Creating synergy between EURISCO and GeneSys

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EURISCO: currently passport data, upload mechanism and network of data suppliers

passport data already included in GeneSys

EURISCO and Genesys

- both want to handle C&E data
- how does it fit how to benefit from each other
 - in any case: avoid double work !



 EURISCO proposals need to be implemented by Bioversity

- Bioversity also responsible for Giga and thus GeneSys
 - will not invest in two competing proposals
- initiative to inform each other about the plans and products
 - Michael created a possibility for Fawzy to visit Theo
 - Frank, Siegfried and CGN staff joined



result: "Note on creating synergy between GeneSys and EURISCO"

- authors: Theo (EURISCO) and Fawzy (GeneSys)
- objective: to analyze both approaches and find overlap and possibilities for synergism

• inventory of required adaptations to become compatible



observations

- GeneSys is database with interface based on Grin & ICARDA data sets
 - technical solutions for practical problems (such as one table per crop/trait/method combination)
 - crop based
 - pragmatic solutions (such as the trait categories)
- EURISCO is a proposal for a repository
 - community support
 - aimed at easiest possible upload
 - EURISCO based (thus no crop-concept)



observations

- GeneSys is poorly defined; there is no documentation with schema and data dictionary (with format rules, etc.)
- structural differences are non-significant
 - only the crop basis of GeneSys might cause problems
 - the format differences could not be inventoried



steps forward

- EURISCO and GeneSys have to make sure that they stay informed about each other - representation in each others planning meetings should be arranged
- GeneSys should develop its documentation to inform EURISCO and others about its structures, format rules, etc - once this is available, a detailed comparison of the two formats should be made and in case of noncompatibility solutions should be discussed and implemented



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steps forward

- GeneSys and EURISCO should select/compile and agree on a list of crops and their names to be used by both
 - should cover (close to) all species occurring in both systems
- GeneSys and EURISCO should jointly develop a system of standardized trait names
 - compatible with the trait categories in GeneSys
 - on existing ontologies (such as those of the Generation Challenge Programme, the Trait Ontology Consortium, etc.).



steps forward

- EURISCO should incorporate the crop concept into the taxonomy system it will develop
 - no part of the uploading format
- EURISCO should expand its upload format with a few optional fields that are essential for the proper use of GeneSys
 - add Methods.Unit & Methods.Options that allow automatic deand recoding of scores
 - information is already contained in the current element TRAIT.TRAIT_METHOD
 - loading them in proposed fields avoids 'manual' atomization



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steps forward

- GeneSys and EURISCO should both consider adding additional optional fields
 - GeneSys: DATASET.UPLOADERCODE and TRAIT.TRAIT_REMARKS
 - EURISCO: Metadata.Institute, Metadata.E_Date, Metadata.Location, Metadata.Alt, Methods.Range



conclusions

- given the obvious complementarity and relatively small differences between GeneSys and the EURISCO initiative it is advisable to let the two approaches converge
- it is hoped that the functionality of GeneSys, using the elements developed by EURISCO, will make the EURISCO activities in this field redundant
 - for the time being the systems are too different in scope and objectives to eliminate one of them

