**Third Session** 

Technical Working Party on Testing Methods and Techniques	TWM/3/4
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Beijing, China, April 28 to May 1, 2025

Original: English Date: March 26, 2025

### COYD-GP ENHANCED DISTINCTNESS CRITERION FOR CROSS-POLLINATED AGRICULTURAL CROPS

Document prepared by an expert from the United Kingdom

Disclaimer: this document does not represent UPOV policies or guidance

The annex to this document contains a copy of a presentation "COYD-GP Enhanced distinctness criterion for cross-pollinated agricultural crops", to be made by an expert from the United Kingdom, at the third session of the TWM.

[Annex follows]

TWM/3/4

### ANNEX





Concern I the new v • Ci • Ci • Ci	y some that in some cas arieties ops like perennial ryegra Large reference collections Population varieties Many measured characteristics v not held by all	ies distinctness is hard	d to achieve despite i	mproved performance of





an Horizon 2020 funded project INVITE gave BioSS the opportunity to look at reference collection
ment using markers and genomic prediction
Included historical DUS field data for perennial ryegrass from Naktuinbouw
TWM/2/4: Reference collection management using molecular markers: a new approach based on genomic prediction
https://www.upov.int/edocs/mdocs/upov/en/twm_2/twm_2_4.pdf
this study, BioSS was asked by a number of parties if we were looking at distinctness using
) Net an ariginal aim of the project
Not an original aim of the project



improve estimates of variety means from the 2 or 3 years of trials rkers help to improve estimates
ng to select breeding material f key traits, such as yield d variable netic data can give a better "prediction" of the trait tter understand the trait type, not genotype

<ul> <li>comparing two varieties</li> <li>This is generally applied when the same varieties are compared over two or three cycles</li> <li>Analysis of variance can be seen as a linear mixed model</li> </ul>	
COYD-GP	
<ul> <li>In COYD-GP, variety is treated as a random effect, with correlations between effect levels</li> <li>Correlations come from a genetic relationship matrix (kinship matrix) calculated from the gen markers → gBLUP</li> </ul>	genetic
<ul> <li>Everything else is the same</li> <li>Differences in variety means compared to a measure of precision (t-test)</li> </ul>	

Initial study based on Perennial Ryegrass DUS data from Naktuinbouw. Netherlands	
• Up to 13 years of trials	
21 characteristics	
119 diploids and 149 tetraploids	
200k SNPs	
<ul> <li>Applied long-term versions of COYD and COY-GP, using the whole data set.</li> <li>Counted number of differences with probability value 1%.</li> </ul>	
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<ul> <li>Applied long-term versions of COYD and COY-GP, using the whole data set.</li> <li>Counted number of differences with probability value 1%.</li> <li>Note:         <ul> <li>It would be better to look at 3 year cycles as normal practice</li> <li>But this was an initial evaluation, and the data set had fewer varieties</li> </ul> </li> </ul>	

	Diploid		Tetraploid	
Ryegrass DUS Characteristic	Distinct with	Increase	Distinct with	Increase with
	COYD	with COYD-	COYD	COYD-GP
		GP		
Plant: vegetative growth habit (without vernalization)	28.8%	1.8%	10.1%	1.3%
Leaf: intensity of green colour (without vernalization)	7.7%	2.2%	2.7%	0.5%
Plant: width (after vernalization)	25.3%	2.0%	15.3%	1.7%
Plant: vegetative growth habit (after vernalization)	37.0%	2.6%	21.2%	6.6%
Plant: height (after vernalization)	46.8%	2.0%	24.2%	5.9%
Leaf: intensity of green colour (after vernalization)	17.0%	1.5%	11.9%	4.8%
Plant: time of inflorescence emergence	77.5%	0.3%	70.0%	1.0%
Plant: natural height at inflorescence emergence	39.0%	2.2%	25.3%	4.6%
Plant: growth habit at inflorescence emergence	15.7%	4.1%	22.0%	9.8%
Flaf leaf: length	13.5%	2.2%	9.8%	3.5%
Flag leaf: width	39.8%	2.2%	28.0%	5.0%
Flag leaf: length/ width ratio	26.7%	2.9%	9.1%	3.4%
Plant: length of longest stem, inflorescence included (when fully expanded)	43.3%	2.9%	31.5%	5.5%
Plant: length of upper internode	17.6%	3.3%	6.4%	1.8%
Inflorescence: length	28.0%	0.8%	27.9%	3.4%
Inflorescence: number of spikelets	37.5%	1.3%	20.0%	2.5%
Inflorescence: density	29.7%	1.4%	29.4%	3.2%
Inflorescence: length of outer glume on basal spikelet	26.2%	1.5%	16.4%	3.5%
Inflorescence: length of basal spikelet excluding awn	22.2%	2.9%	16.8%	4.1%
Inflorescence: spikelet protuberance	22.4%	3.4%	15.7%	5.4%
Inflorescence: glume span	30.3%	3.0%	14.4%	2.9%

	Diploid		Tetraploid	
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# Bacts Ba

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Test in relevant crops	• (	omplete DUS data	
	• T	est in relevant crops	
How many markers needed	• F	ow many markers needed	
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