

**Work plan for UPOV Technical Working Party on Testing Methods and Techniques (TWM), Third Session,
Beijing, China (TWM/3)**

(please note that the schedule is subject to change at any time) – Local time in Beijing (UTC+8)

(UTC+8)	Monday, April 28 Start 08.30	Tuesday, April 29 Start 8.30	Wednesday, April 30 Start 8.30	Thursday, May 1 Start 8.30
8.30 am	<p><u>1. Opening</u></p> <p>(i) Opening remarks [MARA]</p> <p>(ii) Opening remarks [NFGA]</p> <p>(iii) Overviews of China's PVP system [MARA]</p> <p>(iv) Overviews of China's PVP system [NFGA]</p> <p><u>2. Adoption of agenda (TWM/3/1 Rev.)</u></p> <p><u>5. Date and place next session</u></p> <p><u>4. Matters for information</u></p> <p>(a) Reports on developments in UPOV</p> <p>(b) Reports from members and observers (TWM/3/2)</p> <p>(c) Procedures for DUS examination (TWP/9/1)</p> <p>(d) UPOV Information databases (TWP/9/2)</p> <p>(e) Test Guidelines: support for drafters; additional characteristics; and methods of propagating the variety (TWP/9/3)</p> <p>(f) Proposal for a revision of document TGP/7 "Development of Test Guidelines", GN 28 "Example Varieties" (TWP/9/5)</p>	<p>FIELD VISIT</p> <p>Departure from hotel: 8:30</p> <p>Return to hotel: 19:30</p>	<p><u>3.3 (b) Cooperation between international organizations (Joint OECD, ISTA and UPOV workshop on molecular techniques)</u></p> <p>(i) Developments at ISTA (TWM/3/25)</p> <p>(ii) Development at OECD (TWM/3/26)</p> <p>(iii) Developments at UPOV</p>	<p><u>3.3 (g) The use of molecular techniques for enforcement</u></p> <p>(i) Use of DNA techniques for plant variety rights enforcement in Peru (TWM/3/3)</p> <p>(ii) Use of Molecular Markers as a tool to enforce Plant Breeders' Rights (PBR) in Soybean in Uruguay (TWM/3/18)</p> <p><u>3.3 (d) Methods for analysis of molecular data, management of databases and exchange of data and material (cont'd)</u></p> <p>(v) Use of DNA databases at Naktuinbouw to improve DUS work (TWM/3/8)</p> <p>(vi) Shared molecular database (TWM/3/23)</p> <p><u>6. Future program</u></p>
10.00	Coffee Break		Coffee Break	Coffee Break
10.30	<p><u>3.1 Software and statistical analysis methods for DUS examination</u></p> <p>(i) Development of big data platform for DUS examination (TWM/3/19)</p> <p>(ii) Grading criteria of Anthurium DUS quantitative characteristics by multiple comparison (TWM/3/12)</p> <p><u>3.2 Phenotyping and image analysis</u></p> <p>(i) A new perspective on the DUS test of eggplant fruit color based on lab color parameters (TWM/3/13)</p> <p>(ii) Length data collection device pro (TWM/3/14)</p>		<p><u>3.3 (e) Confidentiality, ownership and access to molecular data, including model agreement template</u></p> <p>- Confidentiality of molecular information (TWP/9/6)</p> <p><u>3.3 (f) The use of molecular techniques in examining essential derivation</u></p> <p>(i) Exploration of identification techniques based on SNP markers for essentially derived varieties of wheat (TWM/3/11)</p> <p>(ii) Essentially derived varieties (EDV) threshold development in soybeans (TWM/3/9)</p>	
12.00 pm	Lunch		Lunch	Lunch
14.00	<p><u>3.1 Software and statistical analysis methods for DUS examination (cont'd)</u></p> <p>(iii) COYU development update 2025 (TWM/3/5)</p> <p><u>3.3 (a) Latest developments in molecular techniques and bioinformatics</u></p> <p>(i) Data science activities at Naktuinbouw towards genotyping and phenotyping: an update (TWM/3/16)</p> <p><u>3.3 (d) Methods for analysis of molecular data, management of databases and exchange of data and material</u></p> <p>(i) Exploiting crop haplotype-tag polymorphisms marker for pedigree identification (TWM/3/10)</p> <p>(ii) PAD – an algorithm for progeny-ancestor detection based on genetic profiles (TWM/3/17)</p>		<p><u>3.3 (c) Report of work on molecular techniques in relation to DUS examination</u></p> <p>(i) Guidelines for the validation of a new characteristic-specific molecular marker protocol as an alternative method for observation (TWP/9/4)</p> <p>(ii) Latest developments in characteristic-specific molecular markers at Naktuinbouw: a call for knowledge exchange (TWM/3/7)</p> <p>(iii) The use of biomolecular technology in DUS testing - a case study on barley (TWM/3/20)</p> <p>(iv) Artificial Intelligence and molecular markers in soft fruit: a proof of concept (TWM/3/24)</p>	<p><u>7. Adoption of the Report</u></p> <p><u>8. Closing of the session</u></p>
15.30	Coffee Break		Coffee Break	Coffee Break
16.00	<p><u>3.3 (d) Methods for analysis of molecular data, management of databases and exchange of data and material(cont'd)</u></p> <p>(iii) DurdusTools: Current state and use in DUS-testing (TWM/3/21)</p> <p>(iv) Development of DUS phenotyping tools for and with examination offices: experience gained (TWM/3/27)</p> <p>(v) Phenotyping concept for strengthening the plant variety protection chain via combined use of IA&AI (TWM/3/28)</p>		<p><u>3.3 (c) Report of work on molecular techniques in relation to DUS examination(cont'd)</u></p> <p>(v) Can better understanding of the genetic architecture of wheat DUS characteristics help streamline the DUS processes? (TWM/3/22)</p> <p>(vi) Genomic prediction for variety collection management in wheat (TWM/3/6)</p> <p>(vii) COYD-GP enhanced distinctness criterion for cross-pollinated agricultural crops (TWM/3/4)</p> <p>(viii) CPVO R&D activities (TWM/3/15)</p>	
17.30	End		End	End
	Reception Dinner by MARA (Qin Yuan Fu Hotel)			