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Certified Reference Materials AOCS 1114-B2

Report of the certification process for

NP2391/NP2222 (MZIR098)

Maize Certified Reference Materials

Second Batch

OECD Unique Identifier SYN-ØØØ98-3

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Abstract

This report describes the preparation and certification of the maize Certified Reference Material (CRM) AOCS 1114-B2 produced by AOCS Technical Services in 2020. The CRMs have been prepared by AOCS according to ISO 17034:2016 and are intended to serve as control material for third party testing of maize for transformation events. NP2391/NP2222 (MZIR098) maize seed powder was provided by Syngenta Crop Protection, LLC and was prepared by grinding the bulk source at AVEKA, Inc., Woodbury, MN (an ISO 9001:2015 accredited facility). The NP2391/NP2222 (MZIR098) maize seed powder was then aliquoted and packaged under a nitrogen gas environment at Illinois Crop Improvement Association (an ISO 17025:2017 accredited facility). The certified mass value of MZIR098 in NP2391/NP2222 (MZIR098) maize was based on seed purity and with 95% confidence, it is at least 745 g/kg. Homogeneity testing was performed at Eurofins-GeneScan New Orleans, LA (an ISO 17025:2005 accredited laboratory) using quantitative real-time PCR after the CRM AOCS 1114-B2 was bottled. Homogeneity results indicated that this CRM AOCS 1114-B2 is homogeneous and were used to verify the presence of the Event MZIR098 in this CRM. CRM AOCS 1114-B2 is available in 27mL glass headspace vials. This CRM shall be stored dry in a sealed container at ambient or cooler conditions in the dark.

Acknowledgements

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Glossary

AOCS American Oil Chemists' Society

Conventional Crop Conventional counterpart means a related organism/variety,

its components and/or products for which there is experience

of establishing safety based on common use as food

Cycle threshold (Ct) Number of PCR cycles required for the fluorescent signal to

cross a threshold that exceeds background level

DNA Deoxyribonucleic Acid is the linear, double-helix

macromolecule that makes up the genetic material of most

organisms

Detection Limit Lowest level at which target DNA can be detected in a sample

EC European Commission

Genome The full set of genes and associated DNA characteristic of an

organism

GMO Genetically modified/engineered organism: an organism in

which the genetic material has been changed through modern

biotechnology in a way that does not occur naturally by

multiplication and/or natural recombination

ISO International Organisation for Standardisation

PCR Polymerase Chain Reaction: technique used to determine

whether a sample of plant tissue contains a particular DNA

sequence. PCR relies on primer sets that bind to a particular

target DNA sequence and a special DNA-copying enzyme

(DNA polymerase) that exponentially amplifies the target sequence for identification and measurement

Qualitative PCR PCR methods that determine the presence or absence of a

specific target DNA sequence at a particular level of detection

Quantitation Limit Lowest level at which the amount of target DNA sequence in

a sample can be reliably quantified

Quantitative PCR PCR methods that estimate the relative amount of target

DNA sequence in a mixture of DNA molecules

RSDr Relative standard deviation

SD Standard deviation

Trait: NP2391/NP2222 Line of maize, genetically engineered to provide dual modes

(MZIR098) of action for control of corn rootworm (*Diabrotica* species)

Introduction

Plant biotechnology is an extension of traditional plant breeding. It allows plant breeders

to develop crops with specific traits including insect, disease, and herbicide resistance;

processing advantages; and nutritional enhancement. An important component for

identifying these new traits is a Certified Reference Material (CRM) created from leaf,

seed, or grain containing the new trait as well as a CRM created from the conventionally

bred matrix. The European Commission has mandated that from 18 April 2004, a method

for detecting a new event derived from transgenic technology and Certified Reference

Material must be available before the EC will consider authorization a new crop derived

from transgenic technology. Several nations outside Europe also require grain and

ingredients to be labeled above a threshold level before accepting a shipment.

To meet the above regulatory requirements for GMO determination, CRM AOCS 1114-

B2 was manufactured from maize seed according to ISO 17034:2016 and in accordance

with EC No 1829/2003, EC No 641/2004 and EC No 619/2011. This CRM is available

from AOCS.

Material Processing and Particle Size Analyses

The hemizygous NP2391/NP2222 (MZIR098) maize seed used in the preparation of CRM

AOCS 1114-B2 resulted from the cross of female non-transgenic NP2391 and male

NP2222 (MZIR098). The NP2391/NP2222 (MZIR098) maize seed was first milled and

analyzed for particle size distribution at AVEKA, Inc., Woodbury, Minnesota (an ISO

9001:2015 accredited facility).

Bulk seed received by AVEKA, Inc from Syngenta Crop Protection, LLC was milled in a

Fitzmill cryogenic hammermill using first a 690 µm screen. To further reduce particle size,

this ground material was milled again under the same conditions using a 510 µm

screen. The material was blended in a Patterson-Kelley V-blender, and after homogeni-

zation six samples taken at random were subject to particle size analyses using a Horiba

LA-950 Light Scattering Particle Analyzer. For each sample, the particle size mean and

range, and the percentage of particles below a given size was calculated. On average,

Report of Certification for 1114-B2 Page 8 of 17 | ©AOCS, 2024 Document Version: 1.3 Published: 19 July 2024 the particle size of CRM AOCS 1114-B2 was 167.15 \pm 2.64 μ m, and 90% of the particles were smaller than 355.45 + 5.65 μ m (Table 1).

Table 1. Results of Particle Size Analyses of CRM AOCS 1114-B2 Conducted by AVEKA Stand-Sample Sample Sample Sample ard Sample 1 Sample 2 **Average** 3 4 5 6 Devia-(µm) (µm) (µm) (µm) tion (µm) (µm) (µm) (µm) 169.08 164.70 169.90 166.70 163.48 167.15 2.64 Mean 169.08 0.88-0.88-0.88-0.88-0.88-0.88-

678.50

16.86

147.65

350.56

678.50

18.47

140.45

346.87

678.50

18.59

146.45

359.13

 $N/A^{(a)}$

17.76

145.00

355.45

N/A

0.98

3.42

5.65

678.50

18.59

146.45

359.13

678.50

16.30

140.89

355.61

678.50

17.71

148.10

361.38

Range

D10 (b)

D50 (b)

D90 (b)

The particle size distribution for each of the samples analyzed is presented as a histogram, with the x-axis showing discrete size bins up to 678.5 μ m (Figure 1). Figure 1-A represents the percentage of particles of a given size, and Figure 1-B represents the cumulative particle size distribution, which reflects the total percentage of particles smaller than a given size. For all samples analyzed, 100% of particles were \leq 678.50 μ m.

⁽a) N/A = not applicable

⁽b) D10, D50 and D90 indicate that 10%, 50% or 90% of the particles, respectively, are smaller than size given in table

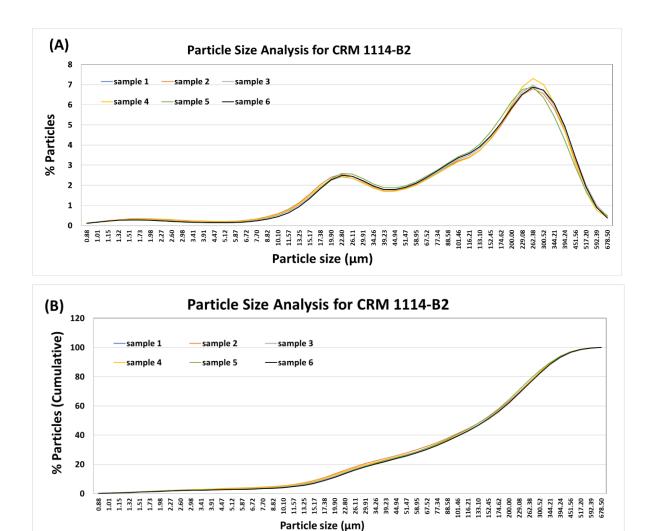


Figure 1. Particle size distribution plots. (A) Percentage of particles of a given size. (B) Cumulative distribution of particle sizes

Bulk NP2391/NP2222 (MZIR098) maize seed powder for the production of CRM AOCS 1114-B2 was delivered to AOCS and it was then aliquoted and packaged in 27-mL glass headspace vials and sealed under a nitrogen gas environment at the Illinois Crop Improvement Association (an ISO 17025:2017 accredited facility).

Certified Value and Measurement Uncertainty

The genetic purity based on the presence of Event MZIR098 in NP2391/NP2222 (MZIR098) maize was assessed by Syngenta Crop Protection, LLC. A total of 379

NP2391/NP2222 (MZIR098) maize seeds were evaluated by qualitative, MZIR098-spe-

cific real time PCR. The results showed that 371 of the 379 seeds tested (97.9%) positive

for the presence of Event MZIR098.

The statistical seed purity of NP2391/NP2222 (MZIR098) was calculated using Seed-

Calc8 (Remund et al., 2008) and corresponded to the lower bound of true % purity. Using

a 95% confidence level, the true % purity of the NP2391/NP2222 (MZIR098) maize seed

lot was at least 74.5%. Consequently, with 95% confidence, the true value is \geq 745 g/kg.

The measurement uncertainty (U_{CRM}) is the expanded uncertainty with a coverage factor

of 2 and a confidence level of 95%. It is obtained by combining the uncertainties from the

purity assessment $(u_{char,rel})$, the homogeneity assessment $(u_{bb,rel})$, the transport stability

assessment $(u_{sts,rel})$ and the long-term stability assessment $(u_{lts,rel})$:

$$u_{CRM,rel} = \sqrt{u_{char,rel}^2 + u_{bb,rel}^2 + u_{sts,rel}^2 + u_{lts,rel}^2}$$

$$U_{CRM} = 2 \times u_{CRM,rel} \times 1000 \, g/kg$$

When using an asymmetric uncertainty, the reported measurement uncertainty is trun-

cated on the right side such that the value does not exceed 1000 g/kg. Consequently, the

expanded measurement uncertainty for this CRM is +21 g/kg, -234 g/kg.

Homogeneity

The material used for the production of CRM AOCS 1114-B2, NP2391/NP2222

(MZIR098) maize, is 96.22% pure and is expected to be homogenous. After

NP2391/NP2222 (MZIR098) maize seed was ground and bottled as described above,

ten samples of CRM AOCS 1114-B2 were randomly selected using the Microsoft Excel

Random Number Generator function and were sent to Eurofins-GeneScan, New Orle-

ans, LA (an ISO 17025:2005 accredited laboratory) for homogeneity testing using

quantitative real-time PCR.

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Homogeneity was assessed after bottling of CRM AOCS 1114-B2 using a MZIR098-**PCR** specific. quantitative real-time method (https://gmocrl.jrc.ec.europa.eu/summaries/EURL-VL-04-17-VM.pdf). A total of 10 samples of CRM AOCS 1114-B2 maize were analyzed, and for each sample, 2 independent DNA extractions and quantifications were performed at Eurofins-GeneScan using a test portion of 2 grams. Extracted DNA was checked for integrity by gel-electrophoresis and quantified prior to using it in quantitative real-time PCR. For each of the DNA extracts, all PCR reactions were done in triplicate.

The cycle threshold (Ct) values for an endogenous adh1 maize gene and for Event MZIR098 were used to calculate the number of copies (cp#) for either target. Subsequently, the ratio between Event MZIR098 copy number and adh1 copy number (MZIR098 cp#/adh1 cp#) was calculated and used to estimate the within-unit relative standard deviation (RSD_w) and between-unit relative standard deviation (RSD_b).

Within-unit relative standard deviation (RSD_w), between-unit relative standard deviation (RSD_b) were calculated as:

Within-unit RSD:
$$RSD_{w} = \frac{\sqrt{MS_{within}}}{\bar{y}}$$
 Between-unit RSD:
$$RSD_{b} = \frac{\sqrt{\frac{MS_{between} - MS_{within}}{\bar{y}}}}{\frac{n}{\bar{y}}}$$

where,

within-unit mean square from an ANOVA MS_{within} between-unit mean square from an ANOVA MS_{between} mean of all results of the homogeneity study \bar{y} mean number of replicates per unit n

Table 2. The within-unit relative standard deviation (RSD _w), and the between-unit relative standard deviation (RSD _b) for vials of AOCS 1114-B2.		
CRM	RSD _w [%]	RSD₀ [%]
AOCS 1114-B2	12.1	11.6

The CRM will be determined to be homogeneous if the within-unit relative standard devi-

ation (RSD_w) and between-unit relative standard deviation (RSD_b) are both \leq 20%. Based

on the quantitative real-time PCR analyses conducted, it was concluded that CRM AOCS

1114-B2 is homogenous (Table 2). These results are in agreement with homogeneity

results from qualitative real-time PCR analyses and with the purity estimate for material

NP2391/NP2222 (MZIR098) calculated in the Certified Value and Measurement

Uncertainty section above.

Trait Verification

The presence of the NP2391/NP2222 (MZIR098) trait was assessed in the same ten

CRM AOCS 1114-B2 samples that were analyzed for homogeneity using MZIR098-

specific quantitative PCR analysis. Quantitative results were converted to qualitative

data, and the results are presented in Table 3. In all instances Event MZIR098 was

present.

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Table 3. Qualitative Results for the verification of CRM AOCS 1114-B2 tested by Eurofins-GeneScan with a MZIR098-specific, quantitative PCR analysis

Sample	NP2391/NP2222 (MZIR098) Presence
AOCS 1114-B2 22	Positive
AOCS 1114-B2 75	Positive
AOCS 1114-B2 105	Positive
AOCS 1114-B2 192	Positive
AOCS 1114-B2 213	Positive
AOCS 1114-B2 280	Positive
AOCS 1114-B2 321	Positive
AOCS 1114-B2 366	Positive
AOCS 1114-B2 414	Positive
AOCS 1114-B2 479	Positive

Stability

Time, temperature and light are regarded as the most relevant influences on the stability of CRM (Linsinger, et al., 2001). The influence of light is mitigated by shipping and storing the vials in boxes, thus minimizing the possibility of degradation due to light. The influence of temperature is mitigated by storing the vials in a temperature-controlled room, and shipping vials at ambient temperature.

The effect of temperature and time are investigated.

A transport (short-term) stability study is conducted to assess the stability of maize CRM

during transport. The temperature and time conditions in the study cover the typical con-

ditions and the not so rare situations. The outcome of the study is considered transferable

to other CRMs of similar property. Samples were subject to 3 different temperatures (4

 $^{\circ}$ C (fridge), 25 $^{\circ}$ C (ambient), 60 $^{\circ}$ C (oven)) for 4 different durations (0, 1, 2, and 4 weeks).

The study concluded that samples are stable at 4 °C (fridge) and 25 °C (ambient) for 4

weeks. The estimated uncertainty contribution from transport (short-term) stability is

1.0%.

A long-term stability study is conducted to assess the stability of maize CRM during stor-

age. Samples are stored at 25 °C (ambient) and the stability of the sample is monitored

as long as the samples is available. The storage temperate studied is 25 °C and the length

of time to be studied is 10 years. The outcome of the study is considered transferable to

other CRMs of similar property. In the initial 1-year stability study, samples were subject

the storage condition for 4 different durations (0, 1, 3, 6 and 12 months). The study con-

cluded that samples are stable at 25 °C (ambient) for 12 months. The estimated uncer-

tainty contribution from long-term stability is 0.42%.

CRM stability over time will be analyzed by repeating the homogeneity study described

above at a chosen shelf life of approximately every 24 months. The 24-month shelf life of

CRM is chosen because the influence of analytical variation can be reduced by increasing

the length of the stability study (Linsinger et al., 2001).

The initial ratio between the number of copies of the GM event and the number of copies

of the endogenous reference gene from the homogeneity study will establish the base

line for the stability study. The ratio at each 24-month interval will be compared to the

ratio established in the homogeneity study. The CRM will be determined to be stable if

the variability of the ratios, determined as relative standard deviation (RSD) is \leq 20%.

Stability of these CRMs has been listed as 2 years from the certification date. The mate-

rials were processed and are stored at ambient temperature, under nitrogen gas, in 27 -

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mL glass headspace vials. These materials are expected to be stable for longer than the estimated expiration date. The stability of the powder material will be reevaluated at time of expiration. If the samples are determined to be stable, the certificates will be extended.

References

AVEKA; Inc., 2045 Wooddale Drive. Woodbury, MN 55125; Telephone: +1 651 730 1729; https://www.aveka.com/

Biosafety Clearing House Living Modified Organism (LMO) Registry http://bch.cbd.int/database/lmo-registry/

Eurofins-GeneScan; 2219 Lakeshore Drive Suite 400, New Orleans, LA 70122; Telephone: +1 504 846 2398; https://www.eurofinsus.com/food-testing/testing-services/gmo/

Illinois Crop Improvement Association, 3105 Research Road, Champaign, IL 61826; Telephone: +1 217 359 4053 Fax: +1 217 359 4075; https://www.ilcrop.com/

ISO 9001:2015, Quality Management Systems – Requirements

ISO 17025:2005 and ISO 17025:2017, General Requirements for the Competence of Testing and Calibration Laboratories

ISO 17034:2016 (E) General requirements for the competence of reference material producers

Linsinger, T., Pauwels, J., van der Veen, A. *et al.* Homogeneity and stability of reference materials. *Accred Qual Assur* **6**, 20–25 (2001). https://doi.org/10.1007/s007690000261

Regulation (EC) No 1829/2003 of the European Parliament and of the Council of 22 September 2003 on genetically modified food and feed (https://eur-lex.europa.eu/legal-content/EN/TXT/PDF/?uri=CELEX:32003R1829&from=en)