranberries, lingonberries, and the fruits of many other members of the heather family, as well as gooseberries (*Ribes* L.), goji berries (*Lycium* L.) and elderberries (*Sambucus* L.)

The fruits of currants (*Ribes* L.), such as blackcurrants, red currants and white currants, are **botanical berries, even though their most commonly used names do not include the word "berry"**.

On the other hand, **several different kinds of fruit commonly called berries are not botanical berries**. Blackberries, raspberries, and strawberries are aggregate fruit; they contain seeds from different ovaries of a single flower.

Because a large number of plant species belongs to berry fruit genetic resources, the activities of the Berry WG will be diverse, and prioritization will take into account the economic importance of the crop.

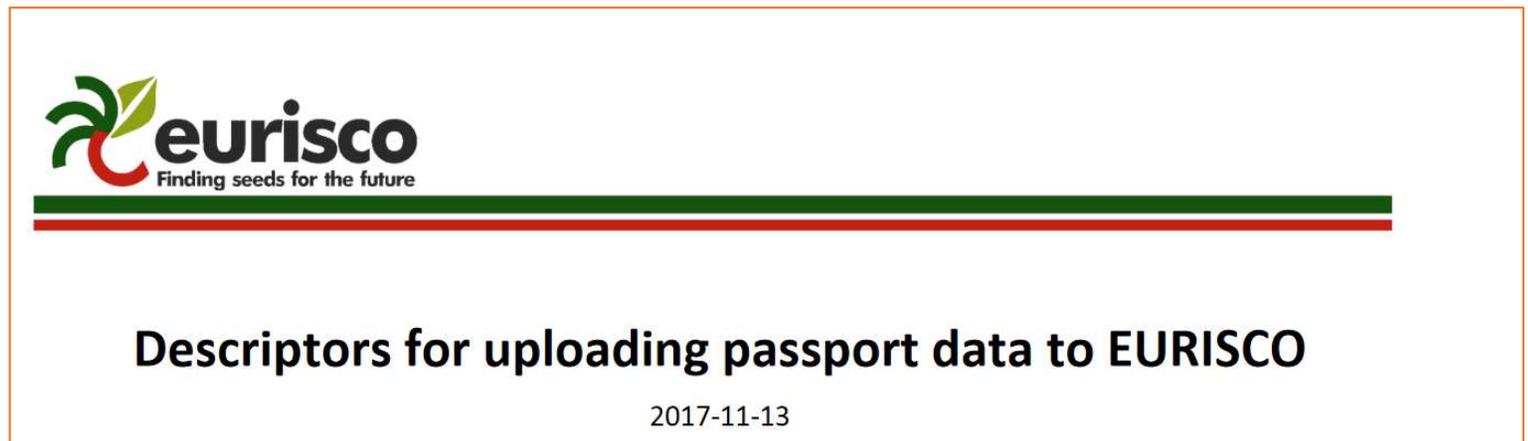




Documentation in **EURISCO** (<http://eurisco.ecpgr.org>)
European Search Catalogue for Plant Genetic Resources

For preparation of the inventories

1. Descriptors for uploading passport data to EURISCO (pdf-file)



2. Template for the inventory (excel-file) - with some additional descriptors important for our future work



NICODE INSTCODE ACCENUMB GENUS SPECIES SPAUTHOR CROPNAME ACCENAME ACQDATE ORIGCTY COLLSITE BREDCODE

DEU	DEU451	ERB0304	Fragaria	×ananassa	Duchesne ex Rozier	strawberry	Senga Sengana	20050801	DEU
DEU	DEU451	ERB0304c	Fragaria	×ananassa	Duchesne ex Rozier	strawberry	Senga Sengana	20050801	DEU
DEU	DEU451	ERB0304vf	Fragaria	×ananassa	Duchesne ex Rozier	strawberry	Senga Sengana	20050801	DEU

BREDNAME SAMPSTAT ANCEST

von Sengbusch, R.	500	Markee/Sieger'
von Sengbusch, R.	500	Markee/Sieger'
von Sengbusch, R.	500	Markee/Sieger'

19. Biological status of accession (SAMPSTAT)

The coding scheme proposed can be used at 3 different levels of detail: either by using the general codes (in boldface) such as 100, 200, 300, 400, or by using the more specific codes such as 110, 120, etc.

- 100) Wild**
 - 110) Natural
 - 120) Semi-natural/wild
 - 130) Semi-natural/sown
- 200) Weedy**
- 300) Traditional cultivar/landrace**
- 400) Breeding/research material**
 - 410) Breeder's line
 - 411) Synthetic population
 - 412) H₁
 - 413) Fo
 - 414) Inl
 - 415) Se
 - 416) Cl
 - 420) Ge
 - 421) M
 - 422) Cytogenetic stocks (e.g. chromosome addition/substitution, aneuploids, amphiploids)
 - 423) Other genetic stocks (e.g. mapping populations)
- 500) Advanced or improved cultivar** (conventional breeding methods)
- 600) GMO** (by genetic engineering)
- 999) Other** (Elaborate in REMARKS field)

20. Ancestral data (ANCEST)

Information about either pedigree or other description of ancestral information (e.g. parent variety in case of mutant or selection). For example a pedigree 'Hanna/7*Atlas//Turk/8*Atlas' or a description 'mutation found in Hanna', 'selection from Irene' or 'cross involving amongst others Hanna and Irene'.

DONOR CODE	DONORN AME	DONOR NUMB	DUPLINST DUPLSITE NAME	STORAGE	MLSSTAT	Virus state	Variety check
						1 = virusfree certified	1 = pomological
						2= virustested	2 = molecular
						3= unknown	3 = pom. + mol.
							4 = not checked
			BSA	20	1		3
				40	1	2	3
				20	1	2	3

26. Type of germplasm storage (STORAGE)
 If germplasm is maintained under different types of storage, multiple choices are allowed, separated by a semicolon (e.g. 20;30). (Refer to FAO/IPGRI Genebank Standards 1994 for details on storage type.)

10) Seed collection
 11) Short term
 12) Medium term
 13) Long term

20) Field collection

30) *In vitro* collection

40) Cryopreserved collection

50) DNA collection

99) Other (elaborate in REMARKS field)

27. MLS status of the accession (MLSSTAT)
 The status of the PGRFA with regard to the Multilateral System of Access and Benefit-Sharing (MLS) of the International Treaty on Plant Genetic Resources for Food and Agriculture. Leave the value empty if the status is not known.

0 No (not available under the MLS)
 1 Yes (available under the MLS)

Project partner providing inventories for the genera *Fragaria*, *Rubus*, *Ribes*, *Vaccinium* and other berry species

Partner ID No.	Member project group	Country	Institutsname	Fragaria	Rubus	Ribes	Vaccinium	Other genera
1	x	Lithuania	Lithuanian Research Centre for Agriculture and Forestry	1		1		
2	x	Germany	Federal Plant Variety Office - Testing Station Wurzen	1	1	1		
3	x	Finland	Natural Resources Institute Finland (Luke)	1	1	1	1	1
4	x	Slovenia	Agricultural Institute of Slovenia		1	1	1	1
5	x	Greece	Institute of Plant Breeding and Genetic Resources – HAO Demeter	1	1	1	1	1
6	coordinator	Germany	Julius Kühn Institute, Institute for Breeding Research on Fruit Crops	1	1			1
7		Czech Republic	Research and Breeding Institute of Pomology Holovousy Ltd.	1	1	1	1	1
8		Estonia	Estonian University of Life Science	1	1	1		1
9		France	INRAE	sent passport data				
10		France	INVENIO	1				
11		Italy	CREA Research Centre for Olive, Fruit and Citrus Crops	1				
12		Latvia	Institute of Horticulture LATHORT	1	1	1	1	1
13		Norway	Njøs Fruit and Berry Centre	1	1	1		1
14		Romania	Research Institute for Fruit Growing	1	1	1	1	1
15		Sweden	SLU/ Swedish National Genebank	1	1	1	1	
16		Schweiz	ProSpeciaRara	1	1	1		1
17		Albania	Agricultural University of Tirana		1		1	
18		Turkey	University of Cukurova	1	1	1		1
Total				15	14	13	8	11

Description of the work step	Who is responsible?	Which genus?
Video conference regarding the further verification work	Höfer, Monika	
1 Verification and completion of the inventories for the respective genera according EURISCO by the person responsible		
	Rugienius, Rytis and Sasnauskas, Audrius	<i>Fragaria</i>
	Schulte, Erik	<i>Rubus</i>
	Karhu, Saila	<i>Ribes</i>
	Koron, Darinka	<i>Vaccinium</i>
	Maloupa, Eleni	Other genera of berries genetic resources
Transfer of the data to EURISCO via the National Focal point of each country	all partners of the project	
2 Draft of crop-specific technical guidelines for genebank management (Manuals) of berry genetic resources	Höfer, Monika	
3 Select accessions for each genera in berries genetic resources for planning further joint projects	all nominated partners	
Project report	Höfer, Monika	

Methology

3 video conferences with all members of the project-working group

1st Activity meeting, online, 21 April 2021

2nd Activity meeting, online, 28 September 2021

3rd Activity meeting, online, 4 May 2022

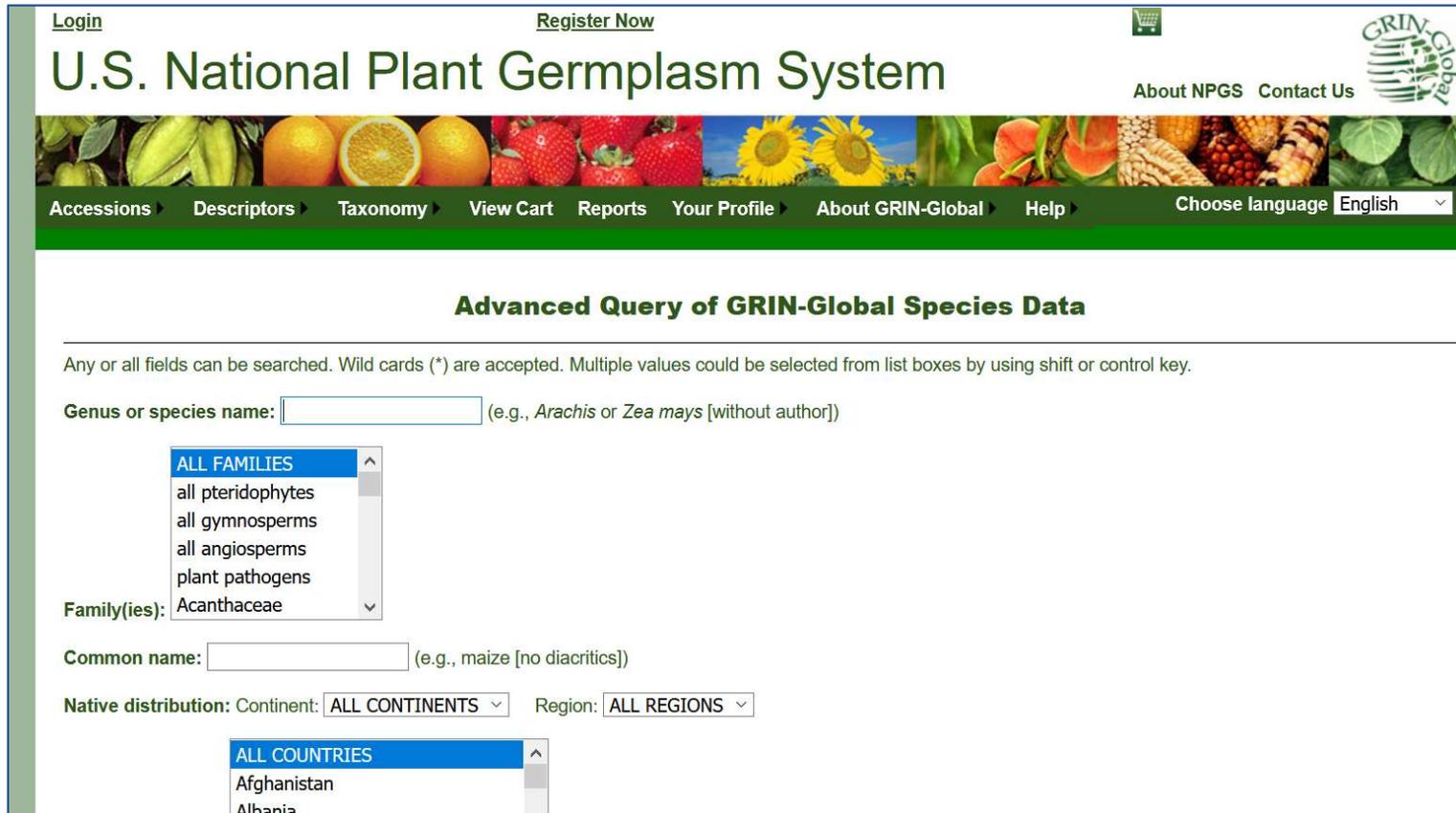
1. Documentation of berries genetic resources in Europe

the members of the project-group have got the inventories of berries genetic, should ask the partners for the missing data of the template or even for additional accessions and **to create a unified list of the accessions for each genus** with a deadline of September 2021.

Revision of the list concerning taxonomy (species) and variety names Integration of passport data

Differing indications of GENUS, SPECIES, and CROPNAME have been adjusted according to GRIN Taxonomy.

The GRIN Taxonomy should be used if possible (<https://npgsweb.ars-grin.gov/gringlobal/taxon/taxonomysearch>).



The screenshot displays the website's header with navigation links: [Login](#), [Register Now](#), [About NPGS](#), and [Contact Us](#). The main title is "U.S. National Plant Germplasm System" with the GRIN-Global logo. A horizontal banner features images of various crops like starfruit, oranges, strawberries, sunflowers, and corn. Below the banner is a green navigation bar with links: [Accessions](#), [Descriptors](#), [Taxonomy](#), [View Cart](#), [Reports](#), [Your Profile](#), [About GRIN-Global](#), [Help](#), and a language dropdown set to "English".

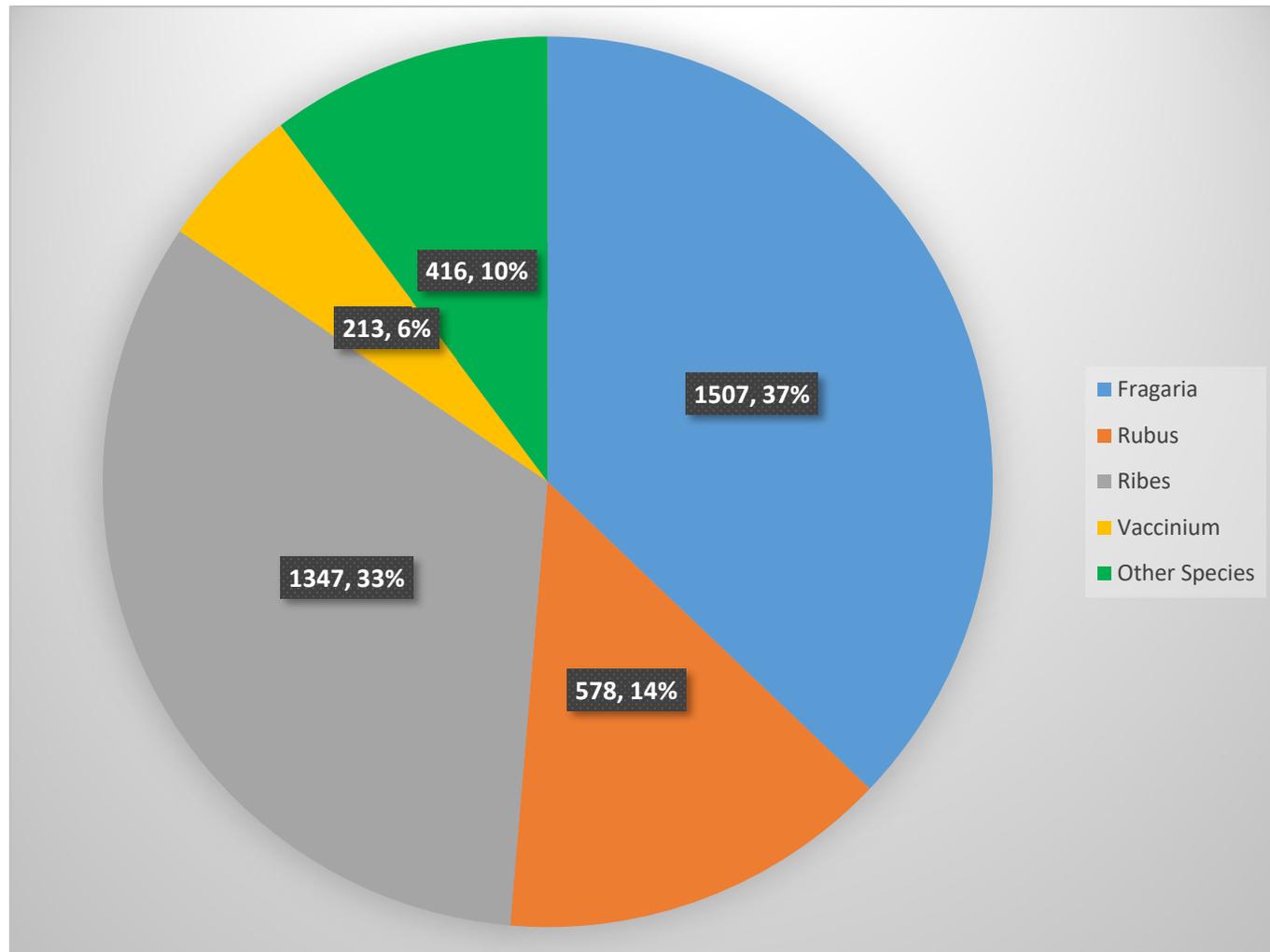
The main content area is titled "Advanced Query of GRIN-Global Species Data". Below the title is a search instruction: "Any or all fields can be searched. Wild cards (*) are accepted. Multiple values could be selected from list boxes by using shift or control key." The search form includes the following fields:

- Genus or species name:** A text input field with the example "(e.g., *Arachis* or *Zea mays* [without author])".
- Family(ies):** A dropdown menu currently showing "Acanthaceae". The menu is open, listing options: "ALL FAMILIES", "all pteridophytes", "all gymnosperms", "all angiosperms", and "plant pathogens".
- Common name:** A text input field with the example "(e.g., maize [no diacritics])".
- Native distribution:** Two dropdown menus: "Continent:" set to "ALL CONTINENTS" and "Region:" set to "ALL REGIONS".
- Country:** A dropdown menu currently showing "ALL COUNTRIES". The menu is open, listing "Afghanistan" and "Albania".

1. **Variety denominations** have been adjusted according to the rule that a variety's name is considered the original one, which is used in the country of its origin,
2. **All other known denominations**, whether submitted by the partners or taken from pomological literature, of a variety have been **listed as known synonyms**.
3. **Passport data have been included**, partly from the individual partners' side (e.g. INSTCODE, ACCENUMB, ACQDATE, ORIGCTY, COLLSITE, SAMPSTAT, DONORNAME, STORAGE, Virus state, and Variety check), or added by the project-group members from the pomological literature (e.g. BREDNAME, ANCEST).
4. Passport data could be used from both the EU GENBERRY project (INRAE, France) project database and the RIBESCO project (Luke, Finland).



Results

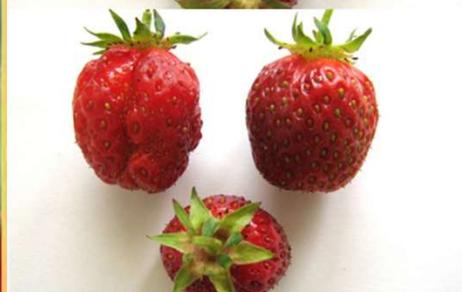
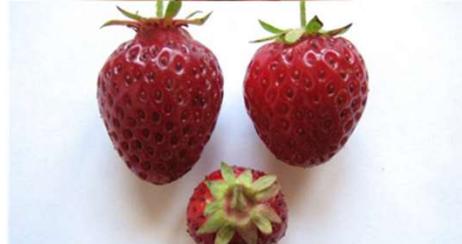
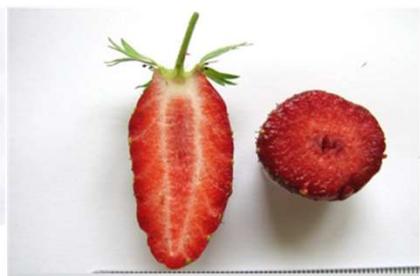


Altogether 4.061 accessions of beery genetic resources were mentioned in the inventories



GENUS	SPECIES	No Acc
<i>Fragaria</i>	<i>chiloensis</i>	11
<i>Fragaria</i>	<i>hybr.</i>	3
<i>Fragaria</i>	<i>moschata</i>	69
<i>Fragaria</i>	<i>spp.</i>	1
<i>Fragaria</i>	<i>vesca</i>	51
<i>Fragaria</i>	<i>virginiana</i>	4
<i>Fragaria</i>	<i>viridis</i>	1
<i>Fragaria</i>	<i>x ananassa</i>	1355
<i>Fragaria</i>	<i>x vescana</i>	12

1507 acc.



Fragaria chiloensis
(photo: K. Olbricht)



X *Fragaria virginiana*



For *Fragaria*, the common list contained **762 different *F. ×ananassa* cultivars**.

- Among them 537 were unique accessions, stored in only one of the collections.
- Some cultivars (Honeoye, Senga Sengana) are held in almost all collections.
- **Information on the virus status of strawberry and the method of cultivar identification in various institutions was also collected.** Among accessions 67.1% virus status was unknown, 30.7 % virus tested, 2.2 % virus free certified.
- **Pomological description of accession was most frequent method for variety identification**, but also molecular technology was in use in some institutes: for 67 % of the mentioned accessions, pomological and/or molecular trueness-to-type determination has already been realised.

<i>Rubus</i>	<i>Idaeus</i> - raspberry	439
<i>Rubus</i>	<i>sect. Rubus</i> - blackberry	86
<i>Rubus</i>	<i>occidentalis</i>	14
<i>Rubus</i>	<i>arcticus</i>	10
<i>Rubus</i>	<i>hybr.</i>	16
<i>Rubus</i>	<i>x binatus</i>	4
<i>Rubus</i>	<i>idaeus x allegheniensis</i>	2
<i>Rubus</i>	<i>phoenicolasius</i>	2
<i>Rubus</i>	<i>allegheniensis</i>	2
<i>Rubus</i>	<i>chamaemorus</i>	1
<i>Rubus</i>	<i>nessensis</i>	1
<i>Rubus</i>	<i>ulmifolius</i>	1

578 acc.



cv. Black Jewel (*R. occidentalis*)

photo Lubera)



cv. Dorman Red (*Rubus parviflorus* x *Rubus idaeus* 'Dorset'),



<i>Ribes</i>	<i>aureum</i>	4
<i>Ribes</i>	<i>dikusha</i>	1
<i>Ribes</i>	<i>nigrum</i> – black currant	451
<i>Ribes</i>	<i>Rubrum</i> – red currant	442
<i>Ribes</i>	<i>x niveum</i>	4
<i>Ribes</i>	<i>spicatum</i>	12
<i>Ribes</i>	<i>uva-crispa</i> - gosseberry	431
<i>Ribes</i>	<i>x nidigrolaria</i>	1
<i>Ribes</i>	<i>spp.</i>	1
<i>Ribes</i>	<i>sp.</i>	2

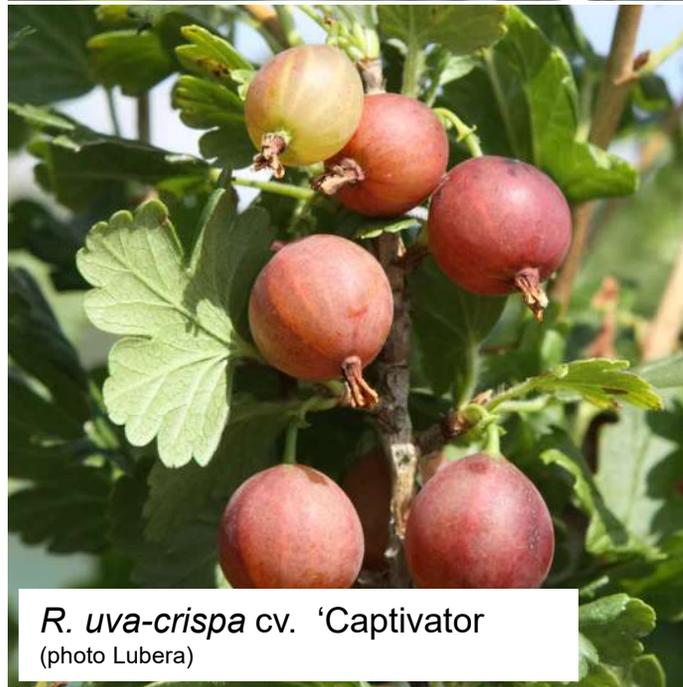
1347 acc.



R. rubrum (red currant cv. ‚Jonkheer van Tets‘
(photo Horstmann))



R. nigrum (black currant) cv. Titania
(photo Lubera)



R. uva-crispa cv. ‚Captivator‘
(photo Lubera)

<i>Vaccinium</i>	<i>angustifolium</i>	2
<i>Vaccinium</i>	<i>corymbosum</i>	113
<i>Vaccinium</i>	<i>hybrid</i>	16
<i>Vaccinium</i>	<i>macrocarpon</i>	8
<i>Vaccinium</i>	<i>myrtilus</i>	51
<i>Vaccinium</i>	<i>oxycoccos</i>	1
<i>Vaccinium</i>	<i>sp.</i>	4
<i>Vaccinium</i>	<i>spp.</i>	3
<i>Vaccinium</i>	<i>uliginosum</i>	1
<i>Vaccinium</i>	<i>vitis-idaea</i>	14

213 acc.



Vaccinium myrtilus - blueberry
(photo Horstmann)



Vaccinium corymbosum 'Bluecrop, - American bluebeery
(photo Horstmann)



Vaccinium vitis-idaea - lingonberry
(photo Horstmann)

At the beginning of the project, the project-working group discussed what are genetic resources in berries and what should be included in particular in the term **‘other genera of berries genetic resources’**.

<i>Actinidia</i>	<i>arguta</i>	7
<i>Amelanchier</i>	<i>alnifolia</i>	5
<i>Amelanchier</i>	<i>ovalis</i>	1
<i>Aronia</i>	× <i>prunifolia</i>	1
<i>Aronia</i>	<i>melanocarpa</i>	4
<i>Cornus</i>	<i>mas</i>	17
<i>Hippophae</i>	<i>rhamnoides</i>	93
<i>Lonicera</i>	× <i>caerulea</i>	2
<i>Lonicera</i>	<i>caerulea</i>	27
<i>Mahonia</i>	<i>aquifolium</i>	1
<i>Morus</i>	<i>alba</i>	112
<i>Morus</i>	<i>nigra</i>	25
<i>Morus</i>	<i>rubra</i>	2
<i>Morus</i>	<i>sp.</i>	22
<i>Rosa</i>	<i>canina</i>	9
<i>Rosa</i>	<i>damascena</i>	1
<i>Rosa</i>	<i>dumalis</i>	1
<i>Sambucus</i>	<i>nigra</i>	51
<i>Sorbus</i>	× <i>hybrida</i>	1
<i>Sorbus</i>	× <i>meinichii</i>	1
<i>Sorbus</i>	<i>aucuparia</i>	24
<i>Sorbus</i>	<i>intermedia</i>	2
<i>Sorbus</i>	<i>latifolia</i>	1
<i>Sorbus</i>	<i>rupicola</i>	1
<i>Sorbus</i>	<i>sp.</i>	5



Morus alba

Based on the discussion about the definition of berries genetic resources another specific request was sent to the partners for the **‘other genera of berries genetic resources’**.

Altogether, 161 acc. of *Morus* (mulberry), 51 acc. of *Sambucus nigra* (elderberry), 93 acc. of *Hippophae rhamnoides* (Sea buckthorn) and 35 acc. of *Sorbus aucuparia* (mountain ash) were reported.

Current status of documentation in EURISCO

No.	Country	Contact	INSTCODE	Institutsname	Fragaria	Rubus	Ribes	Vaccinium	other genus	Confirmation	Remarks
1	Lithuania	Rytis.Rugienius@lammc.lt	LTU006	Lithuanian Research Centre for Agriculture and Forestry	1		1			17.10.2023	
2	Germany	Erik.Schulte@bundessortentamt.de	DEU610	Federal Plant Variety Office - Testing Station Wurzen	1	1	1		1	yes	via German Fruit Genebank
3	Finland	saila.karhu@luke.fi	FIN039	Natural Resources Institute Finland (Luke)	1	1	1	1	1	17.10.2023	First via GenBIS and NordGen
4	Slovenia	Nika.CvelbarWeber@kis.si	SVN019	Agricultural Institute of Slovenia		1	1	1	1	07.11.2023	
5	Greece	maloupa@bbgk.gr	GRC044	Institute of Plant Breeding and Genetic Resources – HAO Demeter	1	1	1	1	1	16.10.2023	Position National Focal Point unfilled
6	Germany	monika.hoefer@julius-kuehn.de	DEU451	Julius Kühn Institute, Institute for Breeding Research on Fruit Crops	1	1			1	yes	via German Fruit Genebank
7	Czech Republic	Boris.KRSKA@vsuo.cz	CZE031	Research and Breeding Institute of Pomology Holovousy Ltd.	1	1	1	1	1	19.10.2023	
8	Estonia	Hedi.Kaldmaee@emu.ee	EST012	Estonian University of Life Science	1	1	1		1		
9	France	beatrice.denoyes@inrae.fr		INRAE	sent passport data						
10	France	j.perrotte@invenio-fl.fr	FRA207	INVENIO	1						
11	Italy	gianluca.baruzzi@crea.gov.it	ITA380	CREA Research Centre for Olive, Fruit and Citrus Crops	1					17.10.2023	
12	Latvia	valda.laugale@llu.lv	LVA015	Institute of Horticulture LATHORT	1	1	1	1	1	18.08.2023	
13	Norway	dag.roen@graminor.no	NOR053	Njøs Fruit and Berry Centre	1	1	1		1	24.11.2023	First via GenBIS and NordGen
14	Romania	sturzeanu1980monica@yahoo.it	ROM009	Research Institute for Fruit Growing	1	1	1	1	1	08.05.2023	
15	Sweden	inger@pomologi.nu	SWE089	SLU/ Swedish National Genebank	1	1	1	1		18.10.2023	First via GenBIS and NordGen
16	Schweiz	claudio.niggli@prospecierara.ch	CHE063	ProSpeciaRara	1	1	1		1	17.10.2023	
17	Albania	albanibrailiu@ubt.edu.al	ALB026	Agricultural University of Tirana		1		1		19.10.2023	
18	Turkey	ebruyasakafkas@gmail.com	TUR029		1	1	1		1	17.10.2023	Position National Focal Point unfilled

Elaboration of crop-specific technical guidelines for genebank management of berries genetic resources

The first query on the existing accessions also included the descriptor '**Type of germplasm storage**' (EURISCO 26 – STORAGE).

- Most of the temperate fruit species are genetically heterozygous and vegetatively propagated.
- A seed collection will not represent the true genotyp of clonal accessions (cultivars).
- The collections of berries genetic resources are maintained in the field or greenhouses as active plantations where the accessions are available for characterization, evaluation, and distribution.
- Backups for the plant material are needed to provide security in case of a disease or an environmental disaster.
- The duplicate collection could be a second field site, an *in vitro* culture or a cryopreservation.





JKI: Potted plants of the *ex situ* strawberry collection



JKI: In vitro cold storage



Federal Plant Variety Office: Field collection



JKI: virus-free material maintained in an insect-protected screen house



JKI: Facility for cryopreservation

Federal Plant Variety Office, Germany: Insect-protected screen house for raspberries, photo: BSA



Natural Resources Institute Finland: Field collection of blackcurrant, photo: Luke



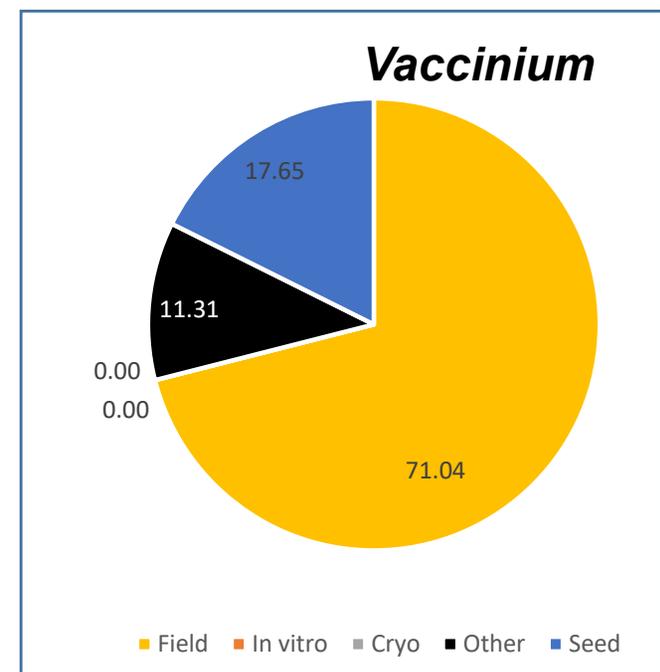
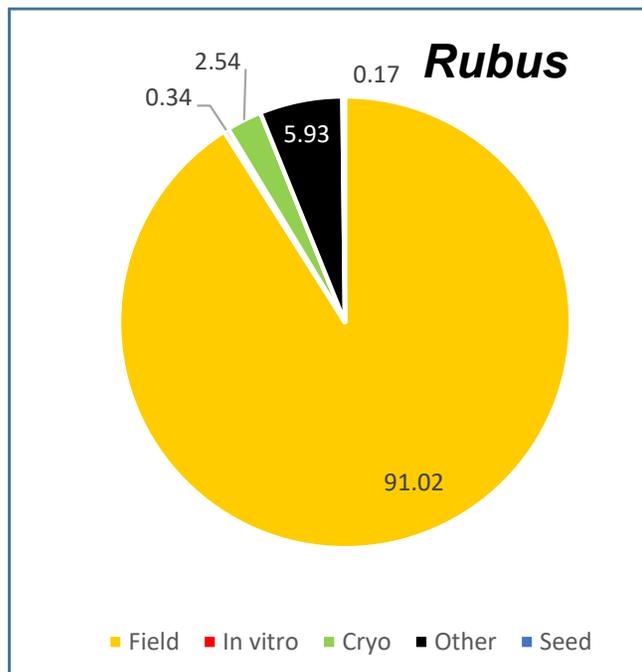
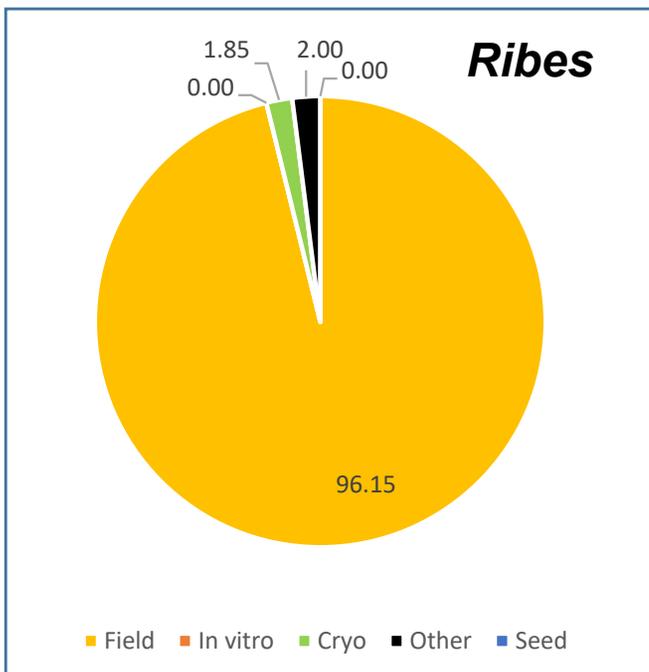
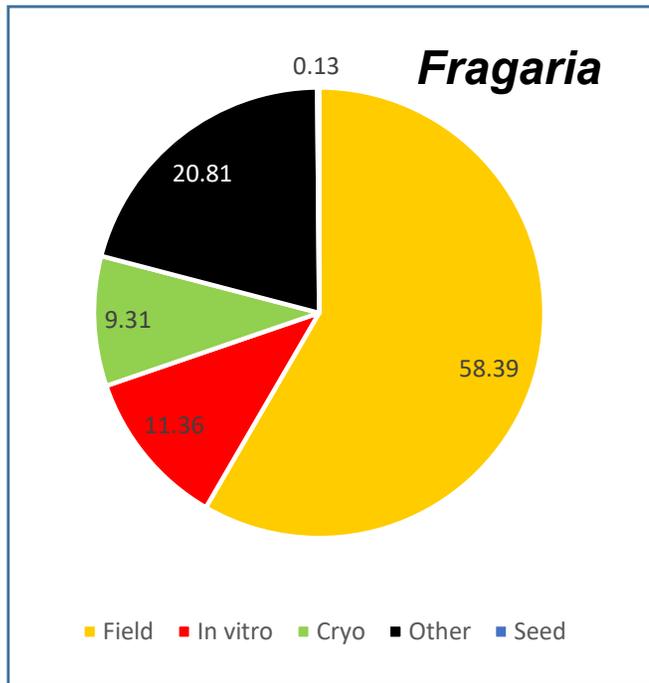
Hellenic Agricultural Organization Demeter, Greece: Dog rose collection, photo: Demeter



F Agricultural Institute of Slovenia: Field collection of blueberries, photo: Agricultural Institute of Slovenia

Percentage distribution of accessions in the various methods of preserving genetic resources in berries

Only in the case of *Fragaria* the *in vitro* culture and cryopreservation makes a significant contribution to build up a duplicate.



- Based on the results of the first request regarding the forms of preservation a **second template** was developed, which was sent to all project partners, **in order to request the detailed methods of conservation.**
- The template contained exact questions on the cultivation of the plants for the *ex situ* collection in the field or in the greenhouse as well as for the *in vitro* and cryo collection.

For the *ex situ* collections the following points were included:

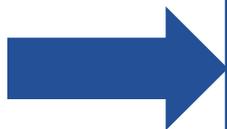
Location of the genebank	Horticultural work
Soil	How is the phytosanitary control organized? Pest and diseases control
Type of planter containers	Disease prevention / integrated pest management
Type of plants	Replanting time of the plantation
Which plant labels are used?	How is the replantion realized?
How many plants per accession?	Periodically verifying the identity of the accession?
Planting time	Could you realize characterization / evaluation work?
Isolation distance of the plants (cm/ m)	Safety duplication
Field management	



Contributions to the query were made by all 18 partners by the end of 2022, so that the preparation of the first draft of crop-specific technical guidelines for genebank management of berries genetic resources could be started.

The *Prunus* specific standards for genebank management served as an example (Giovannini et al. 2016).

This first draft will be sent to all members of the Working group for discussion.



The aim is to establish **quality standards of conservation berries genetic resources (minimal standards)** and to publish these on the webpage of ECPGR.



All objectives of the EUROPE.BERRIES project were achieved.

- An extension to the initially planned time was requested due to different reasons. The main reason was that for the individual partners of the action, working with the EURISCO passport data was new area, so that many explanations were necessary and the work was delayed.
- The situation was the same for the some members of the project-group who had volunteered to revise the inventory / passport data.

RECOMMENDATIONS – PERSPECTIVE

- This action gave the newly established working group the opportunity **to work together for the first time within the framework of the ECPGR** and to make themselves known with its working methods of the ECPGR.
- A grant will be requested for a second group meeting to present the results to a majority of the members and to discuss further projects.
- The results of the project will also be presented at the next Progress Meeting of the EU project 'Breeding value' in order to identify common aims for the further work on genetic resources in berries.
- The project was the basis for establishing the first step towards a long-term work for conservation/ characterisation/ documentation and utilisation of genetic resources in berries.





THANK YOU