

# Exporting results of 35S/NOS/IPC from ABI 7500 FAST to GeneScan Evaluation sheet

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**GMO Testing Facility**

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FREISTATA



**UFS·UV**  
HEALTH SCIENCES  
GESONDHEIDSWETENSAPPE

**Run**

**Analysis**

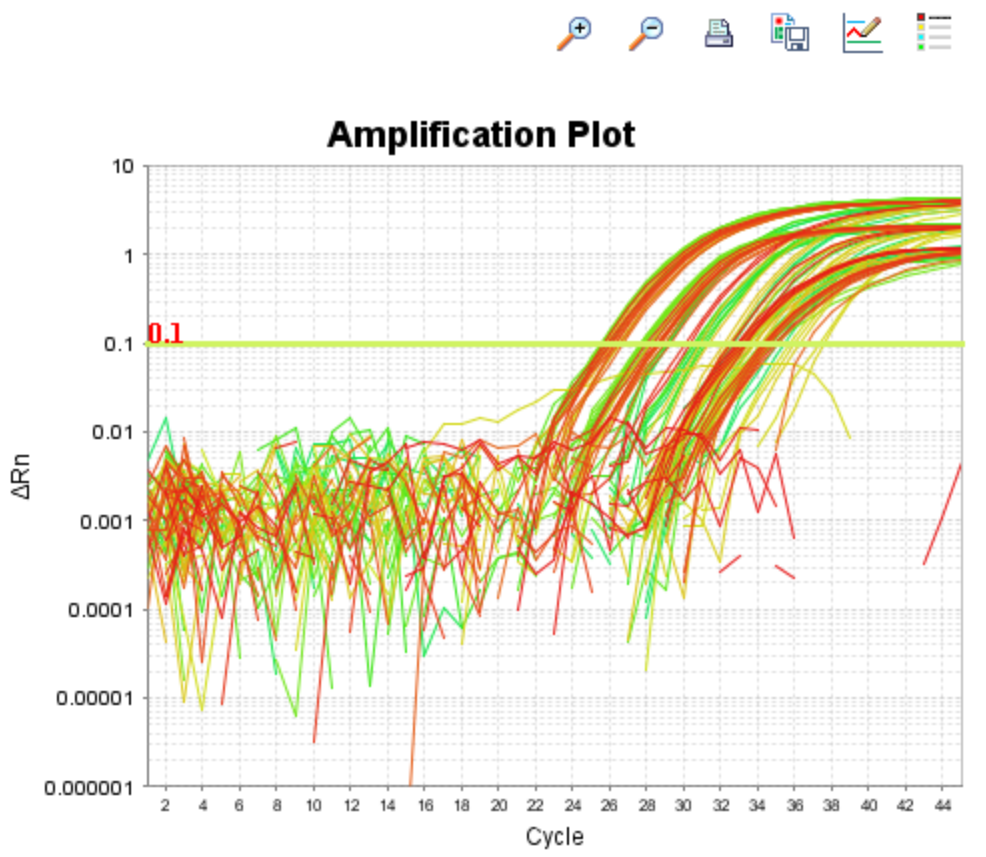
- Amplification Plot
- Standard Curve
- Multicomponent Plot
- Raw Data Plot
- QC Summary
- Multiple Plots View

**Amplification Plot**

Plot Settings

Plot Type:  $\Delta Rn$  vs Cycle | Graph Type: Log | Plot Color: Well

Save current settings as the default



Legend

A B C D E F G H

Options

Target: All | Threshold:  Auto |  Auto Baseline

Show:  Threshold —  Baseline Start: Well  Target ▲  Baseline End: Well ■

Save current settings as the default

View Plate Layout | View Well Table

Select Wells With: - Select Item - | - Select Item -

Show in Wells | View Legend

	1	2	3	4	5	6	7	8	9	10	11	12
A	1 N 50 CT. 3	1 N 50 CT. 3	1 U 5... CT. 3	1 U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	1 U 5... CT. 3	1 U 5... CT. 3
B	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	1 U 5... CT. 3	1 U 5... CT. 3	2 U 5... CT. 3	1 U 5... CT. 3	U 5... CT. 3	U 5... CT. 3
C	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3
D	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3	U 5... CT. 3								
E												
F												
G												

1. Select all wells
2. Under Target: Select "All"





Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

## Export Properties

## Customize Export

1. Select data to export:
- Sample Setup
  - Results
  - Raw Data
  - Multicomponent Data
  - Amplification Data
- Ensure that the "Results" box is selected**
- Include Results in export**

2. Select one file or separate files:  *Select to export all data in one file or in separate files for each data type.*

3. Enter export file properties:

Export File Name:  File Type:

Export File Location:

Open file(s) when export is complete

Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

Export Properties

Customize Export

1. Select data to export:
- Sample Setup
  - Results
  - Raw Data
  - Multicomponent Data

Amplification Data **Select "Amplification Data" box**

Include Amplification Data in export

2. Select one file or separate files: One File *Select to export all data in one file or in separate files for each data type.*

3. Enter export file properties:

Export File Name: Double Screen\_data File Type: (\*.xls)

Export File Location: C:\Applied Biosystems\7500\experiments

Open file(s) when export is complete

**I** Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

Export Properties

Customize Export

1. Select data to export:

- Sample Setup
- Results
- Raw Data
- Multicomponent Data
- Amplification Data

2. Select one file or separate files:

One File  
One File  
Separate Files

Select to export all data in one file or in separate files for each data type.

3. Enter export file properties:

Export File Name: Double Screen\_data File Type: (\*.xls)

Export File Location: C:\Applied Biosystems\7500\experiments

Open file(s) when export is complete

**Select "Separate Files"**

Setup

Run

Analysis

Amplification Plot

Standard Curve

Multicomponent Plot

Raw Data Plot

QC Summary

Multiple Plots View

the export format and to select fields to export. Click "Start Export" to export your data.

Export Properties    Customize Export


Sample Setup     Results  
 Raw Data     Multicomponent Data  
 Amplification Data

1. Select data to export:

2. Select one file or separate files: **Separate Files**    *Select to export all data in one file or in separate files for each data type.*


3. Enter export file properties:

**Results**

Export File Name:     File Type:  

Export File Location:    

**Amplification Data**

Export File Name:     File Type:  

Export File Location:    

Open file(s) when export is complete

Save current settings as the default

em -

	10	11	12
5..	U 5..	1	1
2	U 5..	U 5..	U 5..
3..	CT. 3..	CT. 3..	CT. 3..
5	U 5..	U 5..	U 5..
50	1	U 5..	U 5..
5..	U 5..	U 5..	U 5..
3..	CT. 3..	CT. 3..	CT. 3..
5..	U 5..	U 5..	U 5..
20	CT. 2	CT. 2	CT. 2
3..	CT. 3..	CT. 3..	CT. 3..
5	U 5..	U 5..	U 5..

Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

Export Properties Customize Export

- Sample Setup
- Results
- Raw Data
- Multicomponent Data
- Amplification Data

2. Select one file or separate files: Separate Files Select to export all data in one file or in separate files for each data type.

3. Enter export file properties:

**Results**

Export File Name: 07-11-2019-RESULTS File Type: (\*.xls)

Export File Location: C:\Applied Biosystems\7500\experiments Browse

**Amplification Data**

Export File Name: Double Screen\_amplificationdata File Type: (\*.xls)

Export File Location: C:\Applied Biosystems\7500\experiments Browse

Open file(s) when export is complete

Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

## Export Properties

## Customize Export

1. Select data to export:
- Sample Setup
  - Results
  - Raw Data
  - Multicomponent Data
  - Amplification Data

2. Select one file or separate files:  *Select to export all data in one file or in separate files for each data type.*

3. Enter export file properties:

## Results

Export File Name:  File Type:

Export File Location:

## Amplification Data

Export File Name:  File Type:

Export File Location:

Open file(s) when export is complete

**I** Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

## Export Properties

## Customize Export

1. Select data to

2. Select one file

3. Enter export

### Results

Export File Name

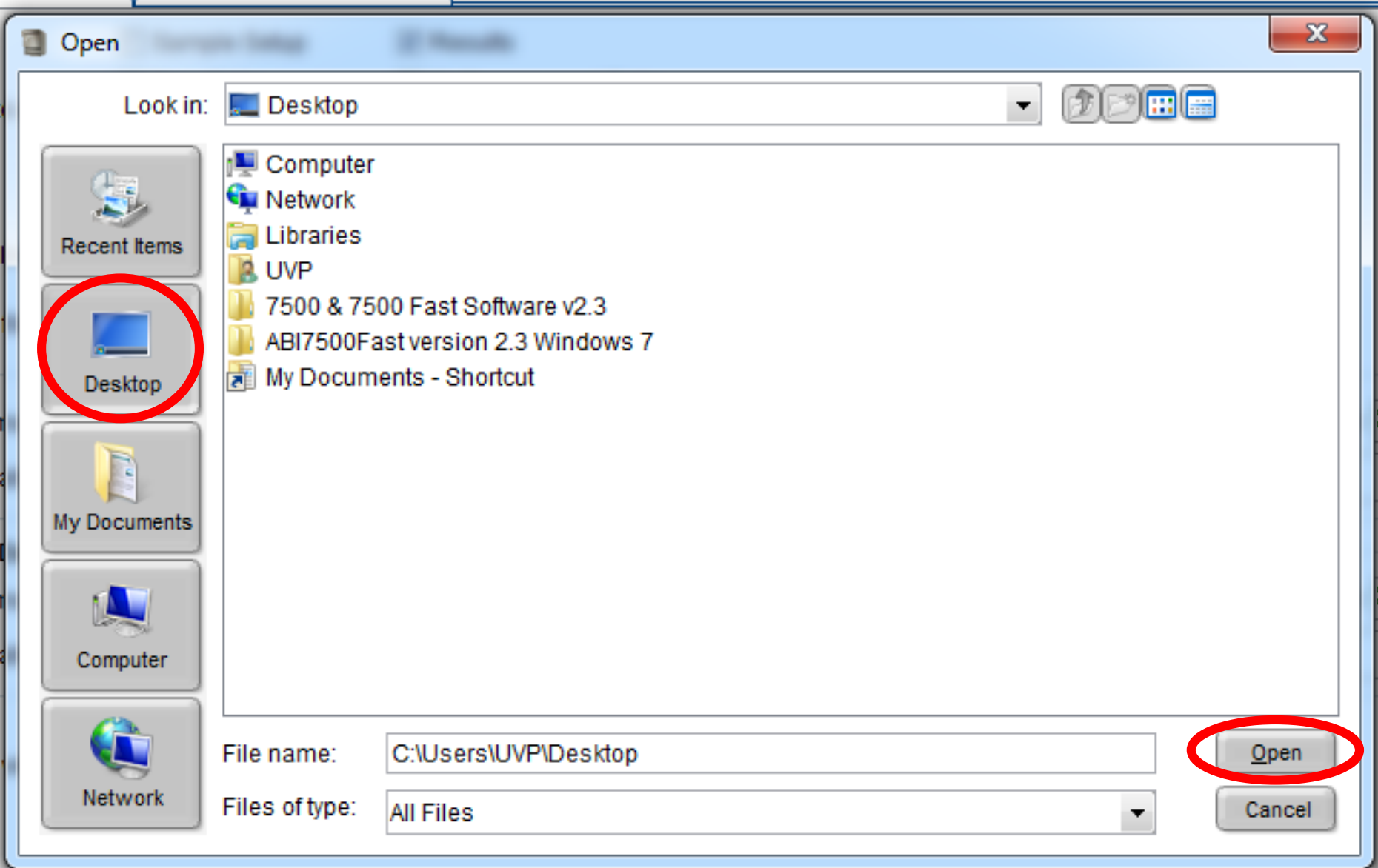
Export File Location

### Amplification

Export File Name

Export File Location

Open file(s)



(\*.xls)

Browse

(\*.xls)

Browse

Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

Export Properties Customize Export

1. Select data to export:
- Sample Setup
  - Results
  - Raw Data
  - Multicomponent Data
  - Amplification Data

2. Select one file or separate files: Separate Files Select to export all data in one file or in separate files for each data type.

3. Enter export file properties:

**Results**

Export File Name: 07-11-2019-RESULTS File Type: (\*.xls)


Export File Location: C:\Users\UVP\Desktop Browse

**Amplification Data**

Export File Name: Double Screen\_amplificationdata File Type: (\*.xls)

Export File Location: C:\Applied Biosystems\7500\experiments Browse

Open file(s) when export is complete

**I** Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data. 

## Export Properties

## Customize Export

1. Select data to export:
- Sample Setup
  - Results
  - Raw Data
  - Multicomponent Data
  - Amplification Data

2. Select one file or separate files:  *Select to export all data in one file or in separate files for each data type.*


3. Enter export file properties:

### Results

Export File Name:  File Type:  (\*.xls)

Export File Location:

### Amplification Data

Export File Name:  File Type:  (\*.xls)

Export File Location:

Open file(s) when export is complete

Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

Export Properties Customize Export

1. Select data to export:
- Sample Setup
  - Results
  - Raw Data
  - Multicomponent Data
  - Amplification Data

2. Select one file or separate files: Separate Files *Select to export all data in one file or in separate files for each data type.*

3. Enter export file properties:

**Results**

Export File Name:  File Type: Excel (\*.xls)

Export File Location:  Browse

**Amplification Data**

Export File Name:  File Type: Excel (\*.xls)

Export File Location:  Browse

Open file(s) when export is complete

Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

Export Properties

Customize Export

1. Select data to export:
- Sample Setup
  - Results
  - Raw Data
  - Multicomponent Data
  - Amplification Data

2. Select one file

3. Enter export

Results

Export File Name

Export File Location

Amplification

Export File Name

Export File Location

Open file(s)

Open

Look in: Desktop

- Computer
- Network
- Libraries
- UVP
- 7500 & 7500 Fast Software v2.3
- ABI7500Fast version 2.3 Windows 7
- My Documents - Shortcut

Recent Items

Desktop

My Documents

Computer

Network

File name: C:\Users\UVP\Desktop

Files of type: All Files

Open

(\*.xls)

Browse

(\*.xls)

Browse



Open selected file

2. Select one file or separate files:  *Select to export all data in one file or in separate files for each data type.*

3. Enter export file properties:

**Results**

Export File Name:  File Type:  (\*.xls)

Export File Location:

**Amplification Data**

Export File Name:  File Type:  (\*.xls)

Export File Location:

Open file(s) when export is complete

Save current settings as the default



Select the type of data to export, select whether to export one file or separate files, then enter export file properties. (Optional) Click "Customize Export" to change the export format and to select fields to export. Click "Start Export" to export your data.

## Export Properties

## Customize Export

1. Select data to export:
- |                                                        |                                              |
|--------------------------------------------------------|----------------------------------------------|
| <input type="checkbox"/> Sample Setup                  | <input checked="" type="checkbox"/> Results  |
| <input type="checkbox"/> Raw Data                      | <input type="checkbox"/> Multicomponent Data |
| <input checked="" type="checkbox"/> Amplification Data |                                              |

2. Select one file or separate files:  *Select to export all data in one file or in separate files for each data type.*

3. Enter export file properties:

## Results

Export File Name:

Export File Location:

## Amplification Data

Export File Name:

Export File Location:

Open file(s) when export is complete

## Export Completed



Your files have been exported to:

Results exported to C:\Users\UVP\Desktop\07-11-2019-RESULTS.xls

Amplification Data exported to C:\Users\UVP\Desktop\07-11-2019-AMPLIFICATION.xls

What do you want to do next?

File Type:

File Type:

- 1. Open the GeneScan Evaluation Sheet**
- 2. Enable editing content / macros as required**
- 3. Select -----> File -----> Save As**
- 4. Save the Excel File using the same name as the Run File**

FILE HOME INSERT PAGE LAYOUT FORMULAS DATA REVIEW VIEW

Clipboard Font Alignment Number Styles Cells Editing

fx GS-P-09.017

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**Macros**

For the function of this sheet macros has to be enabled. Please refer to the links below how to enable macros in Excel sheet, how to change the security settings and how to add, remove or modify a trusted location in Excel 2010.

- [Enable or disable macros in Office](#)
- [Change macro security settings in Office](#)
- [Add, remove or modify a trusted location](#)

**Reporter / Quencher**

Please use for Template-Setup, RealTime PCR Setup etc. the below mentioned Reporter/Quencher Combinations and the recommended Assay-/Detector-Names. If you use other notations (than the mentioned below or in Validation data sheet) for RealTime PCR setup the correct function of this Evaluation sheet couldn't ensure. For Reporter/Quencher Combinations and Assay-/Detector-Names of modules not shown in the tables below please refer to the Validation data sheet submitted with the release of each module or refer to the manual.

Module	GS-P-09.017	Dye:	1	2	IPC
System	GS-P-09.038 event MON89034		1735S	17nos	17IPC
Instrument	GS-P-09.039 bar-gene		FAM	JOE	Cy5
# NTC	GS-P-09.040 event MON88017		NFQ	NFQ	NFQ
# Pos. Control	GS-P-09.041 event 3272				
	GS-P-09.043 eventH7-1				
	GS-P-09.044 event305423				
	GS-P-09.045 nos/DHFR1-T/G				
	GS-P-09.046 Tobacco NIR2				
	GS-P-09.047 Cotton SAH7				

**Platelayou**

Module	GS-P-09.050	35S/NOS V2
System	GS-P-09.052 event MON89034	
Instrument	GS-P-09.053 eventDP098140-6	
# NTC	GS-P-09.055 event KMD1	
# Pos. Control	GS-P-09.061 35S/NOS/FMV	
	GS-P-09.063 CV127-HT-G/G	

	1	2	3	4	5	6	7	8	9	10	11	12
A	N	N	N	N	N	N	P	P	P	P	U	U
B	U	U	U	U	U	U	U	U	U	U	U	U
C	U	U	U	U	U	U	U	U	U	U	U	U
D	U	U	U	U	U	U	U	U	U	U	U	U
E	U	U	U	U	U	U	U	U	U	U	U	U

	H	G	F	E	D	C	B	A
1	N	N	N	N	N	N	P	P
2	P	P	U	U	U	U	U	U
3	U	U	U	U	U	U	U	U
4	U	U	U	U	U	U	U	U
5	U	U	U	U	U	U	U	U

Module

GS-P-09.017

Dye:

	1	2
	1735S	17nos
	FAM	JOE
	NFQ	NFQ

System

GS-P-09.038 event MON89034

Instrument

GS-P-09.039 bar-gene

# NTC

GS-P-09.040 event MON88017

# Pos. Control

GS-P-09.041 event 3272

GS-P-09.043 eventH7-1

GS-P-09.044 event305423

GS-P-09.045 nos/DHFR1-T/G

GS-P-09.046 Tobacco NIR2

GS-P-09.047 Cotton SAH7

GS-P-09.048 event T45

Layout

GS-P-09.050 35S/NOS V2

GS-P-09.052 vip3Aa gene

12 Well format

GS-P-09.053 eventDP098140-6

GS-P-09.055 event KMD1

GS-P-09.061 35S/NOS/FMV

GS-P-09.063 CV127-HT-G/G

1

10

11

12

Module

GS-P-09.050

System

35S/NOS V2

Instrument

~~ABI 7500 SDS Vers 1.4~~

ABI 7500 SDS Vers 2.0

# NTC

Agilent AriaMX

not Validated for this Instrument

# Pos. Control

Agilent MX3005

Biorad CFX96

not Validated for this Instrument

Roche LightCycler480 I

0

not Validated for this Instrument

Platelayou


## Reporter / Quencher

Please use for Template-Setup, RealTime PCR Setup etc. the below mentioned Reporter/Quencher Combinations and the recommended Assay-/Detector-Names. If you use other notations (than the mentioned below or in Validation data sheet) for RealTime PCR setup the correct function of this Evaluation sheet couldn't ensure.

For Reporter/Quencher Combinations and Assay-/Detector-Names of modules not shown in the tables below please refer to the Validation data sheet submitted with the release of each module or refer to the manual.

Module	GS-P-09.050
System	35S/NOS V2
Instrument	ABI 7500 SDS Vers 2.0

# NTC	4
# Pos. Control	4

Dye:  
Reporter dye:  
Quencher dye:

1	2			IPC
5035S	50NOS			50IPC
FAM	JOE			Cy5
NFQ	NFQ			NFQ



36  
37  
38  
39  
40  
41  
42  
43  
44

12 Well format

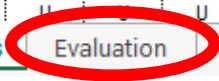
GS-P-09.053	eventDP098140-6
GS-P-09.055	event KMD1
GS-P-09.061	35S/NOS/FMV
GS-P-09.063	CV127-HT-G/G

	1	2	3	4	5	6	7	8	9	10	11	12
A	N	N	N	N	N	N	P	P	P	P	U	U
B	U	U	U	U	U	U	U	U	U	U	U	U
C	U	U	U	U	U	U	U	U	U	U	U	U
D	U	U	U	U	U	U	U	U	U	U	U	U
E	U	U	U	U	U	U	U	U	U	U	U	U

8 Well format

	H	G	F	E	D	C	B	A
1	N	N	N	N	N	N	P	P
2	P	P	U	U	U	U	U	U
3	U	U	U	U	U	U	U	U
4	U	U	U	U	U	U	U	U
5	U	U	U	U	U	U	U	U

Instructions Evaluation Change Log





FILE HOME INSERT PAGE LAYOUT FORMULAS DATA REVIEW VIEW

Clipboard Font Alignment Number Styles Cells Editing

Conditional Formatting Styles Insert Delete Format AutoSum Fill Clear Sort & Filter Find & Select

A B D E F CH I L M N O P Q R U V W X Y Z A AC AD AE AF AG AH AI AK AL AM AN AO AP AI AS AT

1 NO RealTime Data imported.

2 Module GS-P-09.050 Import RealTime PCR Data

3 System 35S/NOS V2

4 Instrument No Data imported No Data imported No Data imported No Data imported IPC

5 Layout ABI 7500 SDS Vers 1.4

6 Format ABI 7500 SDS Vers 2.0

- ABI 7900
- Agilent AriaMX not Validated for this Instrument
- Agilent MX3005
- Biorad CFX96 not Validated for this Instrument
- Roche LightCycler480 I not Validated for this Instrument
- Roche LightCycler480 II not Validated for this Instrument

10 Ct #N/A Out Ct #N/A Out Ct #N/A Out Ct #N/A Out Ct #N/A

11 Run No. Control

12 Lot MM/OM SD

13 Lot BM Delta

14 Lot C+ CutOff

15 Cycles 45

23 Comment: NTC Ct neg Inh pos Ct neg Inh pos Ct neg Inh pos Ct neg Inh pos C+

24 ExCtrl ExCtrl ExCtrl ExCtrl ExCtrl

25

26 Threshold

27

28 35S NOS 0 0 IPC

29 Nr Well Sample Dil. Task Out Ct #N/A Res Ct #N/A Res Ct #N/A Res Ct #N/A Res Ct #N/A

30 1

31 2

32 3

33 4

34 5

35 6

36 7

37 8

38



FILE HOME INSERT PAGE LAYOUT FORMULAS DATA REVIEW VIEW Sign in

Paste Font Alignment Number Styles Cells Editing

Clipboard Font Alignment Number Styles Cells Editing

D5

A B D E F G H I L M N O P Q R U V W X Y Z AA AS AT AU AV AW AX AY AZ

1 NO RealTime Data imported.

2 Module GS-P-09.050 Import RealTime PCR Data

3 System 35S/NOS V2

4 Instrument ABI 7500 SDS Vers 2.0 No Data imported No Data imported IPC | Cy5 | 50IPC

5 Layout

- Format
- 12 Wells Format | 6 NTC | 4 C+
- 12 Wells Format | 4 NTC | 4 C+
- 12 Wells Format | 2 NTC | 2 C+
- 8 Wells Format | 6 NTC | 4 C+
- 8 Wells Format | 4 NTC | 4 C+
- 8 Wells Format | 2 NTC | 2 C+
- Kingfisher 1 - 40 Samples

11 Run No. Control Ct dRN Out Ct dRN Out Ct dRN Out

12 Lot MM/OM SD

13 Lot BM Delta 7.00 0.20 7.00 0.20 4.00 0.33

14 Lot C+ CutOff

15 Cycles 45

21

22 Ct neg Inh pos Ct neg Inh pos Ct neg Inh pos

23 Comment: NTC ExCtrl C+ ExCtrl

24

25

26 Threshold

27

28 35S NOS IPC

29 Nr Well Sample Dil. Task Out Ct dRN Res Ct dRN Res Ct dRN Total

30 1

31 2

32 3

33 4

34 5

35 6

36 7

37 8

38 9

39 10

40 11

41 12

42 13

43 14

FILE HOME INSERT PAGE LAYOUT FORMULAS DATA REVIEW VIEW

Clipboard Font Alignment Number Styles Cells Editing

Calibri 9 B I U A A- A+ Wrap Text Merge & Center \$ % <-0 .00 >0.00 Conditional Formatting Format as Table Cell Styles Insert Delete Format AutoSum Fill Clear Sort & Filter Find & Select

D5

NO RealTime Data imported.

Module: GS-P-09.050  
 System: 35S/NOS V2  
 Instrument: ABI 7500 SDS Vers 2.0  
 Layout: [Dropdown]  
 Format: 12 Wells Format | 2 NTC | 2 C+ (circled in red)

Import RealTime PCR Data

No Data imported (Blue)

No Data imported (Green)

IPC | Cy5 | 50IPC (Pink)

Run No.	Ct	dRN	Out	Ct	dRN	Out	Ct	dRN	Out
Control									
SD									
Delta	7.00	0.20		7.00	0.20		4.00	0.33	
CutOff									

	Ct	neg	Inh	pos	Ct	neg	Inh	pos	C+	ExCtrl
NTC										
ExCtrl										

Threshold: [Input Box]

Nr	Well	Sample	Dil.	Task	Out	35S			NOS			IPC			
						Ct	dRN	Res	Ct	dRN	Res	Ct	dRN	Total	
1															
2															
3															
4															
5															
6															
7															
8															
9															
10															
11															
12															
13															
14															

FILE HOME INSERT PAGE LAYOUT FORMULAS DATA REVIEW VIEW

Clipboard Font Alignment Number Styles Cells Editing

Calibri 9 A<sup>+</sup> A<sup>-</sup> B I U Wrap Text Merge & Center \$ % ←0 →0

Conditional Formatting Format as Table Cell Styles Insert Delete Format AutoSum Fill Clear Sort & Filter Find & Select

D6

A B D E F G H I L M N O P Q R U V W X Y Z AA AS AT AU AV AW AX AY AZ

NO RealTime Data imported.

2 Module GS-P-09.050 Import RealTime PCR Data

3 System 35S/NOS V2

4 Instrument ABI 7500 SDS Vers 2.0 No Data imported No Data imported IPC | Cy5 | 50IPC

5 Layout 12 Wells Format | 2 NTC | 2 C+

6 Format 12 Wells Format 8 Wells Format

	Ct	dRN	Out	Ct	dRN	Out	Ct	dRN	Out
11 Run No.									
12 Lot MM/OM									
13 Lot BM	7.00	0.20		7.00	0.20		4.00	0.33	
14 Lot C+									
15 Cycles	45								

	Ct	neg	Inh	pos	Ct	neg	Inh	pos	Ct	neg	Inh	pos
23 Comment:												
24 NTC												
24 ExCtrl												
26 Threshold												

Nr	Well	Sample	Dil.	Task	Out	35S			NOS			IPC			
						Ct	dRN	Res	Ct	dRN	Res	Ct	dRN	Total	
30	1														
31	2														
32	3														
33	4														
34	5														
35	6														
36	7														
37	8														
38	9														
39	10														
40	11														
41	12														
42	13														

FILE HOME INSERT PAGE LAYOUT FORMULAS DATA REVIEW VIEW

Clipboard Font Alignment Number Styles Cells Editing

Calibri 9 A<sup>+</sup> A<sup>-</sup> B I U Wrap Text Merge & Center \$ % <-0 ->0

Conditional Formatting Format as Table Cell Styles Insert Delete Format AutoSum Fill Clear Sort & Filter Find & Select

D6

A B D E F G H I L M N O P Q R U V W X Y Z AA AS AT AU AV AW AX AY AZ

NO RealTime Data imported.

Module: GS-P-09.050

System: 35S/NOS V2

Instrument: ABI 7500 SDS Vers 2.0

Layout: 12 Wells Format | 2 NTC | 2 C+

Format: 12 Wells Format

Buttons: Import RealTime PCR Data, No Data imported, No Data imported, IPC | Cy5 | 50IPC

Run No.	Ct	dRN	Out	Ct	dRN	Out	Ct	dRN	Out
Control									
SD									
Delta	7.00	0.20		7.00	0.20		4.00	0.33	
CutOff									

Comment:

NTC: Ct neg Inh pos

ExCtrl: Ct neg Inh pos

C+: Ct neg Inh pos

Threshold: [ ] [ ] [ ]

Nr	Well	Sample	Dil.	Task	Out	35S			NOS			IPC		
						Ct	dRN	Res	Ct	dRN	Res	Ct	dRN	Total
1														
2														
3														
4														
5														
6														
7														
8														
9														
10														
11														
12														
13														

FILE HOME INSERT PAGE LAYOUT FORMULAS DATA REVIEW VIEW

Paste Clipboard

Calibri 9 A<sup>↑</sup> A<sup>↓</sup> B I U Font

Alignment Wrap Text Merge & Center

Number \$ % .00 →0

Styles Conditional Formatting Format as Table Cell Styles

D6 : [X] [✓] [fx] 12 Wells Format

	A	B	D	E	F	G	H	I	L	M	N	O	P	Q	R	U	V	W	X	Y	Z	AA	AS	AT	AU
1																									
2	Module		GS-P-09.050																						
3	System		35S/NOS V2																						
4	Instrument		ABI 7500 SDS Vers 2.0																						
5	Layout		12 Wells Format   2 NTC   2 €																						
6	Format		12 Wells Format																						
7																									
8																									
9																									

NO RealTime Data imported.

Import RealTime PCR Data

No Data imported

No Data imported

IPC | Cy5 | 50IPC

FILE HOME INSERT PAGE LAYOUT FORMULAS DATA REVIEW VIEW

Paste Clipboard

Calibri 9 A<sup>↑</sup> A<sup>↓</sup>

B I U

Font

Alignment

Wrap Text Merge & Center

Number

Styles

D6

12 Wells Format

	A	B	D	E	F	G	H	I	L	M	N	O	P	Q	R	U	V	W	X	Y	Z	AA	AS	AT	AU
1			NO RealTime Data imported.																						
2	Module		GS-P-09.050																						
3	System		35S/NOS V2																						
4	Instrument		ABI 7500 SDS Vers 2.0						No Data imported					No Data imported										IPC   Cy5   50IPC	
5	Layout		12 Wells Format   2 NTC   2 €																						
6	Format		12 Wells Format																						
7																									
8																									
9																									

Import RealTime PCR Data





PCR system:

Module

Instrument

NTC / C+

Plateformat

Used Lot#:

PCR run info

Run ID

Export file:

GS-P-09.036	NOS Terminator	
GS-P-09.037	35S Promotor	
GS-P-09.038	event MON89034	
GS-P-09.039	bar-gene	
GS-P-09.040	event MON88017	
GS-P-09.041	event 3272	
GS-P-09.043	eventH7-1	
GS-P-09.044	event305423	
GS-P-09.045	nos/DHFR1-T/G	
GS-P-09.046	Tobacco NIR2	
GS-P-09.047	Cotton SAH7	
GS-P-09.048	event T45	
GS-P-09.050	35S/NOS V2	
GS-P-09.052	vip3Aa gene	
GS-P-09.053	eventDP098140-6	
GS-P-09.055	event KMD1	

Import not possible

Cancel

45

pick file ...

Please fill all red labelled Fields!



**PCR system:**

Module

**Instrument**

- NTC / C+**
- ABI 7500 SDS Vers 1.4
  - ABI 7500 SDS Vers 2.0** ←
- Plateformat**
- Agilent AriaMX not Validated for this Intrument
  - Agilent MX3005
  - Biorad CFX96 not Validated for this Intrument
  - Roche LightCycler480 I
  - Roche LightCycler480 II

**Used Lot#:**

Lot starts with

**Import not possible**

Cancel

**PCR run information:**

**Run ID**

**Cycles**

**Export file:**

Please fill all red labelled Fields!

**PCR system:**

Module

GS-P-09.050

35S/NOS V2

Instrument

ABI 7500 SDS Vers 2.0

NTC / C+

4 NTC | 4 C+

**Plateformat**

6 NTC | 4 C+

4 NTC | 4 C+

**2 NTC | 2 C+**

**Oligo- / MasterMix**

**BasicMix**

**C+**

**Used Lot#:**

Lot starts with

1121

0826

1122

**Import  
not possible**

Cancel

**PCR run information:**

**Run ID**

**Cycles**

45

**SDS V2.0 Results export**

pick file ...

**SDS V2.0 Amplification**

pick file ...

**Please fill all red labelled Fields!**



**PCR system:**

Module

Instrument

NTC / C+

Plateformat

Oligo- / MasterMix  BasicMix  C+

Lot starts with

**Used Lot#:**

**PCR run information:**

Run ID

Cycles

SDS V2.0 Results export

SDS V2.0 Amplification

Please fill all red labelled Fields!



**PCR system:**

Module

Instrument

NTC / C+

Plateformat

	Oligo- / MasterMix	BasicMix	C+
<b>Used Lot#:</b>	<input type="text" value="1121"/>	<input type="text" value="0826"/>	<input type="text" value="1122"/>
Lot starts with	1121	0826	1122

**Import not possible**

Cancel

**PCR run information:**

**Run ID**

**Cycles**

**SDS V2.0 Results export**

**SDS V2.0 Amplification**

Please fill all red labelled Fields!



**PCR system:**

Module

Instrument

NTC / C+

Plateformat

	Oligo- / MasterMix	BasicMix	C+
Used Lot#:	<input type="text" value="1121"/>	<input type="text" value="0826"/>	<input type="text" value="1122"/>
Lot starts with	1121	0826	1122

**Import not possible**

Cancel

**PCR run information:**

Run ID	Cycles
<input type="text" value="Double Screen Run 1"/>	<input type="text" value="45"/>

SDS V2.0 Results export

SDS V2.0 Amplification

Please fill all red labelled Fields!



**PCR system:**

Module	GS-P-09.050	35S/NOS V2
Instrument	ABI 7500 SDS Vers 2.0	
NTC / C+	2 NTC   2 C+	
Plateformat	12 Wells Format	

	Oligo- / MasterMix	BasicMix	C+
<b>Used Lot#:</b>	1121	0826	1122
Lot starts with	1121	0826	1122

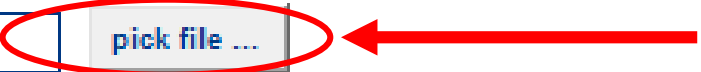
Import not possible

Cancel

**PCR run information:**

Run ID	Cycles
Double Screen Run 1	45

SDS V2.0 Results export	<input type="text"/>	<input type="button" value="pick file ..."/>
SDS V2.0 Amplification	<input type="text"/>	<input type="button" value="pick file ..."/>



Please fill all red labelled Fields!

Select ABI7500 Results export file for Run: Double Screen Run 1 (SDS V2.0)

Documents > GMO Testing Facility

Documents library  
GMO Testing Facility

Arrange by: Folder

- ABI plate setup
- Animal species templates
- BJ Koortzen
- GMO Qualitative screen results
- GMO Qualitative Screen runs
- GMO Quantification results
- GMO Quantification runs
- GMO templates
- GS Evaluation sheets
- Double Screen Run 1 amplification
- Double Screen Run 1 results**

File name: Double Screen Run 1 results

Type: Microsoft Excel 97-2003 Worksheet  
Size: 53.5 KB  
Date modified: 1/11/2019 3:04 PM

Open Cancel

the PCR With IPC NEW 2017 - Excel

Conditional Formatting | Format as Table | Cell Styles | Insert | Delete | Format

X	Y	Z	AA	AB	AC	AD	AE	AF	AG	AH	AI	AJ	AK	AL	AM	AN	AO	AP	AQ	AR	AS	AT	AU	AV	AW

1122

1122

Import not possible

Cancel

21

22

23 Comment:

24

25

26

27

28

Nr	Well	Sample	Dil.	Task
30	1			
31	2			

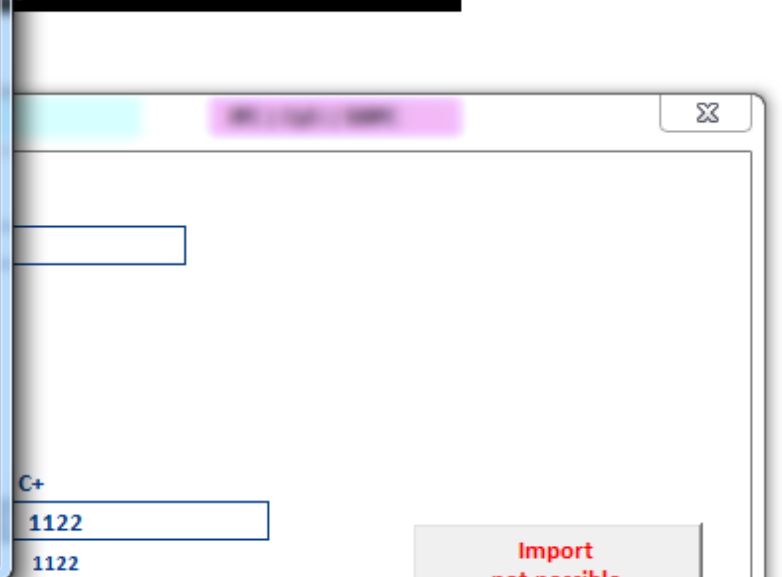
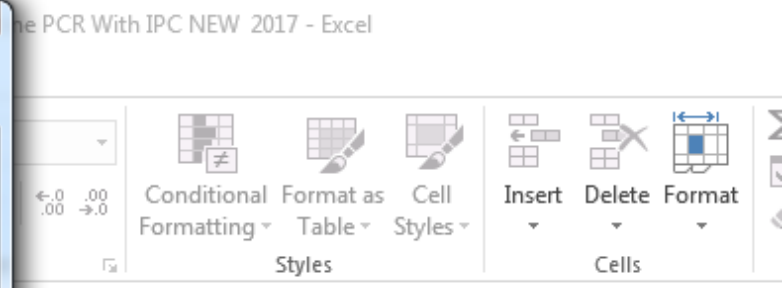
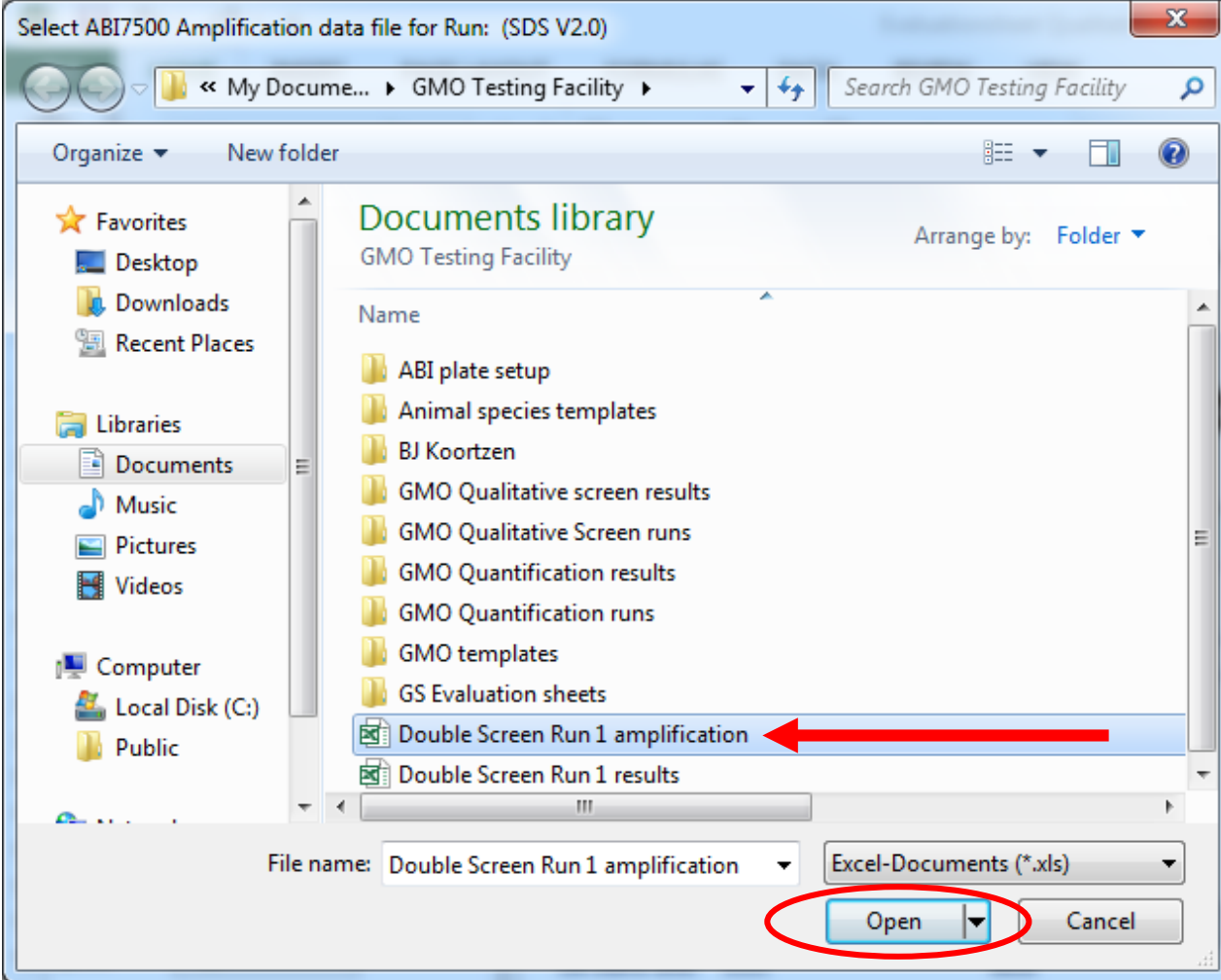
**PCR run information:**

Run ID	Cycles
Double Screen Run 1	45

SDS V2.0 Results export

SDS V2.0 Amplification

Please fill all red labelled Fields!



21					
22					
23	Comment:				
24					
25					
26					
27					
28					
29	Nr	Well	Sample	Dil.	Task
30	1				
31	2				
32	3				

**PCR run information:**

Run ID	Cycles
Double Screen Run 1	45

SDS V2.0 Results export

SDS V2.0 Amplification

Please fill all red labelled Fields!



**PCR system:**

Module

Instrument

NTC / C+

Plateformat


	Oligo- / MasterMix	BasicMix	C+
<b>Used Lot#:</b>	<input type="text" value="1121"/>	<input type="text" value="0826"/>	<input type="text" value="1122"/>
Lot starts with	1121	0826	1122

**PCR run information:**

Run ID	Cycles
<input type="text" value="Double Screen Run 1"/>	<input type="text" value="45"/>

SDS V2.0 Results export

SDS V2.0 Amplification





**PCR system:**

Module

Instrument

NTC / C+

Plateformat

Oligo- / Mast

Used Lot#:

Lot starts with 1121

**PCR run information:**

Run ID

SDS V2.0 Results export

SDS V2.0 Amplification

Import ABI 7500 SDS Software Version 2.0

Ct- und dR/dRN-Values imported sucessfully.

-----

Please assign 'Tasks' to Wells if necessary  
(see Section Evaluation (Instructions)).



**PCR system:**

Module

Instrument

NTC / C+

Plateformat

Oligo- / Mast

**Used Lot#:**

Lot starts with

**PCR run information:**

Run ID

SDS V2.0 Results export

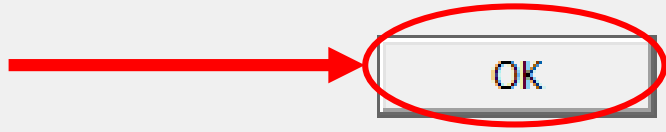
SDS V2.0 Amplification

Import ABI 7500 SDS Software Version 2.0

Ct- und dR/dRN-Values imported sucessfully.

-----

Please assign 'Tasks' to Wells if necessary  
(see Section Evaluation (Instructions)).



1  
2 **Module** GS-P-09.050  
3 **System** 35S/NOS V2  
4 **Instrument** ABI 7500 SDS Vers 2.0  
5 **Layout** 2 NTC | 2 C+  
6 **Format** 12 Wells Format

Import RealTime PCR Data

35S | FAM | 5035S

NOS | JOE | 50NOS

IPC | Cy5 | 50IPC

11 **Run No.** 07-11-2019  
12 **Lot MM/OM** 1121  
13 **Lot BM** 0286  
14 **Lot C+** 1122  
15 **Cycles** 45

	Ct	dRN	Out
Control	30.23	3.70	0
SD	0.11	0.01	
Delta	7.00	0.20	
CutOff	37.23	0.74	

	Ct	dRN	Out
Control	32.95	2.01	0
SD	0.39	0	
Delta	7.00	0.20	
CutOff	39.95	0.40	

	Ct	dRN	Out
Control	33.04	1.19	0
SD	0.10	0	
Delta	4.00	0.33	
CutOff	37.04	0.39	

23 **Comment:**

	Ct	neg	Inh	pos
NTC	45.00	2	0	0
ExCtrl	N/A	N/A	N/A	N/A

	Ct	neg	Inh	pos
NTC	45.00	2	0	0
ExCtrl	N/A	N/A	N/A	N/A

C+	33.02
ExCtrl	N/A

Threshold 0.10

0.10

0.10

Nr	Well	Sample	Dil.	Task
30	1	A1	NC	N
31	2	A2	NC	N
32	3	A3	PC	P
33	4	A4	PC	P
34	5	A5	AS	U
35	6	A6	AS	U

Out	35S
	Ct dRN Res
	45.00 -0.02 -
	45.00 -0.02 -
	30.34 3.68 +
	30.11 3.71 +
	25.65 4.07 +
	25.90 4.12 +

NOS
Ct dRN Res
45.00 0.00 -
45.00 0.00 -
32.56 1.99 +
33.34 2.03 +
28.04 2.12 +
28.35 2.13 +

IPC
Ct dRN Total
32.95 1.21 valid
33.14 1.18 valid
32.80 1.14 valid
33.24 1.16 valid
34.24 1.09 valid
34.91 1.03 valid

	35S					Out	35S		
	Nr	Well	Sample	Dil.	Task		Ct	dRN	Res
28									
29									
30	1	A1	NC		N		45.00	-0.02	-
31	2	A2	NC		N		45.00	-0.02	-
32	3	A3	PC		P		30.34	3.68	+
33	4	A4	PC		P		30.11	3.71	+
34	5	A5	AS		U		25.65	4.07	+
35	6	A6	AS		U		25.90	4.12	+
36	7	A7	MTL		U		25.99	4.03	+
37	8	A8	MTL		U		26.12	4.02	+
38	9	A9	TM		U		26.28	3.98	+
39	10	A10	TM		U		26.36	4.03	+
40	11	A11	DC		U		26.73	3.99	+
41	12	A12	DC		U		26.53	3.95	+
42	13	B1	TMG		U		26.15	3.97	+
43	14	B2	TMG		U		25.82	4.08	+
44	15	B3	BBL		U		31.45	3.53	+
45	16	B4	BBL		U		30.87	3.67	+
46	17	B5	AC		U		32.19	3.61	+
47	18	B6	AC		U		32.48	3.48	+

NOS		
Ct	dRN	Res
45.00	0.00	-
45.00	0.00	-
32.56	1.99	+
33.34	2.03	+
28.04	2.12	+
28.35	2.13	+
28.61	2.10	+
28.60	2.08	+
28.72	2.07	+
28.73	2.10	+
29.22	2.04	+
28.95	2.02	+
28.54	2.03	+
28.20	2.10	+
32.91	2.01	+
32.58	1.99	+
34.55	1.94	+
34.17	1.96	+

IPC		
Ct	dRN	Total
32.95	1.21	valid
33.14	1.18	valid
32.80	1.14	valid
33.24	1.16	valid
34.24	1.09	valid
34.91	1.03	valid
34.67	1.06	valid
34.44	1.08	valid
34.41	1.10	valid
34.50	1.07	valid
35.07	1.05	valid
36.59	0.88	valid
33.70	1.13	valid
34.24	1.03	valid
33.14	1.18	valid
33.26	1.22	valid
33.01	1.18	valid
33.12	1.20	valid

28						Out	35S			NOS			IPC		
29	Nr	Well	Sample	Dil.	Task		Ct	dRN	Res	Ct	dRN	Res	Ct	dRN	Total
46	17	B5	AC		U		32.19	3.61	+	34.55	1.94	+	33.01	1.18	valid
47	18	B6	AC		U		32.48	3.48	+	34.17	1.96	+	33.12	1.20	valid
48	19	B7	AM		U		33.84	3.28	+	36.29	1.78	+	33.44	1.21	valid
49	20	B8	AM		U		37.56	1.95	+	36.02	1.82	+	33.25	1.18	valid
50	21	B9	TJ		U		45.00	-0.18	-	37.30	1.67	+	33.26	1.18	valid
51	22	B10	TJ		U		34.67	2.85	+	35.79	1.82	+	33.44	1.21	valid
52	23	B11	MP		U		26.59	3.96	+	28.69	2.05	+	34.21	1.09	valid
53	24	B12	MP		U		26.50	3.98	+	28.60	2.04	+	34.38	1.10	valid
54	25	C1	KHM		U		25.66	3.92	+	27.90	2.04	+	34.34	0.85	valid
55	26	C2	KHM		U		25.50	4.03	+	27.68	2.04	+	34.91	0.79	valid
56	27	C3	MS		U		25.60	4.24	+	27.70	2.14	+	34.37	1.05	valid
57	28	C4	MS		U		25.58	4.32	+	27.84	2.14	+	34.22	1.08	valid
58	29	C5	AA		U		25.58	4.32	+	27.83	2.17	+	34.72	1.02	valid
59	30	C6	AA		U		25.66	4.41	+	27.89	2.14	+	34.45	1.04	valid
60	31	C7	AHM		U		26.02	4.38	+	28.29	2.23	+	34.41	1.08	valid
61	32	C8	AHM		U		25.87	4.34	+	28.14	2.20	+	34.29	1.11	valid
62	33	C9	RMG		U		29.00	3.76	+	30.59	2.02	+	33.72	0.90	valid
63	34	C10	RMG		U		29.05	3.75	+	30.61	2.03	+	33.47	0.98	valid
64	35	C11	NP		U		31.13	3.70	+	32.99	2.01	+	33.19	1.15	valid
65	36	C12	NP		U		31.22	3.61	+	33.37	1.91	+	33.40	1.18	valid
66	37	D1	A1		U		29.22	3.58	+	30.86	1.99	+	33.44	0.93	valid
67	38	D2	A1		U		29.14	3.66	+	30.84	2.03	+	33.24	1.05	valid
68	39	D3	OP		U		32.59	3.56	+	35.49	1.89	+	34.19	1.13	valid
69	40	D4	OP		U		32.67	3.17	+	35.10	2.18	+	34.44	1.25	valid

- 1. To Edit or make any changes to the sheet,**
- 2. Unprotect the sheet (under Review Tab)**
- 3. Change the “task” if required to “EC” for extraction control**
- 4. The dilutions used can be entered under the “Dil” column as required**
- 5. Ensure NC and PC have passed**
- 6. Check for inconsistent results between duplicates**
- 7. Check for inconsistent Ct values between duplicates**
- 8. Check for inhibition (IPC invalid)**
- 9. Any unusual results must be analyzed simultaneously using the amplification plot**
- 10. For samples indicating inhibition, a further dilution can be made and subjected to re-PCR**