



BMT/15/21

ORIGINAL: English

DATE: May 20, 2016

INTERNATIONAL UNION FOR THE PROTECTION OF NEW VARIETIES OF PLANTS

Geneva

**WORKING GROUP ON BIOCHEMICAL AND MOLECULAR
TECHNIQUES AND DNA PROFILING IN PARTICULAR**

Fifteenth Session

Moscow, Russian Federation, May 24 to 27, 2016

EFFICIENT DUS TEST IN FRENCH BEAN BY USING MOLECULAR DATA

Document prepared by experts from the Netherlands

Disclaimer: this document does not represent UPOV policies or guidance

The Annex to this document contains a copy of a presentation “Efficient DUS test in French bean by using molecular data” to be made at its fifteenth session of the Working Group on Biochemical and Molecular Techniques and DNA-Profiling in particular (BMT).

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[Annex follows]



Efficient DUS test in French bean by using molecular data

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Efficient DUS test in French bean (1)

- Many varieties in same group (TG/12/9 Rev. 2): dwarf, white flower, round, green pod without string, white seed, resistant to BCMNV. And many of them also resistant to *Colletotrichum* and to *Pseudomonas*.

The following have been agreed as useful grouping characteristics:

- (a) Plant: growth type (characteristic 3)
- (b) Flower: color of standard (characteristic 16)
- (c) Pod: shape in cross section (through seed) (characteristic 22)
- (d) Pod: ground color (characteristic 24)
- (e) Pod: stringiness of ventral suture (characteristic 29)
- (f) Seed: number of colors (characteristic 43)
- (g) Seed: main color (largest area) (characteristic 44)
- (h) Seed: secondary color (characteristic 45)
- (i) Resistance to *Bean common mosaic necrosis virus* (BCMNV) (characteristic 50)

- In total 209 varieties known in this group, yearly 6 to 12 new applications at Naktuinbouw.

3

Efficient DUS test in French bean (2)

- Information on other characteristics, as stated in the (national) TQ, is being used for a careful selection of reference varieties for the field trial

- Leaf: green color
- Flower: size of bracts
- Pod: length
- Pod: width
- Pod: intensity of ground color
- Seed: weight

G	2	Plant: growth type	1 dwarf/2 climbing	{ }
	6	Leaf: intensity of green colour	1 very light/3 light/5 medium/7 dark/9 very dark	{ }
	14	Flower: size of bract	1 very small/3 small/5 medium/7 large/9 very large	{ }
G	15	Flower: colour of standard	1 white/2 pinkish white/3 pink/4 violet	{ }
	17	Pod: length (excluding beak)	1 very short/3 short/5 medium/7 long/9 very long (..... cm)	{ }
	18	Pod: width at maximum point	1 very narrow/3 narrow/5 medium/7 broad/9 very broad (..... mm)	{ }
G	21	Pod: shape of cross section (through seed)	1 narrow elliptic/2 elliptic to ovate/3 cordate/4 circular/5 eight shaped	{ }

part of NL TQ

- Information in TQ not always complete and/or accurate: e.g.
 - very dark green leaves (9) and pods 14,5 cm in DUS test,
 - dark green leaves (7) and pods 12-13 cm in TQ

4

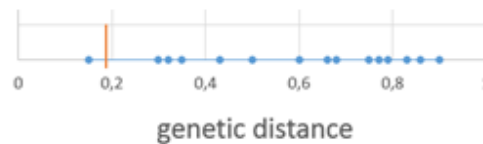
Efficient DUS test in French bean (3)

- Based on grouping characteristics and careful use of other information in TQ **15 to 20** reference varieties selected per application.
 - Expensive (**2 – 3 hours** per application for an expensive DUS expert)
 - Too many to have a good side by side comparison
 - Risk of mistakes in selection due to inaccurate information on TQ.
 - In case of mistakes (2015: 3 cases on 12 new applications) again check on reference varieties, but now based on own, complete description. Risk on 3 years of testing.

5

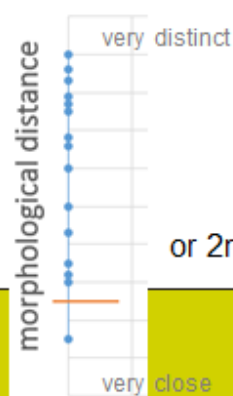
Theory towards more efficiency

- Year 1 test 1



- Year 1 test 2
 - Genetically similar varieties in field
 - Other varieties check on paper

- Conclusion on D after 1 year



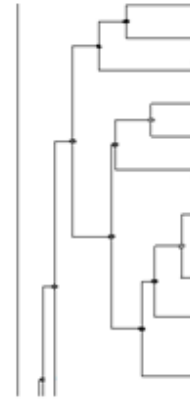
or 2nd year with 'paper' varieties

6



Molecular data

- AFLP database for many French bean varieties available at Naktuinbouw
- 4 primer combinations (approximately 500 bands/loci) 78 bands are polymorph in the database. 230 varieties (528 entries in the database).
- No database yet with SSR or SNP, no whole genome sequencing.
- Dendrogram with genetic distances for 230 varieties



7



Accepted UPOV models

Combining phenotypic and molecular distances in the management of variety collections

Calibrated molecular distances in the management of variety collections

- Not in all crops good correlation
- Large data set needed
- Per crop large preparation phase

Theory as proposed in bean seems not to fit in these models.

8

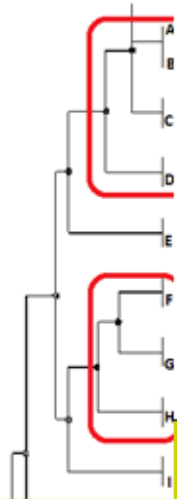
Example French bean (1)

Test 1- year 1:

- seedling check on DNA
- compare DNA pattern with dendrogram
- Set threshold
- decide on 1 to 5 genetically most similar varieties

Example:

- Application A to compare with B, C and D
- Application F to compare with G and H



9

Example French bean (2)

Test 2 – year 1:

- Field trial of the application with as reference varieties:
 - the 1 to 5 genetically most similar varieties
 - the similar variety, variety indicated by the breeder on the TQ, unless this similar variety is in a different group (example different resistance)
- First year of DUS trial with only 1 to 6 reference varieties, instead of 15 to 20: good side by side comparison possible.
- Conclusion of year 1: compare own complete description with all descriptions in database

10



Example French bean (3)

As the description of the application is complete and all descriptions are made by the examination office itself, one can be strict in selecting: not coming to 15 to 20 reference varieties, but none or only a few in a short time (**less than 30 minutes**).

- Clearly distinct in year 1 and (based on check of morphological data in database) no extra reference varieties needed in year 2: 1 year of testing is sufficient to declare the variety Distinct.



Example French bean (4)

Year 2:

- Field trial with
 - one or no similar reference variety (of the 1 to 6) of year 1
 - other reference varieties selected from the database based on own, reliable description made in year 1
- Conclusion on Distinctness

Results in French bean 2015

Application	2 step example French bean			Traditional	
	Number of genetically similar varieties	Number of references in year 1	Number of references to be added in year 2 (similar on paper)	Total number of references in 2 years trials	Total number of references in 2 years trials
A	3	3	5	8	21
B	3	4	7	11	14
C	1	1	2	3	6
D	2	3	1	4	5
E	1	2	1	3	12
F	1	3	0 D year 1	3	25
G	1	4	2	6	13
H	5	7	1	8	15
I	4	5	0	5	13
TOTAL				51	124

13

Implementation

Needed before implementation is possible:

- Availability of own morphological data in database of 'all' varieties in common knowledge
- Availability of database with molecular data of 'all' varieties in common knowledge, based on a well defined and robust marker system. High resolution and validated.
- Validated threshold for similar varieties to put in the field.
- Availability of seed samples of the applications prox. 3 weeks before preparation of the field trial

14



Costs and benefits, example French bean

- + less trial plots ($51/124 = 41\%$ compared to traditional method)
- + less visual observations to make
- + better quality of side by side comparison
- + less hours for the process of selecting reference varieties for the growing trial (at least 1,5 h less)
- +++ when TQ is not very informative (only information on grouping characteristics)

- costs for DNA test (costs will decrease importantly)
- 3 years of test not wanted, but what to do if a 'paper' reference appears to be very close in year 2 and genetical similarity is low?
- Submission of seed should be 3 weeks earlier