



# **Specific Accreditation Criteria Life Sciences ISO/IEC 17025 Annex**

## **Facilities testing for genetically modified organisms (GMO)**

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## **Facilities testing for genetically modified organisms (GMO)**

This document provides additional interpretative criteria and recommendations for the application of ISO/IEC 17025 for both applicant and accredited facilities conducting testing for GM material.

Applicant and accredited facilities must also comply with ISO/IEC 17025, the NATA ISO/IEC 17025 Standard Application Document (SAD) and the Life Sciences ISO/IEC 17025 Appendix.

The clause numbers in this document follow those of ISO/IEC 17025 but since not all clauses require interpretation the numbering may not be consecutive.

Testing for genetically modified sequences in foodstuffs or whole grain and related plant materials relies on the ability to test for the specific DNA sequences associated with modifications and/or the promoting and terminating sequences associated with the inserted sequences.

Accreditation covers qualitative and quantitative analysis of GM material through the use of DNA extraction and Polymerase Chain Reaction (PCR) methods.

In recognition of the continual development of new GM materials NATA operates a flexible scope policy. When a facility has developed the capacity to detect a new GM event or construct this will be added to the scope of accreditation without the need for supporting documentation to be submitted to NATA where the analysis is undertaken using the same technique that is covered by the scope of accreditation. Where a new technique is implemented that has not been assessed an on site variation visit will be required. The competence of a facility to develop and implement tests for new GM events and constructs will be ascertained at the initial assessment. At reassessment a sample of GM events or constructs added since the last technical assessment will be reviewed.

### **5.3 Accommodation and environmental conditions**

In order to reduce the risk of false positive results from cross-contamination or carry-over contamination of samples and reagents by other samples in the laboratory or by amplified material, four physically separate and contained areas with known air conditioning/ventilation airflows are required within a facility undertaking nucleic acid amplification for the detection of GM materials. The four work areas are:

- Sample preparation
- DNA extraction
- Reagent preparation
- Product analysis

## **5.4 Test and calibration methods and method validation**

The following reference documents covering nucleic acid testing should be consulted:

- Former Subcommittee of Animal Health Laboratory Standards (SCAHLs) *Veterinary Laboratory Guidelines for Nucleic Acid Detection Techniques*
- National Pathology Accreditation Advisory Council *Laboratory Requirements for medical testing of human nucleic acids 2013*

### **5.4.5 Method validation**

Where possible consideration should be given to grouping like matrices to reduce the volume of work required to undertake validation.

See the FSANZ website at [www.foodstandards.gov.au](http://www.foodstandards.gov.au) information on the currently approved/pending GM events.

## **5.5 Equipment**

Specialised equipment (i.e. thermocycler, nucleic acid analyser, spectrophotometer) is required for detection of labeled DNA fragments. Many instruments have internal diagnostic checks built into them and, as a minimum, the recommendations from manufacturers should be followed.

## **5.6 Measurement Traceability**

Reference materials used as controls in the validation of new assays and as routine controls must comply with the requirements of NATA's *Metrological Traceability*.

## **5.8 Handling of test and calibration items**

The high sensitivity of methods dictates a higher than usual awareness to the possibility of cross-contamination during transport, storage, preparation and analysis. (See 5.3 Accommodation.)

## **5.9 Assuring the quality of test and calibration results**

Controls that are incorporated with the GMO test system must address inhibition, sensitivity and contamination.

## **5.10 Reporting of results**

Reporting the results of GM material detection must be sufficiently descriptive to allow the reader a clear understanding of what tests have been conducted and to what level of detection.

In addition to the general requirements of ISO/IEC 17025, a test report for GM material testing must also include information on the limitations of the particular detection method used for the particular sample or sample type:

The content of test reports is influenced by the nature of the matrix tested. For unprocessed materials such as soya beans, canola seeds or lupin seeds it is important to include the amount of sample or the number of seeds ground prior to DNA extraction.

Where highly processed materials are tested it is possible that an insufficient quantity of DNA can be extracted to allow the presence of GM material to be detected. In such cases a null result can be reported. Facilities should discuss the possibility of a null result with customers prior to testing if the matrix submitted is highly processed.

No affirmation shall be made stating that there is no GM material present in the sample analysed as determined from test samples.

## References

*Veterinary Laboratory Guidelines for Nucleic Acid Detection Techniques 2008*  
Former Subcommittee of Animal Health Laboratory Standards

*Requirements for medical testing of human nucleic acids 2013*  
National Pathology Accreditation Advisory Council

## NATA Publications

Life Sciences ISO IEC 17025 Appendix

## Amendment Table

The table below provides a summary of changes made to the document with this issue.

<b>Section or Clause</b>	<b>Amendment</b>
New document	This document represents a direct adoption of the former Biological Testing Annex D. The document has been reviewed and updated to reflect the new accreditation criteria documentation structure.