

Final Results of the AHA on Development of specifications for a modern gene bank documentation software (CryoWEB 2.0) and further steps

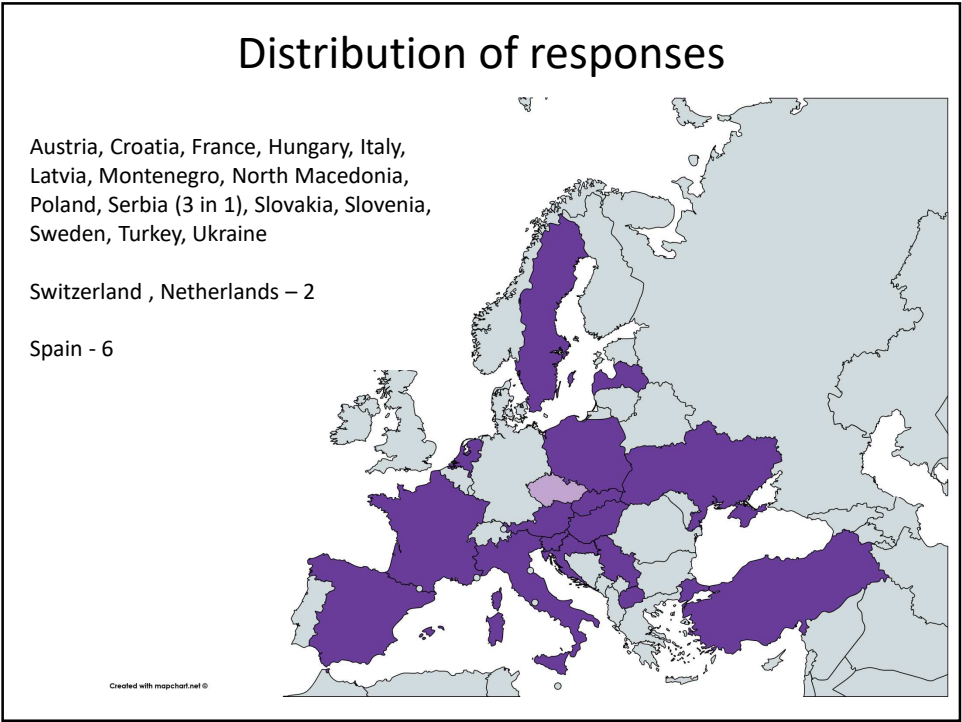
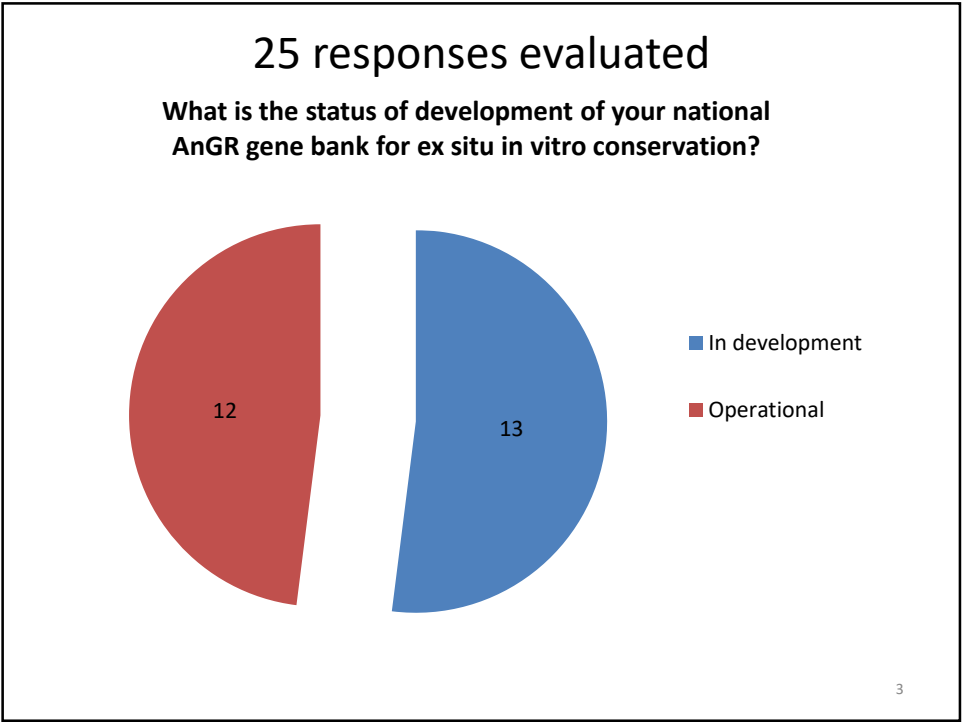
ERFP WG Ex situ conservation
meeting
02.07.2021

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Aim of the survey

- Review current status of documentation of farm animal gene banks (ex situ in vitro) across Europe
- Identify the information needs of the various countries
- Help preparing functional specifications for modern gene bank documentation software

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Respondents grouped by number of breeds

Group A – 2 gene banks

- Breeds >140
- Species >10
- Donors >7000
- Samples >300000

Group C – 7 gene banks

- Breeds 10-20
- Species 2-7
- Donors ?-140
- Samples 250-60000

Group B – 5 gene banks

- Breeds 30-55
- Species 5-8
- Donors 400-4000
- Samples 73000-205000

Group D – 11 gene banks

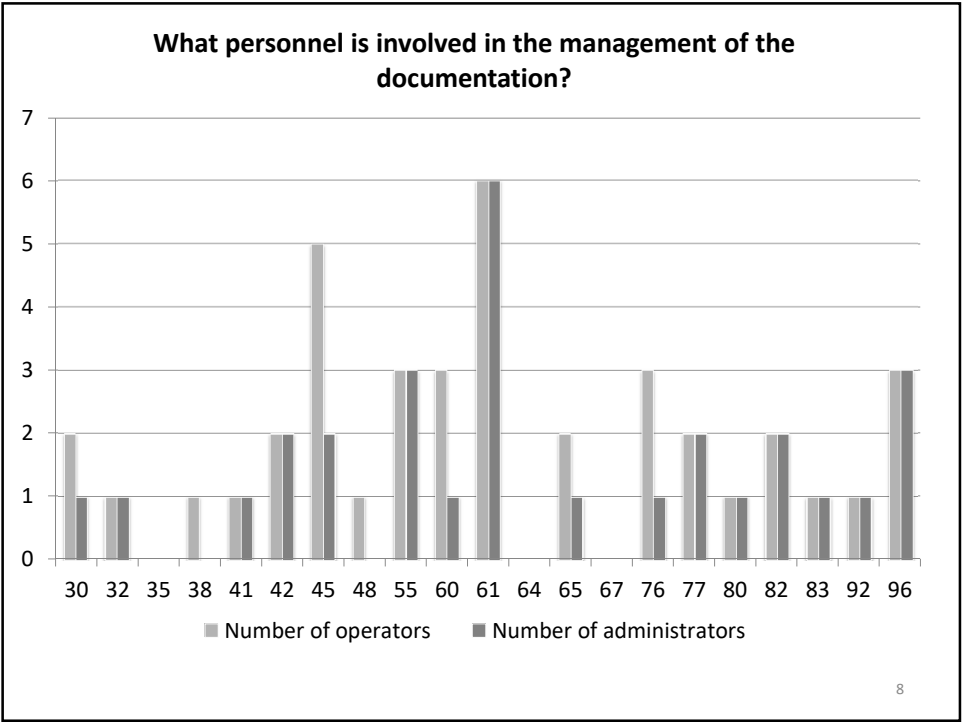
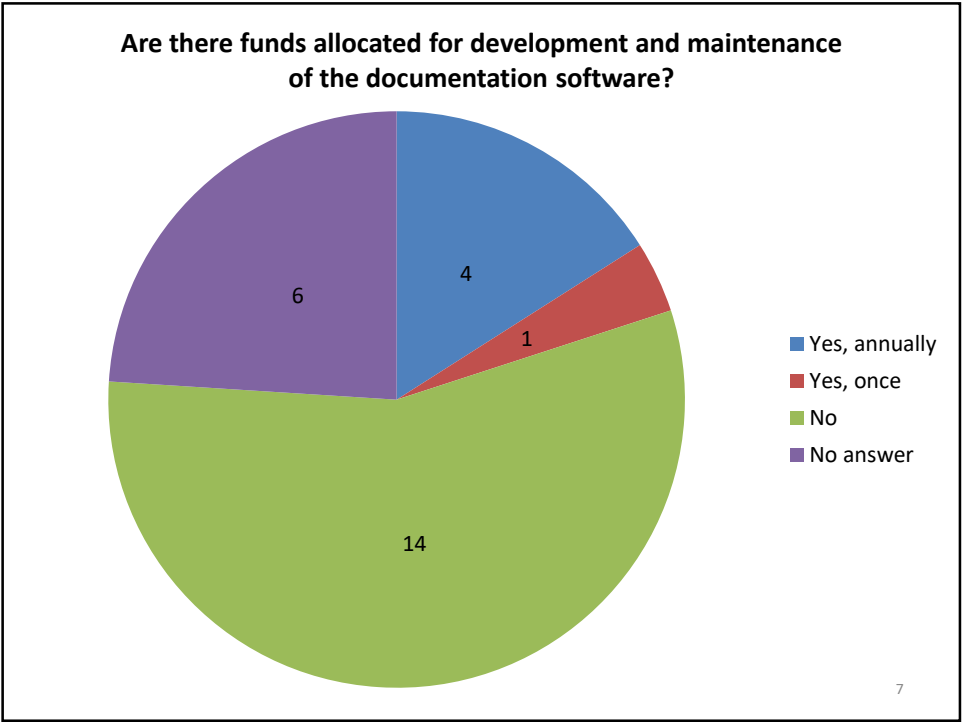
- Breeds 2-9
- Species 1-6
- Donors 20-1600
- Samples ?-52000

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Persons employed by the national gene bank

- Most of the respondent gene banks report:
 - less than 6 employed persons
 - less than 3 full time equivalent persons
- The 2 largest gene banks report 2 or less full time equivalent persons

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Setup and documentation

- Gene banks are very different in size and organization
- Material of choice – semen
- Mostly core collections
- Mostly centralized management by public institution
- Limited personnel
- Documentation is also heterogeneous
- Heavy usage of Spreadsheets
- Some data still on paper
- Data is not open to the general public
- No funds allocated for development and maintenance of documentation software

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Wish list

- Data upload to EFABIS
- More flexibility in reports creation (e.g. reports by organization)
- Control of data access dependent on the user - creation of IS to serve more genebanks, so the users can only see their own data and samples
- Easy export functionality (CSV file)
- Counter to easily determine how many straws are still kept on the tank from a specific session and for each specific animal
- Interface deleting number of doses of multiple samples
- Genomic data storage - link to our genebank database
- The possibility to import data from a template spreadsheet (e.g. format xls)

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No common action

	Country level	ERFP level
Costs	+++	+
Personnel	+++	+
Maintenance	+++	+

Each country develops the software it requires on its own

Individual country joins Animal-Grin as a full-member

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Common action for all EUGENA members

	Country level	ERFP level
Costs	+	++
Personnel	+	++
Maintenance	+	++

ERFP develops simple software for gene bank documentation, in order for the EUGENA member countries to meet minimum data standards of documentation (i.e. CryoWEB 2.0).

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Common action for all ERFP members

	Country level	ERFP level
Costs	+	+++
Personnel	+	+++
Maintenance	+	+++

ERFP develops and maintains a common software that can be used by all countries

ERFP joins Animal-Grin as representative for Europe and provides required permanent personnel.

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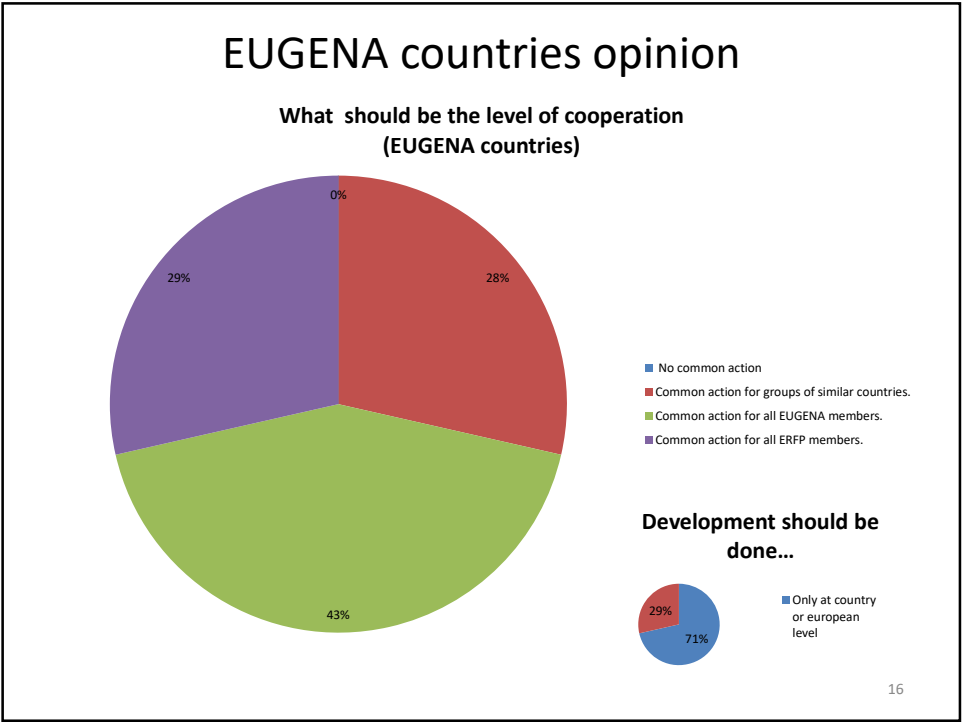
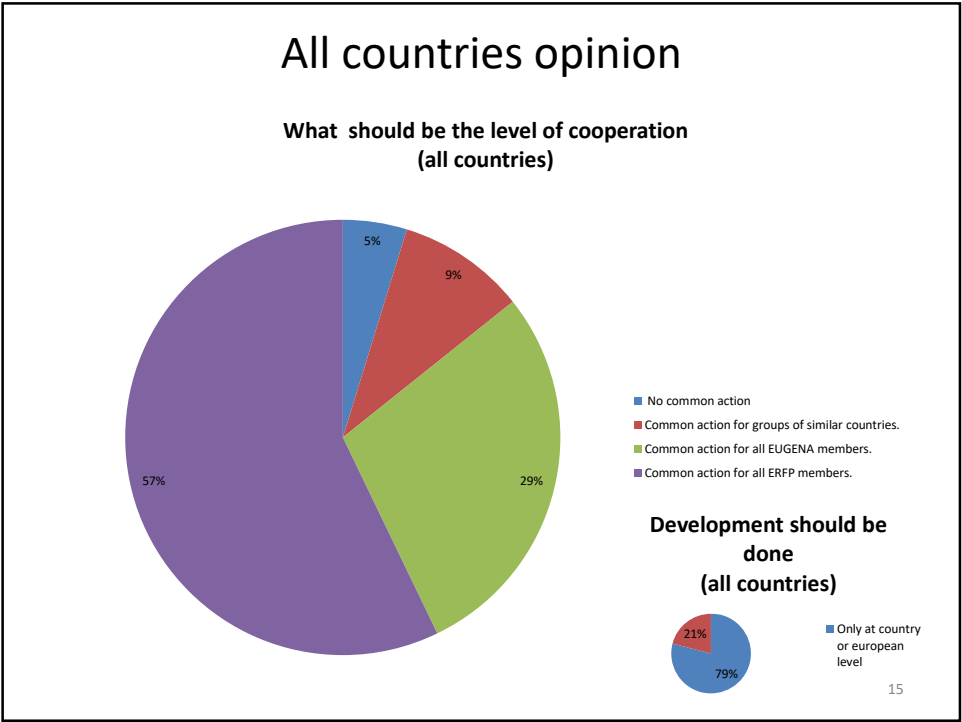
Common action for groups of similar countries

	Country level	ERFP level
Costs	++	+
Personnel	++	+
Maintenance	++	+

Groups of countries with similar requirements develop common software that suits their needs

Groups of countries with similar requirements join Animal-Grin

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BASIC REQUIREMENTS

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General requirements

- Open source
- Follow the FAIR principles
- User interface accessible via LAN or WAN over a secure connection (optional two-factor authentication)
- User interface in the national language
- All user actions should be recorded and kept in history
- Option to revert the last action

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Modules

- Donor animals
- Storage facilities
- Samples
- Reports and queries
- Requesting of material
- Users management
- Journal log
- Backup and restore

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Donor animals

- Registration of new
- Modification of donor data
- Deletion of an animal
- Search for animals with certain characteristics
- Adding other data fields to the donor record
- Visualization of the animals on the map with links to the animal data and stored material

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Storage facilities

- Describe a storage facility by defining the number of levels, their types and identification
- Import storage facility structure from file
- Export selected storage facilities structure to file
- Move selected elements of one facility (with or without their germplasm content) to another
- Modify storage facility data (e.g. rename tank)
- Visualize the storage facilities, allowing the user to browse through the levels

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Samples

- Record new sample with its distribution in the storage facilities
- Import new sample data from file into the database
- Move selected samples from one location another via the web interface
- Move samples to another location reading data from file
- Record sample usage (taking samples out from the gene bank)
- Modify sample data (e.g. correct vessel identification)
- Erase sample from the system
- Keep history of all sample movements and usage
- Search for a sample by number of parameters provided by the user (e.g. samples from a certain donor, all material stored on a certain date, etc.)
- Provide user with information about:
 - The location of a certain sample on a certain date
 - The content of a certain storage facility on a certain date
 - History of all movements of a sample

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Reports and queries

- Drill down browsing of the collection starting from species->donors->samples
- Drill down browsing of the collection starting from certain storage facility
- Browsing the collection by starting from map of donor locations->samples
- Search for germplasm material by set of parameters (e.g. by species, by donor, etc.)
- Generate general statistics about donors per sample type, donors per breed and species, samples per location, samples per breed and year
- Export aggregated statistics data for upload in EFABIS/DAD-IS
- Export aggregated statistics data for upload in EUGENA
- Export all the information related to one level of the storage facilities (e.g. one tank) in structured format
- Export all the information related to one donor in structured format
- Export all the information related to one sample in structured format

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Requesting of material

- Allow organizations and persons to request material from the gene banks
- Allow gene banks managers to approve or reject a request
- Allow reservation of material (after approved request) in order to block it from using for other purposes

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Users management

- Adding new
- Modifying user
- Changing user access rights
- Blocking user

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Journal log

- Provides the gene bank manager and administrator with information about the user activities
- Daily/monthly report with the log in and log out times of all users
- Report about the modifications done by a certain user
- Report about the modifications done in certain module

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Backup and restore

- Archives all the data to external file on a regular interval
- Allows restoring of the database to its state at a certain date from external archive file

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Ad hoc action members

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- Sipke-Joost Hiemstra - Netherlands
- Ina Hulsege - Netherlands
- Eva-Marie Stålhammar – Sweden
- Ewa Sosin-Bzducha - Poland
- Delphine Duclos - France
- Nataliia Reznikova – Ukraine
- Danijela Bojkovski - Slovenia
- Zhivko Duchevev - Bulgaria
- Fernando Tejerina Ampudia - Ex Situ WG

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