



BMT/15/9

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INTERNATIONAL UNION FOR THE PROTECTION OF NEW VARIETIES OF PLANTS

Geneva

**WORKING GROUP ON BIOCHEMICAL AND MOLECULAR
TECHNIQUES AND DNA PROFILING IN PARTICULAR**

Fifteenth Session

Moscow, Russian Federation, May 24 to 27, 2016

**FAST SINGLE-STEP DETECTION AND IDENTIFICATION OF MULTIPLE PHYTOPATHOGENS AND GMO
WITH REAL-TIME PCR-MATRIX TECHNIQUE**

Document prepared by an expert from Russian Federation

Disclaimer: this document does not represent UPOV policies or guidance

The Annex to this document contains a copy of a presentation “Fast Single-step Detection and Identification of Multiple Phytopathogens and GMO with real-time PCR-matrix Technique” to be made at its fifteenth session of the Working Group on Biochemical and Molecular Techniques and DNS-Profiling in particular (BMT).

Alexander Golikov, Science Director, GenBit LLC, Russian Federation

[Annex follows]



**FAST SINGLE-STEP DETECTION AND IDENTIFICATION
OF MULTIPLE PHYTOPATHOGENS AND GMO WITH REAL-
TIME PCR-MATRIX TECHNIQUE**

Alexander Golikov

for

UPOV TECHNICAL WORKING PARTY FOR BIOCHEMICAL AND
MOLECULAR TECHNIQUES, AND DNA-PROFILING IN PARTICULAR
(BMT)

Fifteenth Session, Moscow, Russia, May from 24 to 27, 2016

(with Preparatory Workshop on May 23, 2016)



**FAST SINGLE-STEP DETECTION AND IDENTIFICATION
OF MULTIPLE PHYTOPATHOGENS AND GMO WITH REAL-
TIME PCR-MATRIX TECHNIQUE**

Alexander Golikov

for

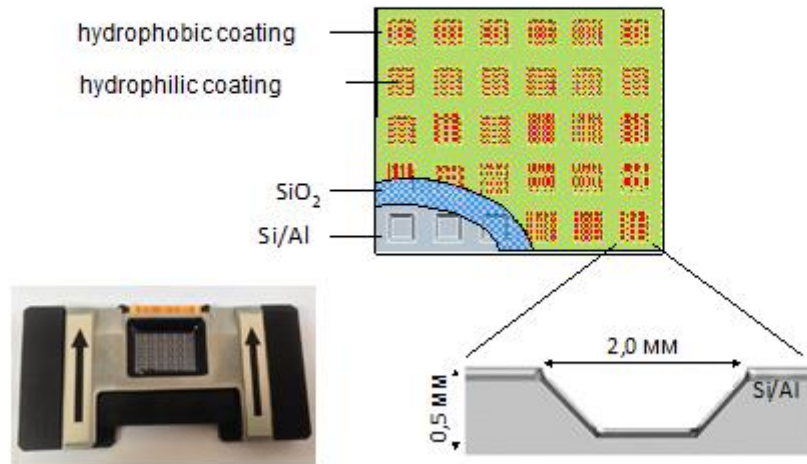
UPOV TECHNICAL WORKING PARTY FOR BIOCHEMICAL AND
MOLECULAR TECHNIQUES, AND DNA-PROFILING IN PARTICULAR
(BMT)

Fifteenth Session, Moscow, Russia, May from 24 to 27, 2016

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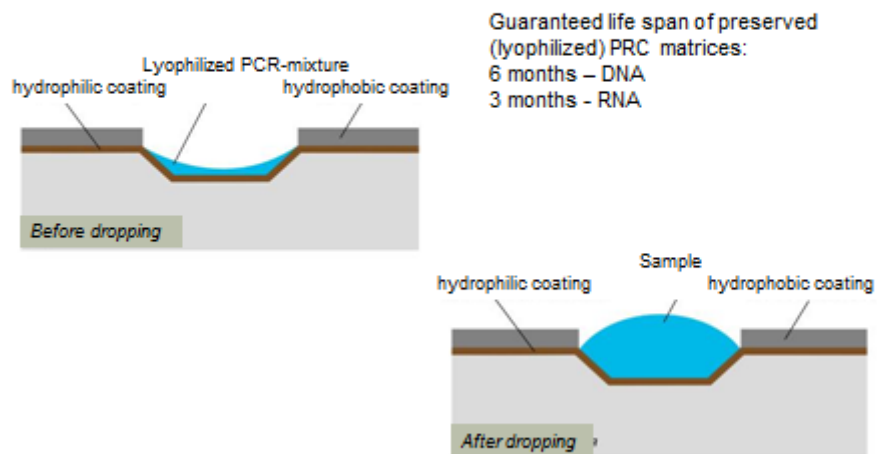
PCR-matrix



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PCR-matrix

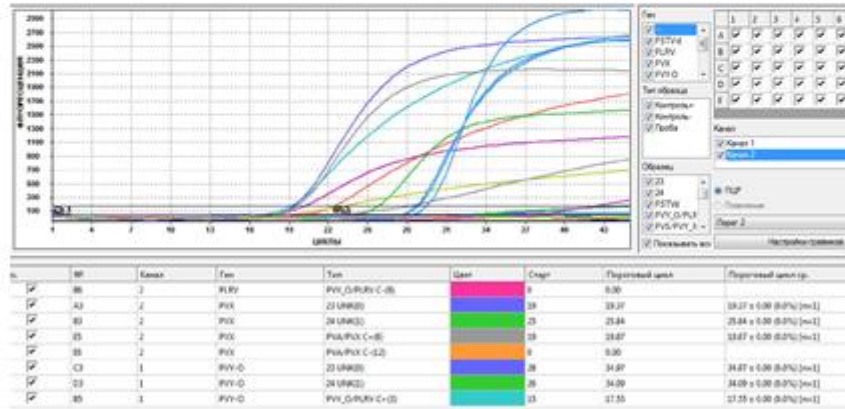


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Interpretation of results

Results are analyzed with the authentic "AriaDNA" software



Proposed vs. Conventional

Proposed

- Target: crop
- Multiple objects in a single test
- Time required: ~2 hrs (including DNA/RNA isolation)
- Could be used "anywhere"
- No need in highly trained personnel
- "Pre-serves" for a chosen range of pathogens that could be kept for ~6 months under room conditions

Conventional

- Target: object/pathogen
- Separate test for a single pathogen
- Time required: > 1 day
- Stationary conditions
- Does require highly trained personnel
- Reaction mixture for each object/pathogen



S.W.O.T.

S - Strength

- User friendliness
- Multiple objects in a single test
- Speed
- High sensitivity and accuracy
- Mobility
- End-users can easily develop their own applications

W - Weakness

- High qualification required for development of the test systems
- Not approved yet by international standards

O - Opportunities

- Use for screening and monitoring "anywhere"
- Use for seed quality assessment and for IPR protection
- End-users can easily develop their own applications

T - Threats

- Possible unpreparedness of the society to instantly accept new, differing significantly from traditional approaches



ISO 9001:2008
№ Q-04.04.18f
from 12.02.2013



Potato - available matrices

| DNA | RNA | Soil |
|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <ul style="list-style-type: none"> • <i>Clavibacter michiganensis</i> subsp. <i>sepedonicus</i> • <i>Pectobacterium atrosepticum</i> • <i>Dickeya dianthicola</i> • <i>Erwinia carotovora</i> subsp. <i>atroseptica</i> • <i>Ralstonia solanacearum</i> • <i>Phytophthora infestans</i> | <ul style="list-style-type: none"> • PLRV • PVY-o+c • PVY-ntv • PVX • PVA • PVM • PVS • PMTV • PSTVd | <ul style="list-style-type: none"> • <i>Globodera rostochiensis</i> • <i>Globodera pallida</i> • Phytoplasma: <ul style="list-style-type: none"> • Aster yellows (16 Sr I) • X-disease (16 Sr III) • Clover proliferation (16 Sr VI) |

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Potato – sampling



Viruses

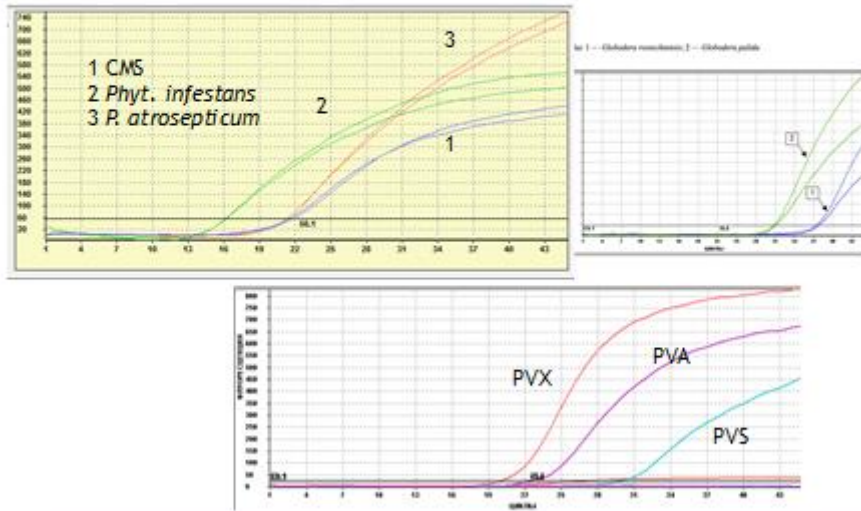
Bacteria



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Potato – PCR-matrices

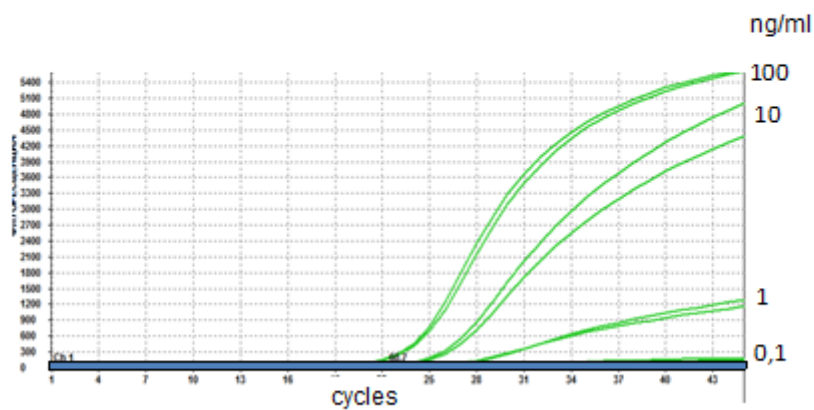


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Potato - PCR-matrices

Analytical sensitivity



12



Potato – PCR-matrices, field tests

Together with the Russian Agricultural Center, Leningrad region, Shushary, 2014

Samples: foliage and stems, 10 potato varieties

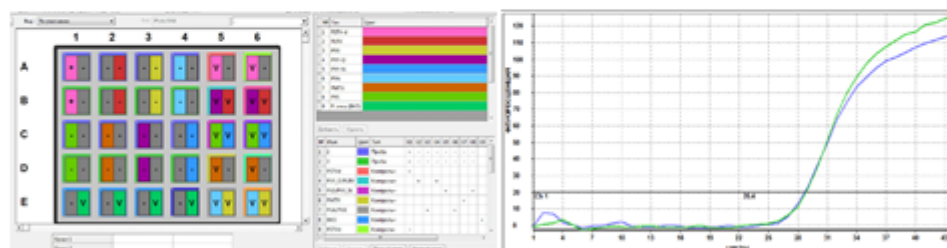
| VARIETY | DESCRIPTION | ELISA | GenBit - (rt) qPCR |
|----------|---------------------------|-------|----------------------------------------------|
| Nevskiy | Suspected: <i>Dickeya</i> | ??? | <i>Erwinia carotovora subsp. atroseptica</i> |
| Nevskiy | Suspected: Y-virus | - | PVY (o) |
| Avrora | Suspected: Y-virus | - | PVY (o+n) |
| Impala | Striated veins | PVM | PVM + PVY (o) |
| Asterisk | Spotted leaves | - | PVY (o+n) |
| Lausnak | Suspected: X-virus | PVX | PVX + PVY (o) |

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Potato – PCR-matrices, PSTV-d

Two samples by 150 mg each were taken from eyes (slices) of two mini-tuber suspected of being infected with the viroid



template

presence of PSTV-d in both samples (~0.5 ng/ml)

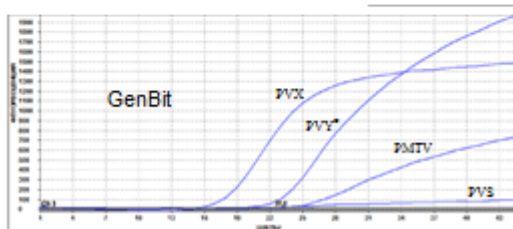
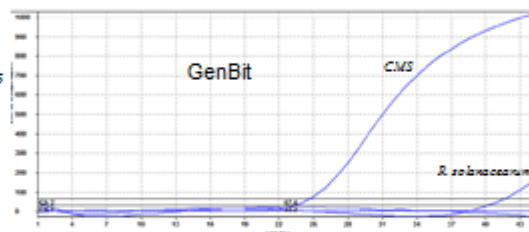
14



Potato – PCR-matrices, multiple pathogens

17.02.2015 Federal Enterprise "The All-Russian Center for Plant Quarantine" (VNIKR), "blind" samples, mix of DNA/RNA of bacterial and viral plant pathogens

VNIKR:
Clavibacter michiganensis subsp. *Sepedonicus*
+
Ralstonia solanacearum



VNIKR:
PMTV+PVS+PVY+PVX

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GMO

[EU Database of Reference Methods for GMO Analysis](#)

[JRC EU: Compendium of reference methods for GMO analysis](#)

[The CropLife International Detection Methods Database](#)

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GM-soybean, element specific detection

GM-soybean lines approved for FFP in the Russian Federation

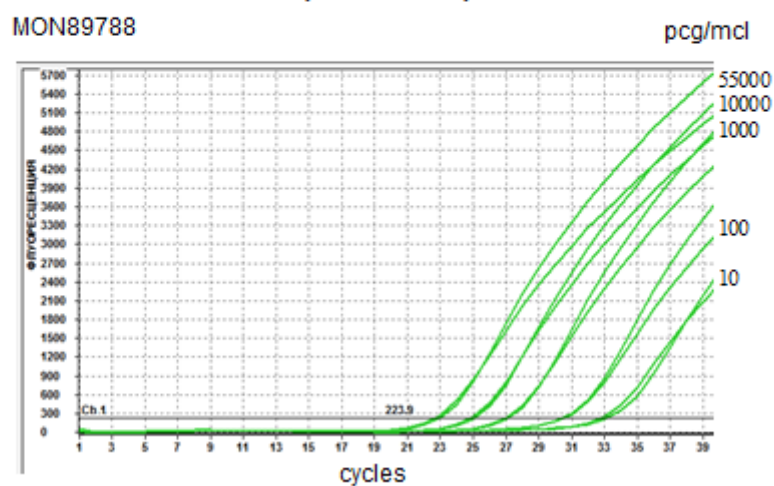
| LINE | CaMV P-35S | P-FMV | T-nos | npt II | bar |
|-------------|------------|-------|-------|--------|-----|
| A 2704-12 | + | - | - | - | - |
| A 5547-127 | + | - | - | - | - |
| BPS-CV127-9 | - | - | - | - | - |
| GTS-40-3-2 | + | - | + | - | - |
| MON87701 | - | - | - | - | - |
| MON89788 | - | + | - | - | - |
| SYHTOH2 | - | - | - | - | - |
| FG72 | - | - | - | - | - |

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GM-soybean, PCR-matrices, event specific

Analytical sensitivity



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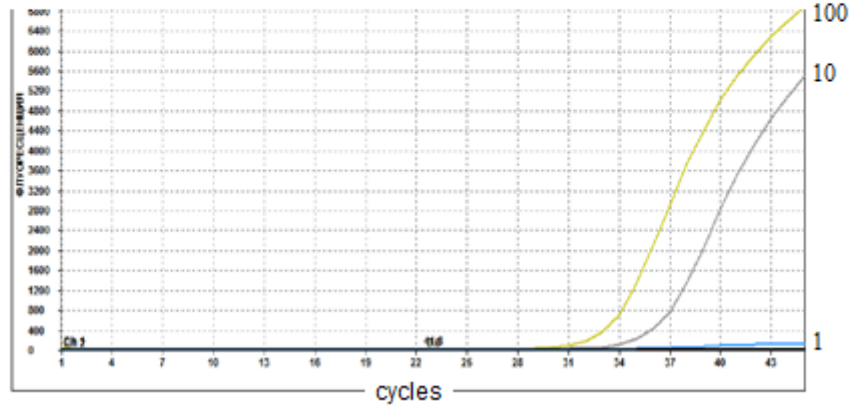


GM-soybean, PCR-matrices, event specific

Analytical sensitivity

SYHT0H2

pcg/mcl



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GM-soybean, PCR-matrices, event specific



20



GM-soybean, PCR-matrices, event specific

17.02.2015 Federal Enterprise "Centre of Molecular Diagnostics (CMD) - The All-Russian State Center for Quality and Standardization of Veterinary Drugs and Feed (VGNI)", three "blind" samples, mix of DNA of 7 GM-soybean lines each, one matrix

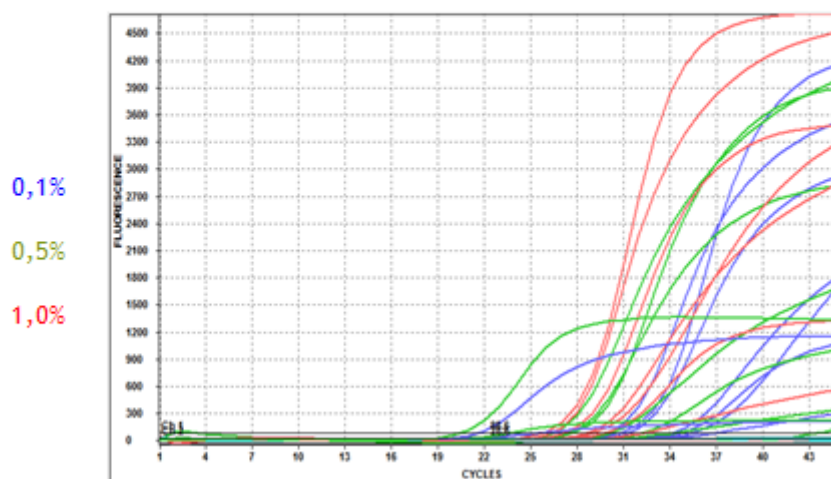
| 0.1% | | 0.5% | | 1.0% | |
|-------------|--------|-------------|--------|-------------|--------|
| LINE | GenBit | LINE | GenBit | LINE | GenBit |
| A 2704-12 | + | A 2704-12 | + | A 2704-12 | + |
| A 5547-127 | + | A 5547-127 | + | A 5547-127 | + |
| BPS-CV127-9 | + | BPS-CV127-9 | + | BPS-CV127-9 | + |
| GTS-40-3-2 | + | GTS-40-3-2 | + | GTS-40-3-2 | + |
| MON87701 | + | MON87701 | + | MON87701 | + |
| MON89788 | + | MON89788 | + | MON89788 | + |
| SYHTOH2 | + | SYHTOH2 | + | SYHTOH2 | + |

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GM-soybean, PCR-matrices, event specific

17.02.2015 CMD-VGNI: A2704-12+A5547-127+BPS-CV127-9+GTS-40-3-2+MON87701+MON89788+SYHTOH2



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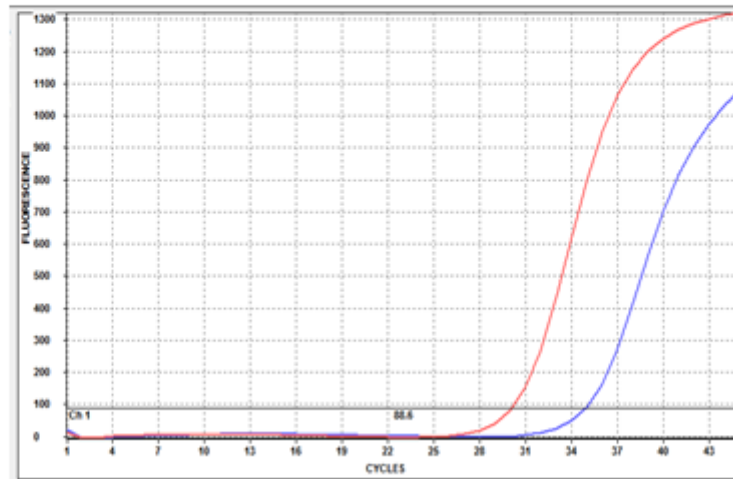


GM-soybean, PCR-matrices, event specific

17.02.2015 CMD-VGNKI: A2704-12 - Ct 30,12 (1%) Ct 34,99 (0,1%)

0,1%

1,0%



23

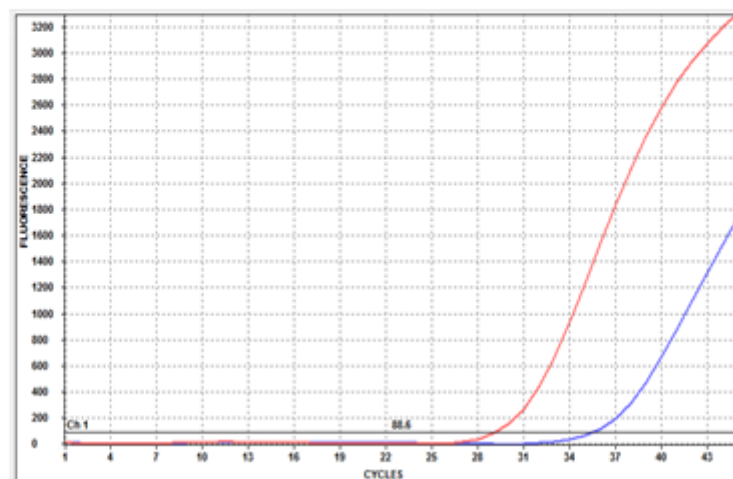


GM-soybean, PCR-matrices, event specific

17.02.2015 CMD-VGNKI: A5547-127 - Ct 29,24 (1%) Ct 30,57 (0,1%)

0,1%

1,0%



24

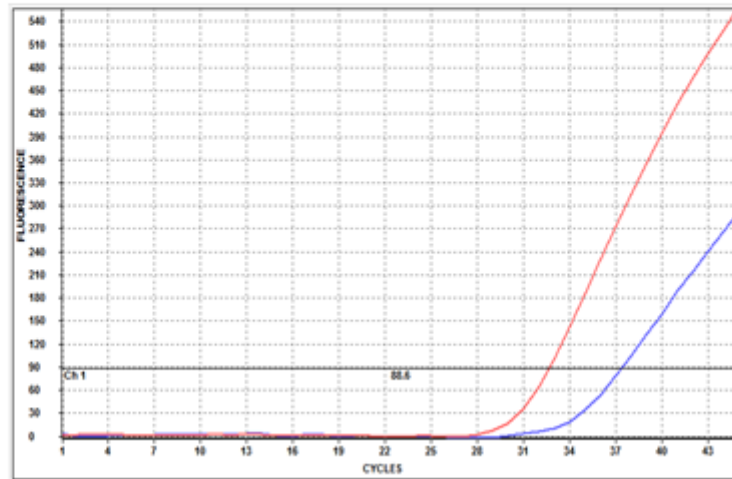


GM-soybean, PCR-matrices, event specific

17.02.2015 CMD-VGNKI: BPS-CV-127 Ct 32,71 (1%) Ct 37,43 (0,1%)

0,1%

1,0%



25

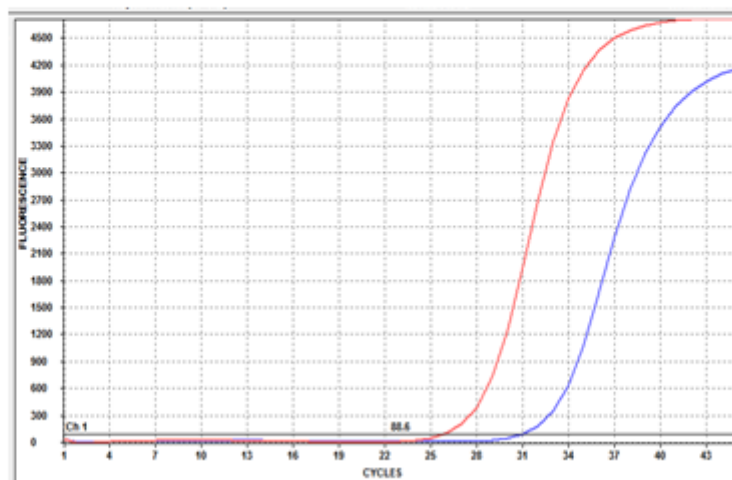


GM-soybean, PCR-matrices, event specific

17.02.2015 CMD-VGNKI: GTS 40-3-2 Ct 26,01 (1%) Ct 31,08 (0,1%)

0,1%

1,0%



26

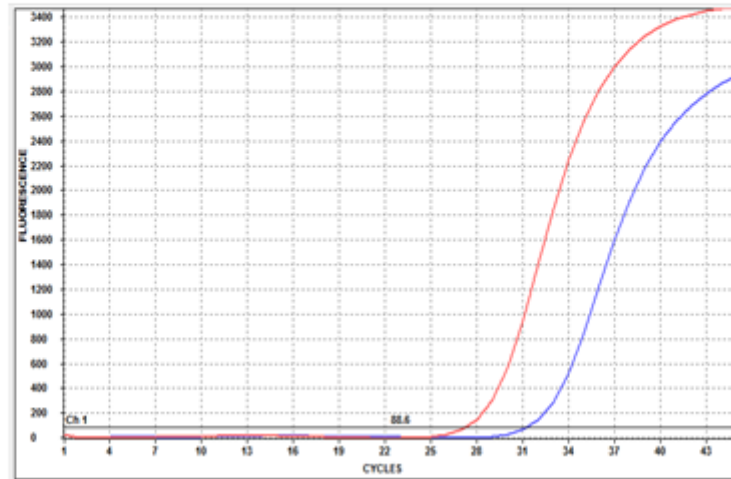


GM-soybean, PCR-matrices, event specific

17.02.2015 CMD-VGNKI: MON87701 Ct 27,36 (1%) Ct 31,41 (0,1%)

0,1%

1,0%



27

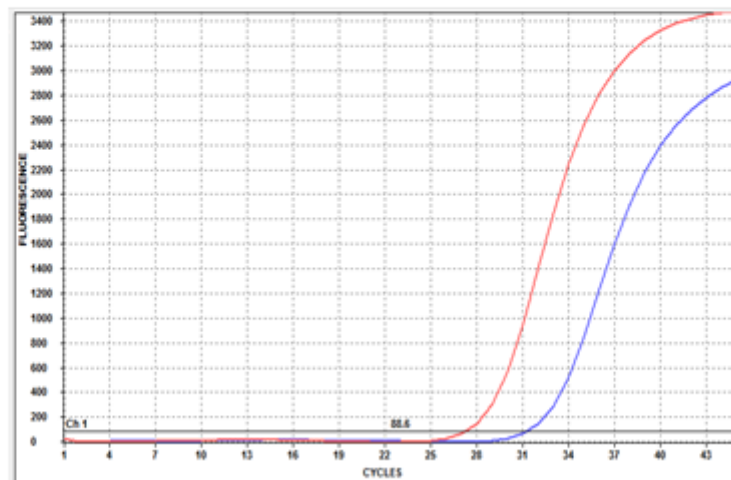


GM-soybean, PCR-matrices, event specific

17.02.2015 CMD-VGNKI: MON89788 Ct 28,95 (1%) Ct 34,39 (0,1%)

0,1%

1,0%



28

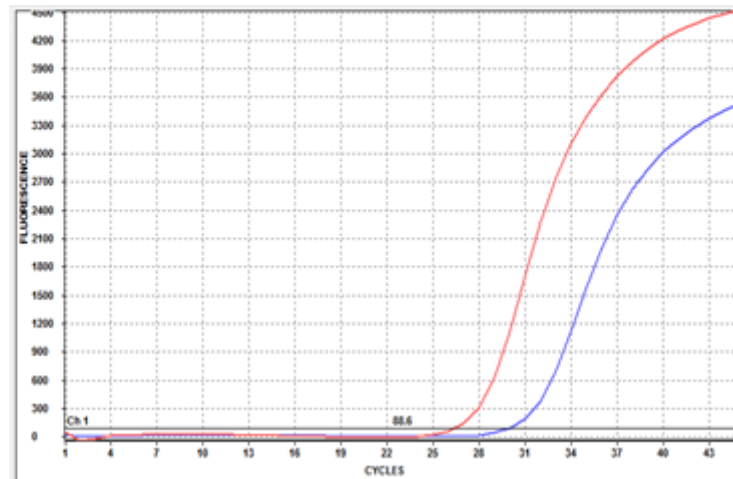


GM-soybean, PCR-matrices, event specific

17.02.2015 CMD-VGNKI: SYHTOH2 Ct 26,50 (1%) Ct 30,10 (0,1%)

0,1%

1,0%



28



GM-soybean, PCR-matrices, event specific (pattern)

17.02.2015 CMD-VGNKI: A2704-12+A5547-127+BPS-CV127-9+GTS-40-3-2+MON87701+MON89788+SYHTOH2

View: Default

File: (NO) .xml

1 2 3 4 5 6

A

B

C

D

E

| # | Gene | Color |
|---|------------|------------|
| 1 | SYH | Blue |
| 2 | Let | Red |
| 3 | MON89788 | Green |
| 4 | MON87701 | Yellow |
| 5 | GTS-40-3-2 | Purple |
| 6 | A2704 | Orange |
| 7 | A5547 | Light Blue |
| 8 | BPS-CV-127 | Dark Blue |

Add

Remove

| # | Name | Color | Type | G1 | G2 | G3 | G4 | G5 | G6 | G7 | G8 | Description |
|---|---------|------------|---------|----|----|----|----|----|----|----|----|-------------|
| 1 | R1 | Blue | Unknown | * | * | * | * | * | * | * | * | |
| 2 | R5 | Green | Unknown | * | * | * | * | * | * | * | * | |
| 3 | I | Red | Unknown | * | * | * | * | * | * | * | * | |
| 4 | Control | Light Blue | Control | - | - | - | - | - | - | - | - | |

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GM-soybean, PCR-matrices, event specific

19.02.2015 Federal Enterprise "Institute of Nutrition", three "blind" samples, DNA mix, one matrix:
MON89788+SYHT0H2+A2704 (№2)
BPS CV-127+SYHT0H2 (№4)
BPS CV-127+SYHT0H2+GTS 40-3-2 (№6)

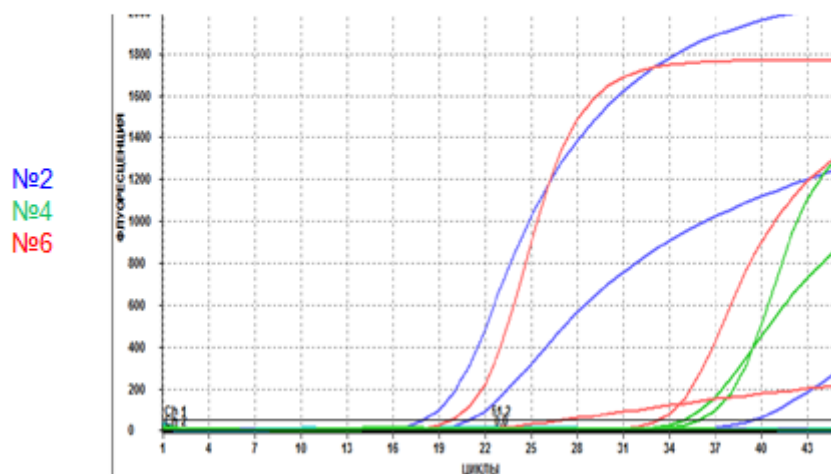
| №2 | | №4 | | №6 | |
|-------------|--------|-------------|--------|-------------|--------|
| LINE | GenBit | LINE | GenBit | LINE | GenBit |
| A 2704-12 | + | A 2704-12 | - | A 2704-12 | - |
| A 5547-127 | - | A 5547-127 | - | A 5547-127 | - |
| BPS-CV127-9 | - | BPS-CV127-9 | + | BPS-CV127-9 | + |
| GTS-40-3-2 | - | GTS-40-3-2 | - | GTS-40-3-2 | + |
| MON87701 | - | MON87701 | - | MON87701 | - |
| MON89788 | + | MON89788 | - | MON89788 | - |
| SYHT0H2 | + | SYHT0H2 | + | SYHT0H2 | + |

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GM-soybean, PCR-matrices, event specific

19.02.2015 Federal Enterprise "Institute of Nutrition", three "blind" samples, DNA mix, one matrix:



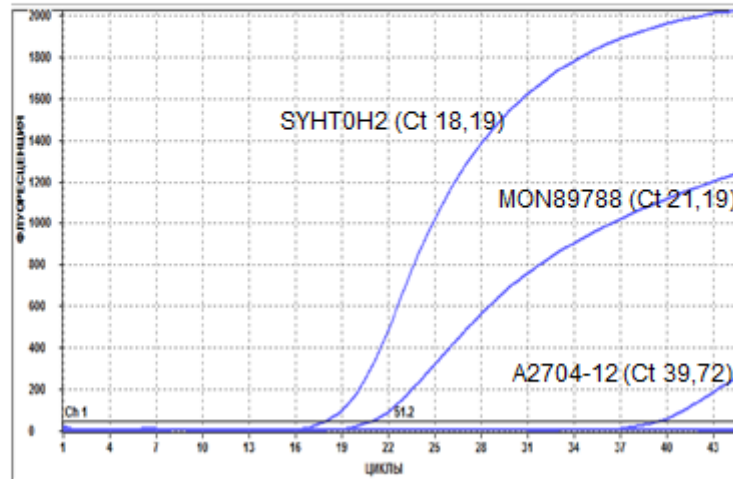
92



GM-soybean, PCR-matrices, event specific

19.02.2015 Federal Enterprise "Institute of Nutrition", three "blind" samples, DNA mix, one matrix:

№2



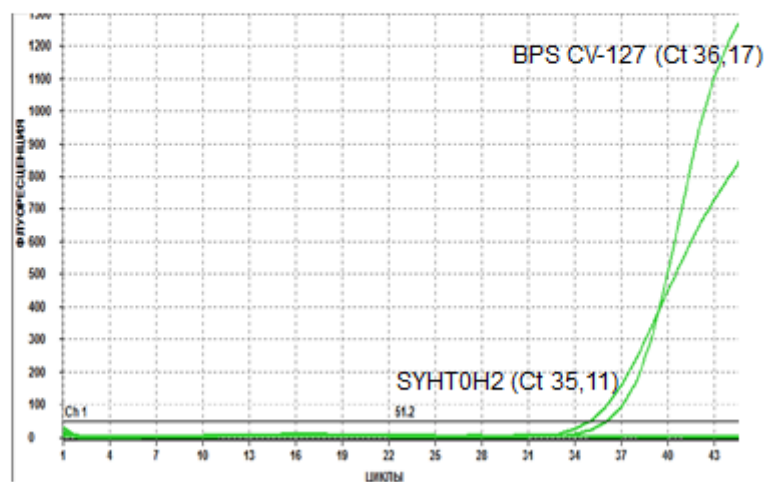
22



GM-soybean, microchips, event specific

19.02.2015 Federal Enterprise "Institute of Nutrition", three "blind" samples, DNA mix, one matrix:

№4



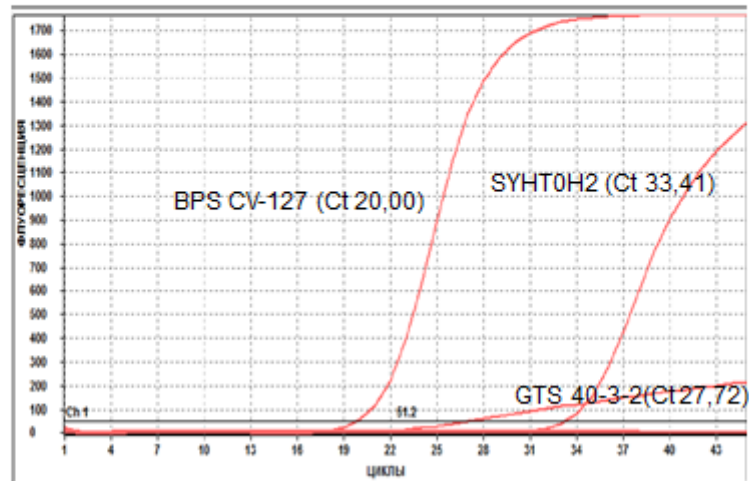
24



GM-soybean, microchips, event specific

19.02.2015 Federal Enterprise "Institute of Nutrition", three "blind" samples, DNA mix, one matrix:

№6

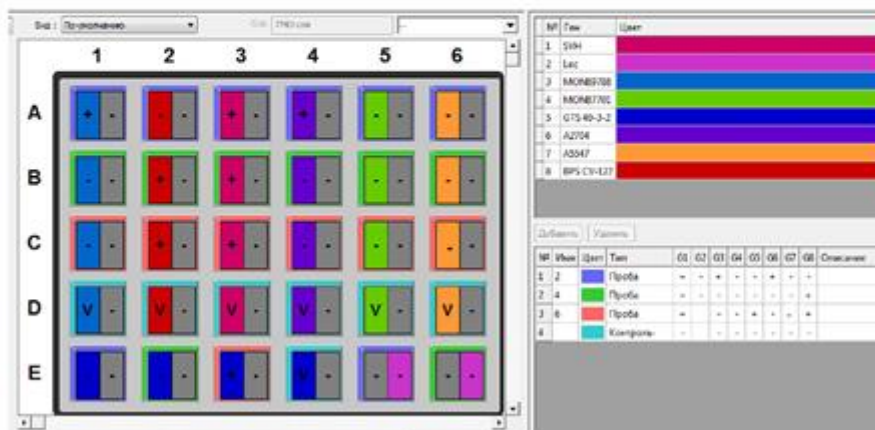


25

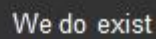


GM-soybean, microchips, event specific (pattern)

19.02.2015 Federal Enterprise "Institute of Nutrition", three "blind" samples, DNA mix, one matrix:



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THANK YOU!



[End of Annex and of document]