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| International Union for the Protection of New Varieties of Plants |  |

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| Administrative and Legal Committee  Seventy-Ninth Session  Geneva, October 26, 2022 | CAJ/79/INF/4  Original: English  Date: October 15, 2022 |

Molecular techniques

Document prepared by the Office of the Union

Disclaimer: this document does not represent UPOV policies or guidance

# Executive summary

The purpose of this document is to report on developments concerning molecular techniques since the

seventy-eighth session of the Administrative and Legal Committee (CAJ).

The structure of this document is as follows:

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ANNEX I SURVEY ON THE USE OF MOLECULAR MARKER TECHNIQUES BY CROP

ANNEX II ELEMENTS FOR DRAFT JOINT DOCUMENT EXPLAINING THE PRINCIPAL FEATURES OF THE SYSTEMS OF THE OECD, UPOV AND ISTA

The following abbreviations are used in this document:

BMT: Working Group on Biochemical and Molecular Techniques, and DNA-Profiling in Particular

ISTA: International Seed Testing Association

OECD: Organization for Economic Co-operation and Development

TC: Technical Committee

TWA: Technical Working Party for Agricultural Crops

TWF: Technical Working Party on Fruit Crops

TWM: Technical Working Party on Testing Methods and Techniques

TWO: Technical Working Party on Ornamental Plants and Forest Trees

TWPs: Technical Working Parties

TWV: Technical Working Party for Vegetables

BACKGROUND

Developments on matters presented in this document at the TC, at its fifty-eighth session, will be reported to the CAJ in document CAJ/79/2 “Report on developments in the Technical Committee”.

# Cooperation between international organizations

## Background

The background to this matter is provided in document CAJ/78/INF/5 “Molecular Techniques”, paragraphs 8 to 31.

Developments concerning this matter since the seventy-eighth session of the CAJ are provided in the following paragraphs.

## Inventory on the use of molecular marker techniques, by crop

The TC, at its fifty-seventh session[[1]](#footnote-2), requested the Office of Union to inform OECD of the result of the survey, which were presented in document TC/57/8 “Molecular Techniques”, and to report on the developments at the TC, at its fifty-eighth session (see document TC/57/25 “Report”, paragraph 48). On December 13, 2021, the Office of the Union informed the results of the survey to OECD.

The TC agreed, at its fifty-seventh session, to continue the survey on the use of molecular markers to obtain information from a greater number of members and to investigate the reasons for members not responding to the first survey.

On February 1, 2022, the Office of the Union issued Circular E-22/009 inviting members of the Union to indicate whether they use molecular marker techniques and continue the survey on their use.

In response to the Circular E-22/009, 28 members of the Union replied, 15 of which were new responses and 8 provided further information on the use of molecular marker techniques. The summary of responses to the surveys in 2020 and 2022 are as follows:

|  |  |
| --- | --- |
| Member | Use of Molecular marker technique (YES/NO) |
| Argentina | YES |
| Australia | NO |
| Austria | YES |
| Belgium | NO |
| Bolivia | NO |
| Brazil | YES |
| Canada | YES |
| China | YES |
| Czech Republic | YES |
| European Union | YES |
| Estonia | YES |
| France | YES |
| Germany | YES |
| Hungary | YES |
| Ireland | YES |
| Israel | NO |
| Italy | YES |
| Japan | YES |
| Jordan | YES |
| Kenya | NO |
| Latvia | NO |
| Member | Use of Molecular marker technique (YES/NO) |
| Lithuania | NO |
| Mexico | NO |
| Moldova | NO |
| Netherlands | YES |
| New Zealand | NO |
| Norway | NO |
| Panama | NO |
| Peru | NO |
| Poland | NO |
| Romania | NO |
| Russian Federation | NO |
| Slovakia | YES |
| Spain | YES |
| Ukraine | YES |
| United Kingdom | YES |
| United States of America | NO |
| Zimbabwe | NO |
| YES | 20 |
| NO | 18 |
| TOTAL | 38 |

The detailed results of the survey are presented in Annex I to this document.

## Joint document explaining the principal features of the systems of OECD, UPOV and ISTA

The TC, at its fifty‑seventh session, noted that it had approved by correspondence the draft joint document explaining the principal features of the systems of OECD, UPOV and ISTA. The TC requested the Office of the Union to inform OECD and ISTA accordingly (see document TC/57/25 “Report”, paragraph 52). The elements of a draft joint document are provided in Annex II to this document.

On December 13, 2021, the Office of the Union transmitted the draft joint document to OECD and ISTA. Responses from OECD and ISTA, when available, will be reported to the TC and the CAJ.

## Lists of possible joint initiatives with OECD and ISTA in relation to molecular techniques

The TC, at its fifty-seventh session, agreed to propose the following topics for a future joint UPOV/OECD/ISTA workshop:

(i) providing information on the use of molecular techniques in each organization;

(ii) procedure for approval of biochemical and molecular methods in each organization; and

(iii) possibilities for harmonizing terms, definitions and methods between UPOV, OECD and ISTA.

The TC agreed to request the Office of the Union to contact OECD and ISTA to explore suitable dates for a future joint workshop, such as in conjunction with the first session on the TWM, to be held in September 2022.

On December 13, 2021, the Office of the Union communicated an invitation to OECD and ISTA to hold another joint workshop on molecular techniques, in conjunction with the first session of the TWM, to be held on September 21, 2022, via electronic means, to include consideration of the topics (i) to (iii) above.

On May 15, 2022, the Office of the Union met with OECD and ISTA at the fringes of the International Seed Federation World Seed Congress and discussed cooperation between OECD, ISTA and UPOV in relation to BMT. UPOV had raised the idea of a joint OECD/ISTA/UPOV workshop in conjunction with the TWM, which would be held in September 2022. It was agreed that there were no significant developments to discuss at this time and that it might be better to plan a joint workshop in 2023 at the earliest.

## International Seed Testing Association

The TWM received a presentation from Ms. Ana Laura Vicario (ISTA) on “ISTA report on the use of molecular techniques”, a copy of which is reproduced in document TWM/1/23.

## Organisation for Economic Co-operation and Development

The TWM received a presentation from Mr. Christophe Rouillard (OECD) on “Latest developments in the application of BMT under the OECD Seed Schemes”, a copy of which is reproduced in document TWM/1/24.

The TWM noted that OECD had formed the Advisory Group on BMT to deal with all BMT issues under the Seed Schemes, including matters relating to cooperation with other international organizations.

# Session to facilitate cooperation in relation to the use of molecular techniques

The background to this matter is provided in document CAJ/78/INF/5 “Molecular Techniques”, paragraphs 32 to 44.

## Technical Working Parties (TWPs)

The TWA, at its fifty-first session[[2]](#footnote-3), held a discussion session to allow participants to exchange information on their work on biochemical and molecular techniques and explore possible areas for cooperation. The TWA considered whether UPOV could support harmonization and cooperation between members already using molecular markers in DUS examination or making information or BMT services available to other UPOV members.

The TWA agreed that the TWPs were a platform for exchanging information about molecular markers in DUS examination, including projects, collaborations and services eventually provided by members. The TWA agreed that UPOV should continue to encourage presentations on using molecular markers in DUS examination, including technical aspects, confidentiality and access to data.

The TWA received a presentation on the “Use of molecular techniques in DUS examination: Report from Argentina” by an expert from Argentina. A copy of the presentation is provided in document TWA/51/4.

The TWA received a presentation on “Developing a strategy to apply SNP molecular markers in the framework of winter oilseed rape DUS testing” by an expert from France. A copy of the presentation is provided in document TWA/51/4 Add.

The TWF, at its fifty-third session[[3]](#footnote-4), received a presentation on “Application of molecular techniques in DUS testing and PBR enforcement of fruit sector in China” by an expert from China. A copy of the presentation is provided in document TWF/53/12.

The TWF noted that molecular markers could be used in China as first instance evidence for enforcement of breeders’ rights, followed by a growing trial in case required.

Following the presentation from China, the TWF had an open discussion about the use of molecular markers in DUS examination and variety identification. The following aspects were mentioned by participants:

* Possibilities for cooperation on the constitution of common databases, including for authorities receiving relatively few applications for particular crops
* Origin of plant material for DNA extraction (e.g. material provided for DUS testing)
* Selection of markers for each crop, according to intended use (e.g. for PBR and/or variety identification).
* Selecting one or more laboratories capable of providing high-quality molecular profiles (e.g. security back-up);
* High cost for harmonizing methodologies for DNA profiling among different laboratories;
* Difficulties to obtain the same results even for laboratories using harmonized methodologies.

The TWO, at its fifty-fourth session[[4]](#footnote-5), noted that no reports were made on the use of biochemical and molecular techniques in DUS examination of ornamental plants.

The TWM, at its first session[[5]](#footnote-6), held a discussion session to allow participants to exchange information on their work on biochemical and molecular techniques and explore possible areas for cooperation.

The TWM agreed that sufficient time should be allocated for discussing the agenda items in the future work plan for the meeting and agreed that it would not be necessary to have an open discussion session.

# Confidentiality AND Ownership of Molecular Information

The background to this matter is provided in document CAJ/78/INF/5 “Molecular Techniques”, paragraphs 45 to 55.

The TWPs and the BMT, at their sessions in 2021, received a presentation on “Confidentiality & Ownership of Molecular Information” by an expert on behalf of the African Seed Trade Association (AFSTA), the Asia and Pacific Seed Association (APSA), the International Community of Breeders of Asexually Reproduced Horticultural Plants (CIOPORA), CropLife International, Euroseeds, the International Seed Federation (ISF) and the Seed Association of the Americas (SAA). The TWPs and the BMT considered the proposal to revise document TGP/5, Section 3: Model Application Form, to include a request for confidentiality of molecular information of candidate varieties as follows:

*“I/We request that molecular information pertaining to the variety remains confidential and exchange to another UPOV member or examination office is subject to approval by the applicant.”*

At their sessions in 2022, the TWV[[6]](#footnote-7), TWA[[7]](#footnote-8), TWO[[8]](#footnote-9), TWF[[9]](#footnote-10) and TWM[[10]](#footnote-11), noted discussions held at the TWPs and the BMT, at their sessions in 2021, on “Confidentiality & Ownership of Molecular Information”.

The TWA, at its fifty-first session, noted the report from the joint breeders’ organizations that a survey on confidentiality of molecular data was being conducted among plant breeding companies across different organizations. The TWA noted that the outcomes of the survey would be presented to the TWM, at its first session. The TWA agreed to invite the joint breeders’ organizations to report developments at its fifty-second session.

The TWM, at its first session, received a presentation from Mr. Marcel Bruins (CropLife International) on “Confidentiality and ownership of molecular information”, a copy of which is reproduced in document TWM/1/22.

The TWM noted the concern expressed by breeders’ organizations that molecular information provided for the examination of a variety should not be provided to others outside the authority that received the application without the permission of the breeder. The TWM further noted the concern that breeders were lacking clarity and information about how molecular information was being used, and especially shared.

The TWM agreed that further clarification was required on the type of information and purpose of use of data to be shared (e.g. molecular distances between varieties, genotype sequences) that would require permission before being shared by PVP offices with other PVP offices.

The TWM agreed to invite members and observers to report existing policies on confidentiality of molecular information at the second session of the TWM.

[Annexes follow]

SURVEY ON THE USE OF MOLECULAR MARKER TECHNIQUES BY CROP

*Please see the Excel spreadsheet for all the responses received*

[Appendix to Annex I follows]

Response from the European Union:

USE OF MOLECULAR MARKER TECHNIQUES FOR DUS TESTING IN THE FRAMEWORK OF CPVR

Elements of context

The Technical Committee (TC) of UPOV, at its fifty-fifth session, held in Geneva, on October 28 and 29, 2019, agreed to invite members of the Union to complete a survey as a basis to develop an inventory on the use of molecular marker techniques, by crop, in coordination with the OECD Seed Schemes (see document TC/55/25 “Report”, paragraphs 184 and 185).

The information on molecular marker techniques used by members of the Union will be used to develop a joint UPOV/OECD/ISTA document containing that information, in a similar format to document UPOV/INF/16 “Exchangeable Software”, subject to the approval of the Council and in coordination with OECD and ISTA.

The present document constitutes the contribution of the CPVO to this survey and, as such, describes the molecular marker techniques that can be used for DUS testing aiming at the granting of Community Plant Variety Rights (CPVR) in the framework of the CPVO policy.

1. Legal framework on the use of molecular techniques in DUS testing for CPVR

The legal framework for conducting DUS testing for CPVR includes the CPVO Basic Regulation 2100/94 (BR) and its Implementing Rules, the CPVO Technical Protocols (TPs) and the guidance documents adopted by UPOV.

The CPVO does not undertake DUS testing himself but, as stated in Article 56 (BR) “shall arrange for the technical examination [...] to be carried out by the competent office or offices in at least one of the Member States entrusted with responsibility for the technical examination of varieties of the species concerned by the Administrative Council”.

When carrying out a technical examination, “[…] the Examination Offices shall, for the purposes of the technical examination, grow the variety or undertake any other investigations” (Article 56-BR). Molecular techniques may thus be used to support DUS testing by the entrusted EOs provided that the technical examination is conducted in accordance with the test guidelines issued by the CPVO Administrative Council.

As the same guidelines are used both for variety protection and for registration in the EU, the Directives on the Common Catalogues (Council Directives 2002/53/EC and 2002/55/EC) are also to be taken into account. According to them, the acceptance of varieties shall be based on the results of official examinations, particularly growing trials, covering a sufficient number of characteristics for the variety to be described. Therefore, molecular techniques may be used only as complementary tools in addition to the growing trials.

As a UPOV member, the CPVO respects the agreed framework on the use of molecular techniques in DUS testing as laid down in documents UPOV/INF/18 (adopted by the Council of UPOV in 2011) and UPOV/TGP/15/3 (adopted by the Council of UPOV in 2020). More specifically, the CPVO supports the application by the network of its entrusted EOs of molecular tools according to the models positively assessed as regards their conformity with the UPOV convention.

1. Models supported by the CPVO and examples of application
   1. Characteristic-specific markers

Molecular markers can be used as an alternative to the phenotypic observation, as predictors of traditional characteristics that are difficult or cumbersome to assess, if a clear link exists. They can be either fully or partly correlated to the phenotype. These methods are included in technical protocols of CPVO on the basis of an evaluation/validation and suggestion of the CPVO crop-expert groups.

* + 1. Markers 100% correlated to a given state of expression of the characteristic

In that case, the marker may replace the phenotypic observation.

Examples of characteristics concerned:

- Resistances to mono- or oligogenic diseases resistances (e.g. diseases in vegetables, resistance to nematode *Heterodera schachtii* in sugarbeet)

- CMS (cytoplasmic male sterility) in cabbages

- Herbicides (e.g. sunflower, rapeseed)

So far, none of these markers have been included in CPVO TPs.

* + 1. Markers providing incomplete information on the state of expression of the characteristic

In that case, the marker is only partially linked to the characteristic and give an incomplete information on the level of expression of the trait. Its use has to be described in an assessment scheme that precise the situations where it can be used and when it needs to be completed by a phenotypical observation.

Examples of characteristics concerned: quantitative diseases resistances in vegetables, such as

- Tomato mosaic virus (ToMV)

- Tomato spotted wilt virus (TSWV)

The two sets of co-dominant markers developed for these two tomato viruses have been included in the CPVO TPs for tomato (4.4-2) and tomato rootstocks (1.4) as a potential alternative to the biotests in specific cases.

* 1. Management of variety collections
     1. Combining molecular and phenotypic thresholds to exclude super-distinct varieties from the second growing trial

In this model, two independent thresholds are set for the selection of similar varieties to be included into the growing trial. The first threshold is based on the information of morphological characteristics and the second relies on a genetic distance calculated using a set of markers distributed throughout the genome. Except for morphologically very similar varieties, reference varieties exceeding the two thresholds do not need to be included into the growing trial (they are considered as “super-distinct”).

This model is routinely applied by certain entrusted EOs for species like maize, lettuce, wheat and barley, and is currently being tested for oilseed rape through R&D projects co-financed by CPVO.

* + 1. Genetic selection of similar varieties for the first growing trial

The candidate variety is genotyped using a defined set of markers, and its profile is compared to the varieties from the reference collection. All reference varieties with a genetic similarity to the candidate higher than a certain percentage (e.g. 80%) are to be included in the first growing cycle, all the others being excluded.

During the first cycle, the candidate variety is assessed on uniformity and described morphologically according to the technical protocol. Its morphological description is then compared *in silico* to the descriptions of all the reference varieties.

*Remark: the morphological descriptions of the reference varieties used for the in silico comparison shall be based on observations made by the EO (in-house variety descriptions). If the variety descriptions used are not made in-house, they can be used only if notation scales have been harmonized between the examination offices producing and using them (through ring tests for example).*

The reference varieties identified to be morphologically similar to the candidate will be included for comparison into a second growing trial. If the variety is clearly distinct from the similar varieties in the first growing cycle and no similar varieties are detected based on the variety description after the first growing cycle, a positive decision on distinctness can be taken after one growing cycle.

This model is under application by certain entrusted EOs for species like French bean and potato. It is currently under test for durum wheat and will be explored for tomato and hemp through R&D projects co-financed by CPVO.

The methods are evaluated by CPVO crop expert groups

* 1. Other uses
     1. Identification in support to the maintenance of variety collections

All the molecular markers used in the above examples can be used for identification purposes in support to the maintenance of reference collections.

In addition, other molecular marker sets can be used also for identification purposes by certain entrusted EOs for species like rose, cherry tree, peach, grapevine, citrus… These sets can be harmonized between EOs (e.g. potato, thanks to a project supported by CPVO) or not.

* + 1. Detection of GMOs (in the sense of Directive 2001/18/EC)

In specific cases, markers are used by EOs to detect varieties produced with the help of transgenesis or targeted mutagenesis techniques for:

- Confirmation of the presence of a declared genetic transformation (classical transgene insertion, or point mutations triggered by gene editing technologies).

- Detection of adventitious presence of GM seeds in the submitted reference lots.

Conclusion

In summary, many molecular methods are presently being used, or under development, by the CPVO network of entrusted EOs in support of DUS testing.

However, only 2 sets of characteristic-specific molecular markers are officially described in CPVP TPs on the basis of evaluation/validation of experts in the CPVO expert groups.. These markers are publically available.

Consequently, the CPVO leaves to its entrusted EOs the role of describing the molecular tools they use in respect of the CPVO policy for the methods used in relation to variety reference collections.

[Annex II follows]

ELEMENTS FOR DRAFT JOINT DOCUMENT explaining the principal features of the systems of the OECD, UPOV and ISTA

The Organisation for Economic Co-operation and Development (OECD)

*What are the OECD Seed Schemes?*

The OECD Seed Schemes provide an international framework for the varietal certification of agricultural seed moving in international trade. The Schemes were established in 1958 driven by a combination of factors including a fast-growing seed trade, regulatory harmonisation in Europe, the development of off-season production, the seed breeding and production potential of large exporting countries in America (North and South) and Europe, and the support of private industry. Membership of the Schemes is voluntary and participation varies. There are seven agricultural Seed Schemes.

*Participating countries*

59 countries from Europe, North and South America, Africa, the Middle-East, Asia and Oceania currently participate in the OECD Seed Schemes:

|  |  |  |  |
| --- | --- | --- | --- |
| ALBANIA | (2) | LITHUANIA | (2) |
| ARGENTINA | (2) | LUXEMBOURG | (1) |
| AUSTRALIA | (1) | MEXICO | (1) |
| AUSTRIA | (1) | MOLDOVA | (2) |
| BELGIUM | (1) | MOROCCO | (2) |
| BOLIVIA | (2) | NETHERLANDS | (1) |
| BRAZIL | (2) | NEW ZEALAND | (1) |
| BULGARIA | (2) | NORWAY | (1) |
| CANADA | (1) | POLAND | (1) |
| CHILE | (1) | PORTUGAL | (1) |
| CROATIA | (2) | ROMANIA | (2) |
| CYPRUS1 | (2) | RUSSIAN FEDERATION | (2) |
| CZECH REPUBLIC | (1) | SENEGAL | (2) |
| DENMARK | (1) | SERBIA | (2) |
| EGYPT | (2) | SLOVAKIA | (1) |
| ESTONIA | (1) | SLOVENIA | (1) |
| FINLAND | (1) | SOUTH AFRICA | (2) |
| FRANCE | (1) | SPAIN | (1) |
| GERMANY | (1) | SWEDEN | (1) |

1 Source OECD “Note by Turkey

The information in this document with reference to ‘Cyprus’ relates to the southern part of the Island. There is no single authority representing both Turkish and Greek Cypriot people on the Island. Turkey recognises the Turkish Republic of Northern Cyprus (TRNC). Until a lasting and equitable solution is found within the context of the United Nations, Turkey shall preserve its position concerning the ‘Cyprus issue’.

Note by all the European Union Member States of the OECD and the European Union

The Republic of Cyprus is recognised by all members of the United Nations with the exception of Turkey. The information in this document relates to the area under the effective control of the Government of the Republic of Cyprus.”

|  |  |  |  |
| --- | --- | --- | --- |
| GREECE | (1) | SWITZERLAND | (1) |
| HUNGARY | (1) | TUNISIA | (2) |
| ICELAND | (1) | TURKEY | (1) |
| INDIA | (2) | UGANDA | (2) |
| IRAN | (2) | UKRAINE | (2) |
| IRELAND | (1) | UNITED KINGDOM | (1) |
| ISRAEL | (1) | UNITED STATES | (1) |
| ITALY | (1) | URUGUAY | (2) |
| JAPAN | (1) | ZIMBABWE | (2) |
| KENYA | (2) |  |  |
| KYRGYZSTAN | (2) | (1) OECD Member Country |  |
| LATVIA | (2) | (2) Non OECD Member Country | |

Figure 1 Map of Participating Countries in the OECD Seed Schemes (2016)



*Objectives*

The objectives of the Schemes are to encourage the production and use of “quality-guaranteed” seed in participating countries. The Schemes authorise the use of labels and certificates for seed produced and processed for international trade according to agreed principles ensuring varietal identity and purity.

The Schemes facilitate the import and export of seed, by the removal of technical barriers to trade by assuring identification and origin through internationally recognised labels (“passports”) for trade. They also lay down guidelines for seed multiplication abroad, as well as for the delegation of some control activities to the private sector (“authorisation”). The quantity of seed certified through the OECD Schemes has grown rapidly in recent years and now exceeds 1 million tonnes.

*How do the Seed Schemes operate*

The success of international certification depends upon close co-operation between maintainers, seed producers, traders and the designated authority (appointed by the government) in each participating country. Frequent meetings allow for a multi-stakeholder dialogue to exchange information, discuss case studies, revise rules and update the Schemes. A wide range of international and non-governmental organisations as well as and seed industry networks participate actively in the Schemes.

*Benefits of the Schemes*

* + To facilitate international trade by using harmonised certification procedures, crop inspection techniques and use of control plots. The varietal purity standards for the appropriate species are also agreed and standardised by all member states.
  + To provide a framework to develop seed production with other countries or companies.
  + To participate in the elaboration of international rules for seed certification.
  + To develop collaboration between the public and private sectors.
  + To benefit from regular exchanges of information with other national certification agencies and Observer organisations.

*Annual List of Varieties*

The Annual List of Varieties eligible for OECD certification includes varieties which are officially recognized as distinct, uniform and stable, and possess an acceptable value in one or more participating country. The List contains the seed varieties internationally traded using the OECD seed Schemes. The number of varieties included has grown steadily over the last thirty years. Currently, the number of listed varieties amounts to over 62 000, corresponding to 200 species. The List is available online and updated frequently.

*Outlook*

As seed “consumers” become more demanding, there are greater needs for uniform seed standards, while at the same time public financial resources for regulation and quality control are limited.

Co-operation among countries and stakeholders in the framework of the Schemes is a response to the concern for a market-responsive regulatory approach*.* Every country is confronted with a different legal framework, institutional barriers and trade relations whilst the different approaches must remain consistent between countries entering international markets as importers or exporters of seed.

Maintainers and seed companies are responsible for ensuring their varieties remain pure and true to the description and the definitive sample (which is the ‘living description’ of the variety) not only domestically, but also across borders. However, there is a need for minimum criteria to be commonly defined, endorsed and enforced when multiplying seed in large quantities for the trade. The OECD Seed Schemes provide this legal framework at international level.

*Status of Biochemical and Molecular Techniques (BMT) in the OECD Seed Schemes*

The OECD Seed Schemes do not specifically endorse any laboratory method for determining varietal identity or for determining varietal purity. The traditional OECD methods of using field inspection techniques together with pre- and post- control plots are to be regarded as the required methods of determining varietal identity and varietal purity.

However, the OECD Seed Schemes do recognise that there are occasions where these traditional methods limit the certainty of the varietal determination, and in some cases varieties of some species cannot be identified with certainty using these traditional methods. In these specific circumstances, it might be beneficial to use non-field based techniques such as BMT, which must be seen as supplementing and not replacing the more traditional methods.

For more information on the OECD Seed Schemes see: [**www.oecd.org/tad**/**seed**](http://www.oecd.org/tad/seed)

International Union for the Protection of New Varieties of Plants (UPOV)

Type of Organization: Intergovernmental

Membership

[List of UPOV members](http://www.upov.int/export/sites/upov/members/en/pdf/pub423.pdf)  / [Situation in UPOV](http://www.upov.int/export/sites/upov/images/worldmap_en.jpg)

*What is UPOV?*

The International Union for the Protection of New Varieties of Plants (UPOV) is an intergovernmental organization based in Geneva, Switzerland. UPOV was established in 1961 by the International Convention for the Protection of New Varieties of Plants (the "UPOV Convention").

The mission of UPOV is to provide and promote an effective system of plant variety protection, with the aim of encouraging the development of new varieties of plants, for the benefit of society.

The UPOV Convention provides the basis for members to encourage plant breeding by granting breeders of new plant varieties an intellectual property right: the breeder’s right.

*What does UPOV do?*

UPOV’s mission is to provide and promote an effective system of plant variety protection, with the aim of encouraging the development of new varieties of plants, for the benefit of society. The main objectives of UPOV are, in accordance with the UPOV Convention, to:

* provide and develop the legal, administrative and technical basis for international cooperation in plant variety protection;
* assist States and organizations in the development of legislation and the implementation of an effective plant variety protection system; and
* enhance public awareness and understanding of the UPOV system of plant variety protection.

*What are the benefits of plant variety protection and UPOV membership?*

The UPOV Report on the Impact of Plant Variety Protection demonstrated that in order to enjoy the full benefits which plant variety protection is able to generate, both implementation of the UPOV Convention and membership of UPOV are important. The introduction of the UPOV system of plant variety protection and UPOV membership were found to be associated with:

(a) increased breeding activities,

(b) greater availability of improved varieties,

(c) increased number of new varieties,

(d) diversification of types of breeders (e.g. private breeders, researchers),

(e) increased number of foreign new varieties,

(f) encouraging the development of a new industry competitiveness on foreign markets, and

(g) improved access to foreign plant varieties and enhanced domestic breeding programs.

In order to become a UPOV member the advice of the UPOV Council in respect of the conformity of the law of a future member with the provisions of the UPOV Convention is required. This procedure leads, in itself, to a high degree of harmony in those laws, thus facilitating cooperation between members in the implementation of the system.

*Does UPOV allow molecular techniques (DNA profiles) in the examination of Distinctness, Uniformity and Stability (“DUS”)?*

It is important to note that, in some cases, varieties may have a different DNA profile but be phenotypically identical, whilst, in other cases, varieties which have a large phenotypic difference may have the same DNA profile for a particular set of molecular markers (e.g. some mutations).

In relation to the use of molecular markers that are not related to phenotypic differences, the concern is that it might be possible to use a limitless number of markers to find differences between varieties at the genetic level that are not reflected in phenotypic characteristics.

On the above basis, UPOV has agreed the following uses of molecular markers in relation to DUS examination:

(a) Molecular markers can be used as a method of examining DUS characteristics that satisfy the criteria for characteristics set out in the General Introduction if there is a reliable link between the marker and the characteristic.

(b) A combination of phenotypic differences and molecular distances can be used to improve the selection of varieties to be compared in the growing trial if the molecular distances are sufficiently related to phenotypic differences and the method does not create an increased risk of not selecting a variety in the variety collection which should be compared to candidate varieties in the DUS growing trial.

The situation in UPOV is explained in documents TGP/15 “Guidance on the Use of Biochemical and Molecular Markers in the Examination of Distinctness, Uniformity and Stability (DUS)” and UPOV/INF/18 “Possible use of Molecular Markers in the Examination of Distinctness, Uniformity and Stability (DUS)”.

<https://www.upov.int/about/en/faq.html#QB80>

International Seed Testing Association (ISTA)

ISTA’S VISION: UNIFORMITY IN SEED TESTING

Founded in 1924, with the aim to develop and publish standard procedures in the field of seed testing, ISTA is inextricably linked with the history of seed testing. With member laboratories in over 80 countries/distinct economies worldwide, ISTA membership is truly a global network.

Our association produces internationally agreed rules for seed sampling and testing, accredits laboratories, promotes research, provides international seed analysis certificates and training, and disseminates knowledge in seed science and technology on behalf of our membership and governed by its member countries/distinct economies. This facilitates seed trading nationally and internationally, and therefore contributes to food security.

ISTA’S MEMBERSHIP 2019

With member laboratories in 82 countries/distinct economies worldwide, ISTA membership is a truly global network. Currently, ISTA membership consists of:

* 235 Member Laboratories, out of which 136 are ISTA accredited
* 63 Associate Members
* 39 Personal Members

ISTA’S TECHNICAL WORK

The principle objective of ISTA Technical Committees is to develop, standardise and validate methods for sampling and testing of seed quality, using the best scientific knowledge available. They enhance the **ISTA ‘International Rules for Seed Testing’** and develop ISTA Handbooks on seed methods including sampling and testing. Further they are responsible for the organisation of Symposia, Seminars and Workshops. ISTA Technical Committees regularly hold workshops which provide a platform for training as well as the exchange of information, experience and ideas.

There are 20 Technical Committees in ISTA:

|  |  |
| --- | --- |
|  | Technical Committees |
| 1. | Advanced Technologies Committee |
| 2. | Bulking and Sampling Committee |
| 3. | Editorial Board of Seed Science and Technology |
| 4. | Flower Seed Testing Committee |
| 5. | Forest Tree and Shrub Seed Committee |
| 6. | Germination Committee |
| 7. | GMO Committee |
| 8. | Moisture Committee |
| 9. | Nomenclature Committee |
| 10. | Proficiency Test Committee |
| 11. | Purity Committee |
| 12. | Rules Committee |
| 13. | Seed Health Committee |
| 14. | Seed Science Advisory Group |
| 15. | Statistics Committee |
| 16. | Seed Storage Committee |
| 17. | Tetrazolium Committee |
| 18. | Variety Committee |
| 19. | Vigour Committee |
| 20. | Wild Species Working Group |

ISTA ACCREDITATION PROGRAMME:

ISTA Accreditation verifies whether a laboratory is technically competent to carry out seed sampling and testing procedures in accordance with the [ISTA International Rules for Seed Testing](https://www.seedtest.org/en/international-rules-for-seed-testing-2019-_content---1--1083--1065.html). Accredited laboratories must run a quality assurance system, fulfilling the requirements of the [ISTA Accreditation Standard](https://www.seedtest.org/upload/cms/user/ISTAAccreditationStandardforSeedTestingandSeedSamplingV6.11.pdf). Accreditation can be granted for:

* entities performing sampling only
* laboratories performing testing only
* laboratories performing sampling and testing.

ISTA CERTIFICATES: PASSPORT FOR INTERNATIONAL SEED TRADING

Only ISTA-accredited laboratories are authorised to issue ISTA certificates for seed analysis.

By reporting seed test results on ISTA Certificates, the issuing laboratory assures that the sampling and testing has been carried out in accordance with the ISTA Rules. ISTA Certificates are accepted by most authorities and are mentioned in the seed Acts of several countries.

The ISTA certificates are assuring that the results are reproducible, true and represent the quality of the seed.

More than 200,000 ISTA Orange and Blue Certificates are issued every year, facilitating trading of seed internationally.

**THE STATUS OF BIOCHEMICAL AND MOLECULAR TECHNIQUE (BMT) IN ISTA.**

The ISTA International Rules for Seed Testing have included BMTs for many years. For example, BMTs are acceptable for GMO testing under a "performance-based approach"; methods that are frequently used include qualitative and quantitative protein detection analyses and various DNA-based methods. BMTs are used as diagnostic and quantitative assessment tools in seed health testing methods. Testing for species and varieties verification also makes use of BMTs by analysing storage protein profiles for sunflower, maize, oat, barley, wheat, rye grass and pea or by DNA fingerprint using molecular markers for maize and wheat. As the versatility of these methods increases and the cost of utilizing them decreases, they may in the future play an even larger role in seed testing.

To learn more about ISTA, visit our website: [www.seedtest.org](http://www.seedtest.org)

[End of Annex II and of document]

1. Held via electronic means on October 26 and 27, 2021 [↑](#footnote-ref-2)
2. hosted by the United Kingdom and held via electronic means, from May 23 to 27, 2022 [↑](#footnote-ref-3)
3. held via electronic means, from July 11 to 15, 2022 [↑](#footnote-ref-4)
4. hosted by Germany held via electronic means, from June 13 to 17, 2022 [↑](#footnote-ref-5)
5. held via electronic means, from September 19 to 23, 2022 [↑](#footnote-ref-6)
6. at its fifty-sixth session, held via electronic means, from April 18 to 22, 2022 [↑](#footnote-ref-7)
7. at its fifty-first session, hosted by the United Kingdom and held via electronic means, from May 23 to 27, 2022 [↑](#footnote-ref-8)
8. at its fifty-fourth session, , hosted by Germany held via electronic means, from June 13 to 17, 2022 [↑](#footnote-ref-9)
9. at its fifty-third session, held via electronic means, from July 11 to 15, 2022 [↑](#footnote-ref-10)
10. at its first session, held via electronic means, from September 19 to 23, 2022 [↑](#footnote-ref-11)