



Netherlands Food and Consumer
Product Safety Authority
Ministry of Economic Affairs

Obtaining an ISO17025 accreditation for PCR-sequencing

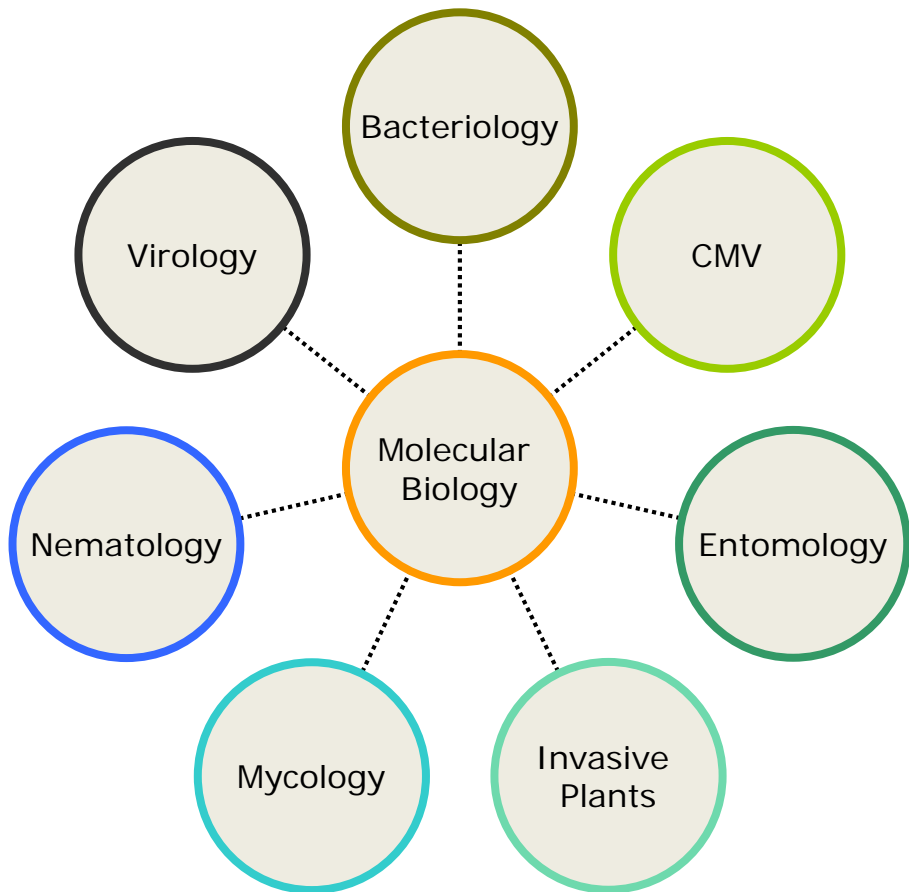
Experiences of the Dutch NPPO

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Molecular Biology at NPPO-NL



- MolBio is a separate group of NRC
- Diagnostics and applied research
- Conventional (RT-)PCR (RFLP)
- Real-time (RT-)PCR
- PCR-sequencing (Sanger)
- At present ISO17025 accredited for several real-time PCR tests under a fixed scope



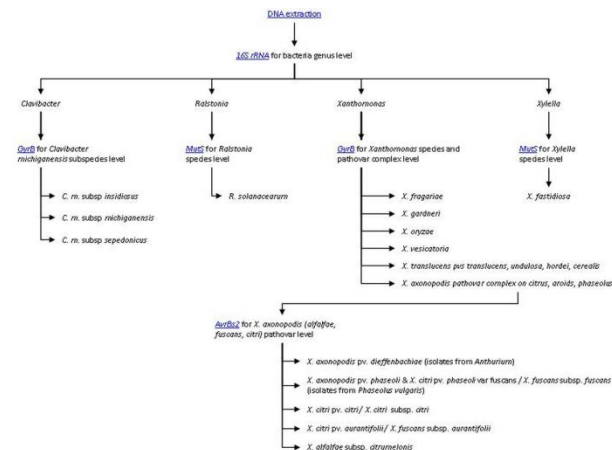
Ambitions of the NPPO-NL

- Accreditation of phytosanitary diagnostics under flexible scope
- Emphasis on overall species identification instead of individual tests
- Flexibility the detection and identification of organism A in matrix B using tests X, Y, Z
- Covering all methods (not tests) performed in-house
- For MolBio, this includes molecular identification using PCR-sequencing according to Sanger



PCR-Sequencing (DNA Barcoding)

- Diagnostic method that uses a short standardised genetic marker in an organism's DNA to aid species identification
- Plant pest DNA Barcoding protocols from the QBOL project, Q-bank and EUPHRESKO DNA Barcoding project and selected virological tests
- EPPO standard on DNA barcoding for plant pests (viruses excluded) under preparation
- Tests are not used stand-alone, but always in support of the diagnosis,





PCR-sequencing at NPPO-NL



Aromia bungii



Validation: a requirement for accreditation

- One major challenge for accreditation of PCR-sequencing
- Validation: is the test fit for purpose?
- Determination of performance criteria (EPPO PM 7/98)
 - › Analytical sensitivity, Analytical specificity, Selectivity, Repeatability, Reproducibility, Robustness
- Well described standard operating procedures (A-SOP) are the object of validation
 - › Not available for consensus sequence preparation and analysis of consensus sequence(s)



Regular review of validation status

- ISO17025 requirement: Regular review of performance criteria should be carried out to verify that a test is still fit for purpose
- Challenging when using online databases with changing content:

BOLDSYSTEMS Databases | Taxonomy | Identification | Workbench | Resources

Animal Identification [COI] | Fungal Identification [ITS] | Plant Identification [rbcL & matK]

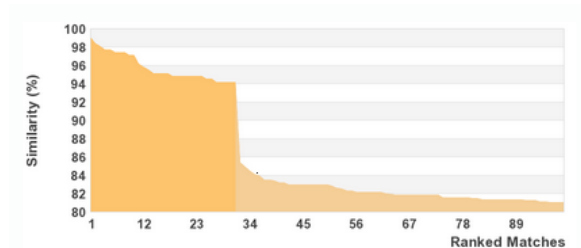
Historical Databases: [Jul-2013](#) [Jul-2012](#) [Jul-2011](#) [Jul-2010](#) [Jul-2009](#)

Species Level Barcode Records (1,540,956 Sequences/138,653 Species/56,354 Interim Species)

▼ Query: `unlabeled_sequence` Top Hit: No match

Unable to match any records in the selected database

2009 - 2012



Search Result:

The submitted sequence has been matched to *Anthonomus grandis*. This identification is solid unless there is a very closely allied congeneric species that has not yet been analyzed.

2013 - present



Remarks made during the ISO17025 audit underline the challenges of PCR-sequencing validation and verification





How to tackle the problem?

- Separation of PCR-test and sequence analysis
- Determine performance criteria for the two parts separately
- PCR test
 - › Standardised procedures in A-SOP
 - › Determine relevant performance criteria once, and update at regular intervals (5 years) following PM7/98
- Sequence analysis
 - › General guidance: depends on expertise technician/researcher
 - › Usability of the generated sequence for molecular identification on a certain taxonomical level is determined on *ad hoc* basis
 - › Last analysis report is the up-to-date validation status for an “organism-matrix-locus (loci)” combination



PCR test versus sequence analysis

DNA extraction

PCR

Cycle sequencing

Sequencing

Consensus sequence

Data-analysis

Test specific
A-SOP

Expertise
technician

Determination or verification of Performance criteria

Analytical sensitivity – mainly relevant for viruses, viroids and phytoplasmas

Analytical specificity – A (small) subset of target organisms (the full range of target organisms is not known)

Selectivity – relevant for viruses, viroids and phytoplasmas

Repeatability and Reproducibility

Robustness – relevant when introducing small changes to the test



PCR test versus sequence analysis

DNA extraction

PCR

Cycle sequencing

Sequencing

Consensus sequence

Data-analysis

Test specific
A-SOP

Expertise
technician

Analytical specificity - can change from day to day depending on the content of the (online) sources consulted.

Determined during each analyses

Inter- versus intraspecies variation

Clustering in taxon specific clades

Using one or more loci

All results and observations have to be documented to ensure traceability

Conclusions are drawn using conservative terms

Repeatability and reproducibility – first and second line controls



First line controls

- PIC, NIC, PAC, NAC and a positive cycle sequence control
- PAC is a synthetic construct per organism group
- PAC is used as a process control to monitor all steps from amplification to sequence analysis (also used for reproducibility and repeatability)

Negative isolation control

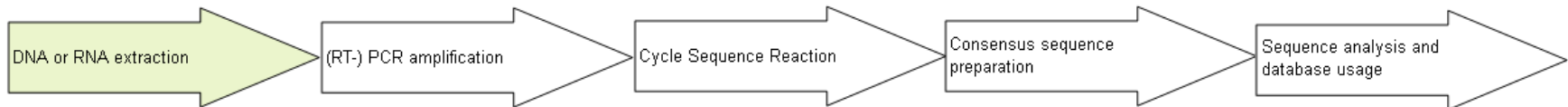
Positive isolation control
(amplicon generated with
diagnostic sample using
a generic PCR test)

Negative amplification control

Positive amplification control - synthetic construct (sequence in NCBI)

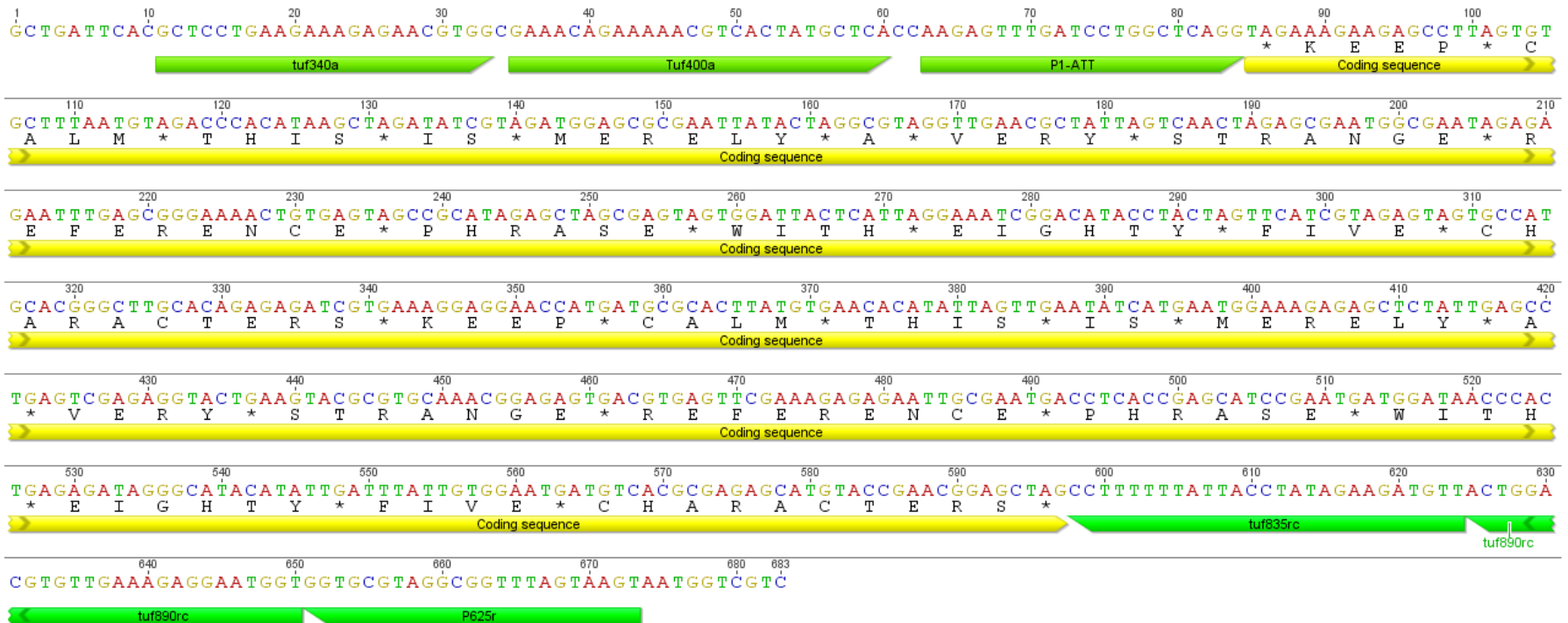
Also used to determine Reproducibility and Repeatability

Positive cycle sequence control





Finding the Easter egg





Internal Control Program

- How to make sure that technicians are proficient?
 - › Internal training sessions
 - › Internal workshops discussing “though” cases
 - › Second assessor in particular cases (new organism-matrix-locus (loci) combinations)
 - › 2nd line of controls (blind samples)
 - › 3rd line of controls (proficiency tests) (e.g. Pospiviroids)
- Discussing results with taxonomical specialists increases the level of expertise
 - › Being able to estimate the coverage of taxa in databases
 - › Knowledge on organism-matrix-locus (loci) combinations
 - › Knowledge on synonyms used



Conclusions

- We believe it is possible to acquire an accreditation for PCR-sequencing
- The Dutch accreditation board (RvA) is open to accreditation of methods in which expertise plays an important role
- We need to:
 - › add extra validation data on the PCR test part of the method (mainly sensitivity and selectivity where relevant)
 - › Keep track of repeatability and reproducibility of the sequence analysis part of the method over time (synthetic PACs)
- Report back to RvA in the coming eight months with our solutions for shortcomings found during the audit



Thank you very much for your attention

Are there any questions?



Acknowledgements

- Maureen Bruil, Eveline Metz-Verschuren, Jeanette Teunisse, Esther van Veen, Joris Voogd, Tim Warbroek
- QBOL and EUPHRESCO DNA barcoding partners
- Technical assessors ISO17025 audit