

Variety Committee Report

ISTA Annual Meeting 2024



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AGENDA

Members

Markers for variety identification / verification and laboratory accreditation

Update on the development on new markers for detection of annual types in perennial rye grass varieties

Use of neuronal networks for variety identification

Handbook

Acknowledgments

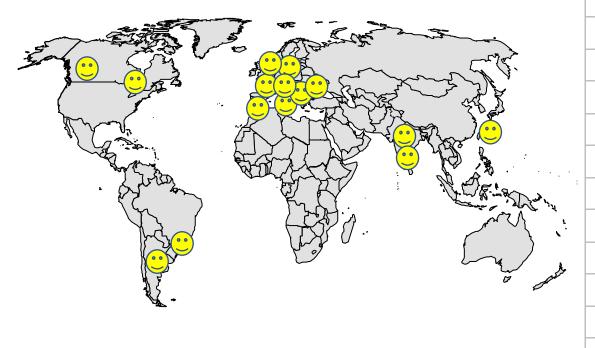








Members



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COMMITTEE MEMBERSHIP LIST Chair: Ana Laura Vicario Member since 2007 1 Vice-Chair: Marie-Claude Member since 2020 2 Gagnon Anne Bernole Member since 2016 З. Member since 2004 Berta Killermann 4 Member since 2007 5 Chiara Delogu Kae-Kang Hwu Member since 2007 6 Ksenija Markovic Member since 2013 7 Ksenija Taski-Ajdukovic Member since 2010 8 Keshavulu Kunusoth Member since 2010 9 Ana Patricia Fernandez Getino Member since 2021 10 Mariana Menoni 11 Member since 2021 12 Sean Walkowiak Member since 2022 13 Umashankar Bellan Member since 2023 Member since 2023 14 Stephanie Guillet Member since 2023 15 Lorella Andreani

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Markers for variety identification / verification and laboratory accreditation



VARCOM SCOPE

The purpose of species and variety verification is to determine the extent that the submitted sample conforms to the species or variety as requested by the applicant, using methods not permissible in a purity test according to Chapter 3.

WHAT WE HAVE NOW

CONVENTIONAL METHODS: morphological traits, possible to see under daylight or UV light, by naked eye or using some magnifier, with or without previous treatment.

PROTEIN-BASED: refer to standard methods for obtaining storage proteins band patterns that allow variety verification.

DNA-BASED: refer to a semi-performance based methods for obtaining DNA patterns that provide a tool for assessing laboratory performance aimed to laboratory accreditation.

WHAT IS UNDER DEVELOPMENT AND NEW TO COME

DNA-based test for specific traits for variety identification: Lolium test to discriminate ARG from PRG

Morphological based test using New Technologies Equivalent to the Human Analyst for Seed Analysis: Use of neuronal networks for variety identification





Project leader: Giovanny López from ATC

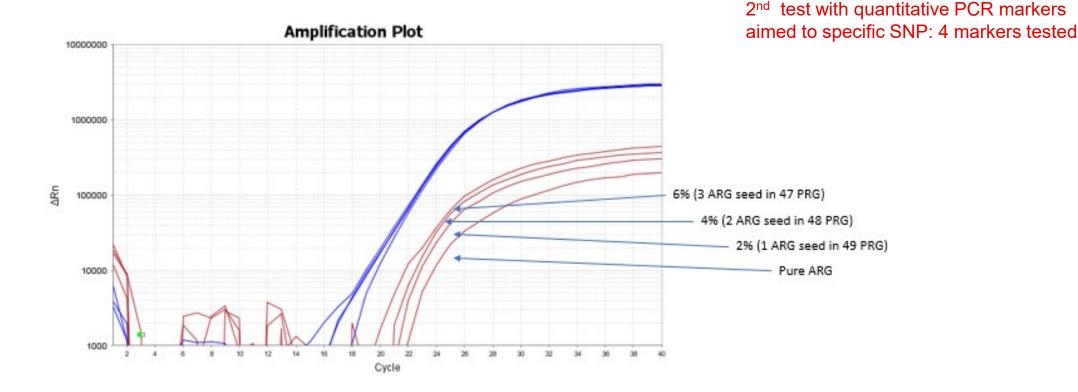
Collaboration with Shaun Bushman from USDA who sequenced ARG and PRG varieties and developed about 10 markers.

Daniel Curry from Oregon State University who provided seeds samples for the test and technical support.

- 1st test done using KASP system (results presented last year).
- 2nd test with quantitative PCR markers aimed to specific SNP: 4 markers tested.
- 3rd test on a larger set of annual, perennial and hybrid seed samples.
- 4th test on spiked samples (undergoing).
- 5th test interlaboratory trial with best markers (next year).



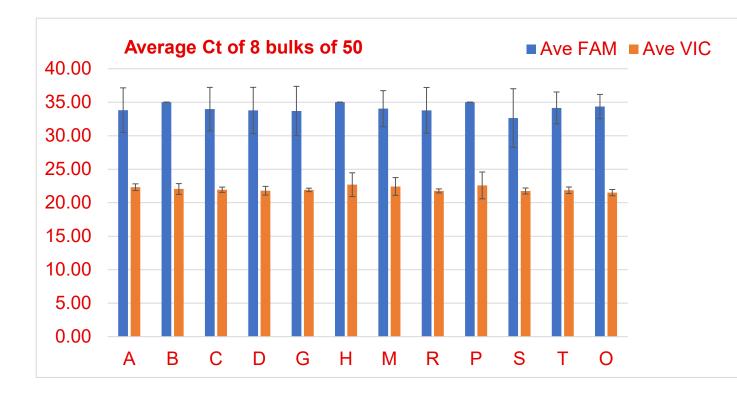












3rd test on a larger set of annual, perennial and hybrid seed samples

Test on 12 perennial varieties

FAM fluorophore is for the annual ryegrass

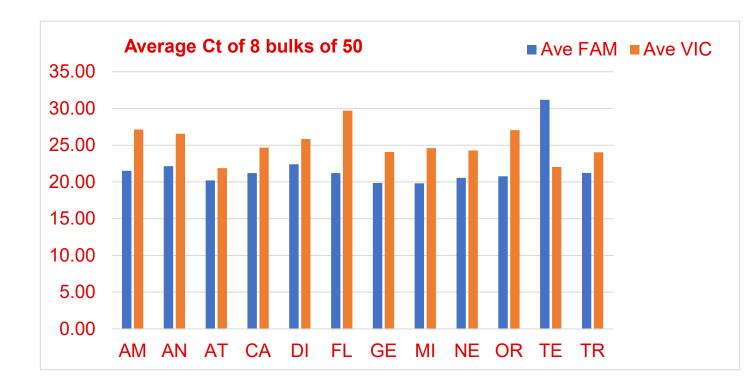
VIC fluorophore is for the perennial ryegrass

Example for 1 of the 3 markers tested









3rd test on a larger set of annual, perennial and hybrid seed samples

Test on 10 annual varieties + last two hybrids

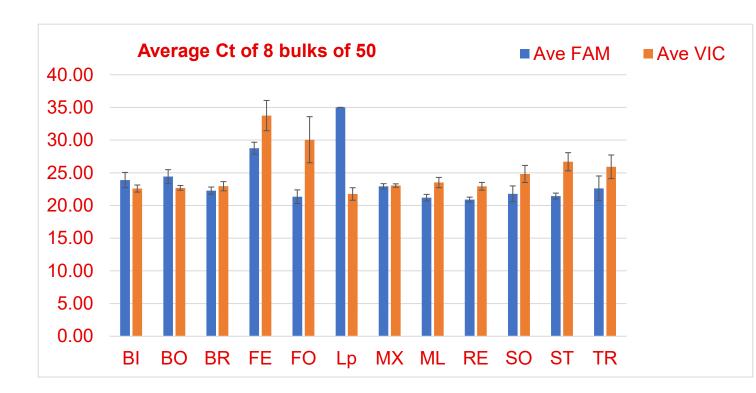
FAM fluorophore is for the annual ryegrass

VIC fluorophore is for the perennial ryegrass

Example for 1 of the 3 markers tested







3rd test on a larger set of annual, perennial and hybrid seed samples

Test on 11 hybrids varieties + 1 perennial (Lp)

FAM fluorophore is for the annual ryegrass

VIC fluorophore is for the perennial ryegrass

Example for 1 of the 3 markers tested







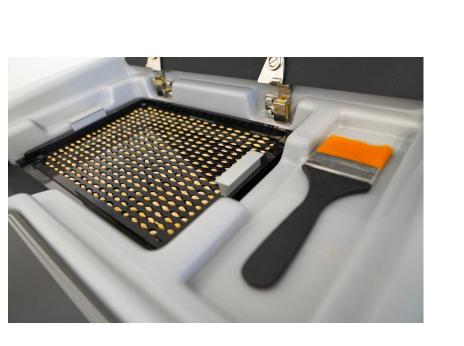
Use of neuronal networks for variety identification



Method used in 3 countries for wheat and 15 countries for barley

We ran a spiking test and a repeatability test.

Advice from STACOM: run again the spiking test in a different laboratory for checking on robustness, repeatability and reproducibility. This second test is undergoing.



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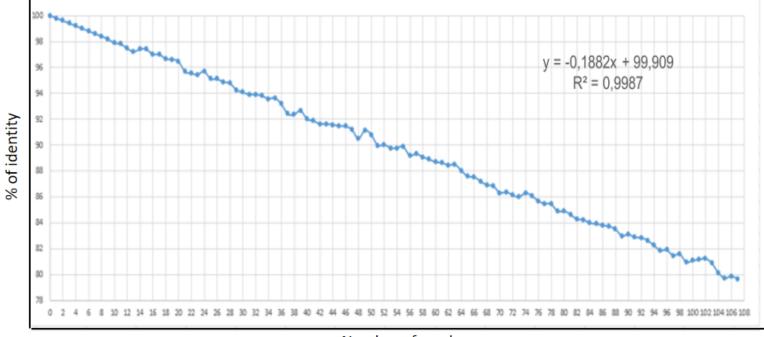






Use of neuronal networks for variety identification

Spiking test: testing the impact of 1 single seed on the % of identity



Each dot correspond to the identity value obtained after taken out one seed from the grid and replace it by a seed from another variety. The "intercept" value is closed to the expected 0.2% value, which is the % of each seed in a 500 seeds grid (1/500).



Number of seeds





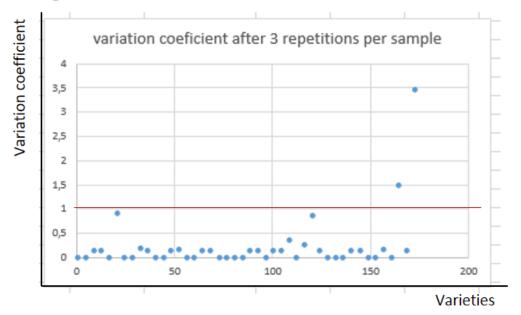
Use of neuronal networks for variety identification



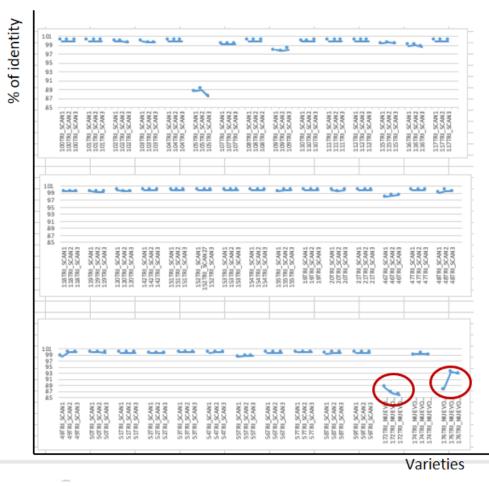
Repeatability test

191A

Each group correspond to a variety, tested 3 times. Less than 5% of the samples presented variation of results higher than 1.



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Handbook

6 members in the working group

HANDBOOK ON DNA BASED TESTS - Table of Contents

i. Preface

ii. Health and Safety Information

iii. Acknowledgments

iv. Contributors

Content

Introduction

- 1.1 Summary
- 1.2 History of DNA-based varietal identification and ISTA
- 1.3 The purpose of DNA-based testing in the ISTA Rules

1.4 Goal and scope of the ISTA "Handbook on DNA testing" Development: guidelines for Comparative Tests (CTs) organization

Considerations on the varieties

Considerations on the sample size

Considerations the on markers

Other considerations



HANDBOOK ON DNA BASED TESTS – Table of Contents

Validation: validation of DNA based markers (guide to following ISTA SOP for validation of methods) Adoption as an official method Reference material collection (RMC) Statistical approaches for results analysis (including : evaluation of performance method and proficiency tests) draft to be presented to STATCOM ISTA accreditation: evaluation of performance method and proficiency tests Statistical approaches for results analysis ISTA accreditation for DNA based testing (to be discussed with Acc department) Auditing laboratories for DNA-based Testing (to be discussed with Acc department) Appendices





Acknowledgments



We would like to thank all VARCOM members and collaborators for their contribution to the committee work aimed to achieving ISTA goals









Thank you for your attention



