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How to continue the work, some

- proposals

 ➤ The IMAGE self-diagnostic tool seems to be a good way to identify strengths and points to be developed. The total score can indicate whether the process is advanced or not (e.g.: more than 60 points, the process is almost complete)
- ➤ Facilitate the exchanges of document templates needed for a QMS (for example: table of relevant stakeholders, PESTEL analysis for external factors, SWOT analysis...).
 - => More generally, what kind of documents/protocols could be shared (translated) between genebanks for inspiration at ERFP level?
- > Organise a peer review/on-site visit system for



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To be continued?

- Seems interesting to pursue with new genebanks (3 / 2 years?)
- ➤ New ad hoc action needed or part of EUGENA budget?
- ➤ Who will coordinate it?

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New web tools in EUGENA

Issue: Breeds names in EUGENA and DAD-IS/EFABIS are not synchronized

Goals:

Use of the FAO persistent identifiers in EUGENA

Automated exchange of the breeds names from DAD-IS to EUGENA

Entering data for breeds which are currently not present in DAD-IS

Option to export genebanks collection data at country level in a format suitable for upload in EFABIS

Led by Z. Duchev

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New web tools in EUGENA

Objectives and... all done!

- ✓ Extending the database with fields to link DAD-IS data
- ✓ Develop automated loading of breeds and species names in EUGENA
- ✓ Develop automated process for checking for changes and applying the updates
- ✓ Development of additional form for indexing non DAD-IS species and breeds by the administrator
- Extend EUGENA database with a field indicating implementation of QMS system and including this field in the data entry form and the gene bank contact data

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