

Rose

- Most important ornamental crop
- More than 25,000 varieties of modern rose (Cairns, 2000)
- More than 10,000 hybrid tea varieties
- Rose list 2006: more than 13,000 varieties in commercial trade
- Large collection of roses in "common knowledge"

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STMS	linkage group	No. of alleles in 23 varieties	No. of allele phenotypes	Selected for	Scoring quality
RhO517	1	5	14	gr/ht	1
RhEO506	2	12	19	gr/ht	1
RhD221	4	8	12	gr/ht	1
RhE2b	6	7	12	gr/ht	1
RhB303	unknown	6	14	gr/ht	1
RhP519	unknown	7	15	gr/ht	1
RhAB40	4	11	18	gr/ht	1
RhD201	unknown	7	10	gr/ht	1
RhAB22	6	12	15	gr/ht	1
RhP50	3	11	13	Gr	1
RhP518	5	7	15	Gr	1
RhAB73	7	9	18	Gr	1
RhM405	unknown	5	13	Ht	1
RhAB15	2	10	5	Ht	1
RhO507	4	14	18	Ht	1



Standardisation between laboratories Proved to be very difficult, poor DNA quality from garden roses Often weak amplification Differences in signal intensity resulted in scoring of a peak in one lab as a marker and not scoring the same peak in the other lab Missing values Mis-scoring of alleles Marker data from one lab only

Database content 400 varieties included 314 varieties on behalf of the CPVO Morphological data available for all At least one photograph for 215 varieties 193 single pictures and 184 composite pictures Molecular profiles for 364 varieties.



Evaluation of the database (1)

Morphological descriptors

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- are useful, although usefulness varies with rose type
- greenhouse cut flower roses, most varieties currently fall into the same flower and plant growth type
- Flower color group frequently (38% of the case in greenhouse roses) wrongly indicated on TQ
- A ring test will be useful to ensure continued consistency of scoring

Evaluation of the database (2)

Photographs

- Very important
- For greenhouse roses the composite photo is not so informative as there is very little variation in the extra characteristics photographed
- For the garden varieties the composite adds very useful information
- Point for consideration: cost involved

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Evaluation of the database (3)

Molecular data

- Seedling varieties show unique patterns
- Mutants and mutant groups show the same molecular profile
- For two pairs of varieties there is still uncertainty about their nature
- Useful for spotting mutants
- More effort is needed to harmonize the molecular marker analysis between different laboratories

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Use of the database

- Characterization and cataloging of the reference collection
- Pre-screening and selection of appropriate reference varieties
- Exchange of data on current candidate varieties between testing stations
- To reduce permanent living reference collections at testing stations
- Quality assurance within examination offices (verification of identity/authenticity)

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Database and molecular profiles based on material submitted for DUS testing

- Identification label for tracing infringements
- Evidence in EDV cases

Advantage for breeders

Better possibilities to enforce rights

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