Confirmation of Salmonella isolates by Real-time Polymerase Chain Reaction (PCR)

This method is applicable to the confirmation of pure cultural isolates as *Salmonella* spp. Real-time PCR assembly and data analysis protocols are described below for two instrument platforms: SmartCycler II and Applied Biosystems 7500 Fast Real-time PCR thermocycler. Use of other platforms and protocols must first be validated per FDA microbiological methods validation guidelines

(https://www.fda.gov/downloads/ScienceResearch/FieldScience/UCM298730.pdf) or other internationally recognized validation guidelines such as AOAC International's Appendix J

(http://www.eoma.aoac.org/app_j.pdf) or the International Organization for Standardization's 16140 (www.iso.org).

A. Equipment and Materials

- 1. SmartCycler II PCR thermolcycler (Cepheid, Sunnyvale, CA) capable of performing cycling parameters described below and simultaneous real-time sequence detection for FAM, Texas Red, and Cy5 dyes.
- 2. ABI 7500 Fast Real-time PCR thermocycler (SDS version 1.4) (ThermoFisher Scientific, Waltham, MA), capable of performing cycling parameters described below and simultaneous real-time sequence detection for FAM, Texas Red, and Cy5 dyes.
- 3. SmartCycler PCR reaction tubes (minimum reaction volume of 25 µl; Cepheid # 900-0003) and racks (Cepheid # 900-0087) compatible with PCR thermocycler
- 4. MicroAmpTM Fast Optical 96-Well Reaction Plates (ThermoFisher Scientific # 4346906)
- 5. MicroAmpTM Optical Adhesive Film for 7500 Fast Plates (ThermoFisher Scientific # 4311971)
- 6. MicroAmpTM Fast Reaction 8-Tube Strips (ThermoFisher Scientific # 4358293)
- 7. MicroAmpTM Optical 8-Cap Strips (ThermoFisher Scientific # 4323032)
- 8. Appropriate ABI 7500 Fast Plate Holder (specific for 96-well tray or 8-strip well tubes)
- 9. Ice bucket and ice
- 10. Microcentrifuge tubes (0.5 to 2.0 mL)
- 11. Pipets (1-1000 µL volume)
- 12. Pippete tips (0.2 to 1000 µL volume) (aerosol resistant tips)
- 13. QIAprep Spin Miniprep Kit (QIAgen # 27104)
- 14. Qubit dsDNA BR Assay Kit (ThermoFisher Scientific # 32850 or 32853)
- 15. Sterile gloves
- 16. Vortex Mixer
- 17. Water bath or heat block capable of maintaining 100°C

B. Media and Reagents

- 1. Sterile Molecular Grade water
- 2. illustraTM PuReTaqTM Ready-To-GoTM PCR beads (GE Healthcare # 27955801)
- 3. Applied BiosystemsTM TaqMan Fast Advance Master Mix (ThermoFisher Scientific # 4444556)
- 4. *Salmonella* primers and probes listed in Table 1 are specific to real-time PCR platforms being used.
 - a. Primers 10 μ M working solution of each primer listed in Table 1. Stock (1000 μ M) and working solutions can be prepared from commercially synthesized primers with basic desalt purification

(Biosearch Technologies or equivalent) by rehydrating with sterile molecular biology grade water to appropriate concentrations. Store at -20° C to -70° C non-frost-free freezer.

- b. Probes 10 μ M working solution of each probe listed in Table 1. Dual hybridization probes should be purchased as Dual HPLC-purified and labeled as indicated in Table 1. Stock (100 μ M) and working solutions can be prepared from commercially synthesized probes (Biosearch Technologies or equivalent) by rehydrating with sterile molecular biology grade water to appropriate concentrations. Working solutions should be aliquoted in small amounts and stored frozen (-20 to -70°C) and away from light until use to avoid fluorophore degradation.
- c. Exogenous Internal Amplification Control (IAC)

IAC is a synthetic 100 bp sequence: 5'-AGTTGCAGTGTAACCGTCATGTACCAGTAATCTGCGTCGCACGTGTGCACCTAGTCTA ATCACTTATGACTCAGATAACTTAACAGCAGAGTCTCGTCGA.

IAC plasmid pCR2.1-InC (Plasmid #83959) is available through Addgene at https://www.addgene.org/83959. Host strain containing pCR2.1-InC plasmid can be cultured in LB (Lysogeny broth) or BHI (Brain Heart Infusion) broth containing either kanamycin (50µg/mL) or ampicillin (100µg/mL) at 37°C for 16 hours.

The pCR2.1-InC plasmid can be extracted from overnight culture by using QIAprep Spin Miniprep Kit. The plasmid concentration can be quantitated with Qubit dsDNA BR Assay Kit.

Primers ¹	Gen Bank #	Bases	5' → 3' Sequence
<i>Sal</i> 1598 F	U43273	20	AACGTGTTTCCGTGCGTAAT
Sal1859 R	U43273	20	TCCATCAAATTAGCGGAGGC
IAC F		22	AGTTGCAGTGTAACCGTCATGT
IAC R		22	TCGACGAGACTCTGCTGTTAAG
Probes ¹			
Sal1631PFAM		20	FAM-TGGAAGCGCTCGCATTGTGG-BHQ
IAC30PCy5		20	Cy5-ATCTGCGTCGCACGTGTGCA-BHQ

Table 1. Primer/probe sequences for use on SmartCycler II and ABI 7500 Fast platforms

¹Primer/Probe name composed of target (*Sal* = *Salmonella* species targeting *invA* gene, IAC = Internal Amplification Control), 5' base position of oligonucleotide in the respective gene sequence specified in column 3 and forward primer (F), reverse primer (R) or probe (P).

C. PCR Controls

- 1. For a positive PCR control, include a template prepared from *Salmonella enterica*, such as ATCC 8324, *Salmonella* Gaminara.
- 2. Always include a no template (water) negative control in every run.

D. DNA Template Preparation

- Transfer 250 μL of overnight culture in BHI broth at 35±2°C to a 1.5 mL microcentrifuge tube; or suspend a colony from a 24 h fresh prepared BHI plate in 250 μL sterile water in a 1.5 mL microcentrifuge tube.
- 2. Centrifuge $10,000 \times g$ for 5 min.
- 3. Remove the supernatant and completely resuspend pellet in $250 \,\mu$ L sterile water.
- 4. Boil templates for 20 min.
- 5. Immediately cool down on ice for 5 min.
- 6. Centrifuge at $10,000 \times g$ for 5 min. Remove and save the supernatant as DNA template (This may be frozen, minimum -20°C, for future PCR tests).

E. Preparation of qPCR Salmonella Master Mix

1. Preparation of dehydrated qPCR Salmonella Master Mix

- **a.** Mix all components in Table 2 in 1.5 mL microcentrifuge tube by vortex at top speed and centrifuge briefly.
- b. Aliquot 5 μL Master Mix solution for an evaluation run with *Salmonella* and negative controls on SmartCycler II, and a second 5 μL aliquot for evaluation run on ABI 7500 Fast (see F for Reaction assembly).
- **c.** When both evaluation runs were satisfactory, dispense 10 μL Master Mix solution per tube to sterile 1.5 mL black microcentrifuge tubes.
- d. Dry the qPCR *Salmonella* Master Mix solution with tube lids open in a vacuum chamber for 2 to 3 days and shield the vacuum chamber completely from light.
- e. The vacuum dried qPCR *Salmonella* Master Mix can be stored in aluminum porch with silica gel at ambient for 2 years.
- f. Each tube of qPCR *Salmonella* Master Mix can carry out forty 25 μL reactions on SmartCycler II or fifty 20 μL reactions on ABI 7500 Fast systems.

Component	Volume (µL)
Primer Sal1598 F (1000 µM Solution)	16.0
Primer Sal1859 R (1000 µM Solution)	16.0
Primer IAC F (1000 µM Solution)	6.7
Primer IAC R (1000 µM Solution)	6.7
Probe Sal1631PFAM (100 µM Solution)	200.0
Probe IAC30PCy5 (100 µM Solution)	250.0
IAC DNA template (0.75 pg/µL)	25.0
20% sucrose (0.22 µM filter-sterilized)	130.0

Table 2. Recipe for 100 tubes of qPCR Salmonella Master Mix

PCR grade water	349.6
Total	1000.0

2. Preparation of wet qPCR Salmonella Master Mix

- a. Mix all components in Table 3 in 1.5 mL black microcentrifuge tube by vortex at top speed and centrifuge briefly.
- b. Dilute to 1X or 2.5X for reaction set up accordingly (see F for Reaction assembly).
- c. Perform a QA run for every batch of wet master mix on SmartCycler II and ABI 7500 Fast.
- d. Shield the wet qPCR Salmonella master mix from light and stored at 4°C up to two months.

Table 3. Recipe for 1 tube of qPCR Salmonella Master Mix (10X)

Component	Volume (µL)
Primer Sal1598 F (10 µM Solution)*	16.0
Primer Sal1859 R (10 µM Solution)*	16.0
Primer IAC F (10 µM Solution)*	6.7
Primer IAC R (10 µM Solution)*	6.7
Probe Sal1631PFAM (10 µM Solution)**	20.0
Probe IAC30PCy5 (10 µM Solution)**	25.0
IAC plasmid DNA template (0.075 pg/µl)***	2.5
20% sucrose (0.22 µM filter-sterilized)	1.3
PCR grade water	5.8
Total	100.0

* 10 µM primer working solutions were used.

** $10 \,\mu$ M probe working solutions were used.

*** 10-fold diluted IAC template was used.

F. Real-Time PCR assembly

1. Reaction assembly for SmartCycler II (Table 4)

Table 4. SmartCycler II Amplification Reaction Components

Volume (µL) /rxn	Component
22.5	1X working solution qPCR Salmonella Master Mix (1X MM) ¹
1.5 beads	illustra PuReTaq Ready-To-Go PCR beads
2.5	Template (Sample or control)

¹Resuspend one tube of dehydrated *Salmonella* Master Mix in 900 μ L of PCR grade water to make 1X MM. ¹Add 800 μ L of PCR grade water to 1 tube of wet qPCR *Salmonella* master mix to make 1X MM. **Note:** Each 900 μ L of 1X qPCR *Salmonella* master mix tube will need 60 PCR beads for total 40 reactions.

2. Reaction Assembly for ABI 7500 Fast (Table 5)

Volume (µL) /rxn	Component
8.0	2.5X working solution of qPCR <i>Salmonella</i> Master Mix $(2.5X \text{ MM})^2$
10.0	TaqMan Fast Advance Master Mix
2.0	Template (Sample or control)

Table 5. ABI 7500 Fast Amplification Reaction Components

²Resuspend one tube of dehydrated Salmonella Master Mix in 400 µl of PCR grade water to make 2.5X MM.

²Add 300 µl of PCR grade water to 1 tube of wet qPCR *Salmonella* master mix (10X) to make 2.5X MM.

Note: Each 400 μl of 2.5X qPCR *Salmonella* master mix tube will need 500 μl TaqMan Fast Advance Master Mix for total 50 reactions.

G. Real-Time PCR run

1. Run on SmartCycler II

a. Create run on SmartCycler II.

Give each run a unique run name, select Dye set FCTC25, select 2-step PCR protocol as described below and assign appropriate sites on SmartCycler block.

Initial Activation	Each of 50 Cycles			
95°C, 180 sec	95°C, 15 sec	60°C, 30 sec		
Optics Off	Optics Off	Optics On		

b. Qualitative analysis setting

On SmartCycler II Instrument, set the following analysis settings for FAM, and Cy5 channels. Update analysis settings if they are changed before recording results.

Usage	Assay
Background subtraction	ON
Background Min. Cycle	5
Background Max. Cycle	40
Curve Analysis	Primary
Threshold Setting	Manual
Manual Threshold Fluorescence Units	20.0
Auto Min Cycle	5

Auto Max Cycle	10
Valid Min Cycle	3
Valid Max. Cycle	60
Boxcar Avg. Cycles	0

- c. Report of Result
 - i. Primary fluorescence curves that cross the threshold will be recorded as "POS".
 - ii. The cycle the sample crossed the threshold will be displayed in the Results Table view.
 - iii. Results can also be viewed graphically.
 - iv. A report can be generated or a screen capture of the results table view can be used to record data (see Figure 1 for an example).
- d. Interpretation of Results

If *inv*A gene target in the FAM and IAC in the Cy5 channels are POS, the isolate is a *Salmonella* spp. If only the Cy5 channel is POS, the isolate is a *non-Salmonella* spp.

2. Run Real-Time PCR on ABI 7500 Fast

- a. Turn on the computer and ABI 7500 Fast.
- b. Launch ABI 7500 Fast System Software v1.4.2.
- c. Create run by using New Document Wizard.
- d. Highlight all wells and select "Detector Manager" under "Tool".
- e. Shift select Salmonella (FAM) and IAC (Cy5) and click "Add to Plate Document.
- f. Under "Well Inspector" in "View", mark both "Salmonella" and "IAC", and select "ROX" for Passive Reference.
- g. Under "Instrument", select 2-step PCR protocol as described below.

Activation of UDG Initial Activation		Each of 50 Cycles		
50°C, 2 min	95°C, 5 min	95°C, 3 sec	60°C, 30 sec	

- h. Save the newly created file as template for *Salmonella* in **.sdt** format. The *Salmonella*_template.**sdt** file can be used as template for future .sds run file with all preset parameters.
- i. Save the file again in .sds format as a run file with a different name such as "Salmonella_Test.sds".
- j. Assign appropriate sites with sample names on corresponding wells and save the file.
- k. Load the samples and start the run under "Instrument". If the "Start" button under the "Instrument" is not highlighted, close and re-open the "*Salmonella*_Test.sds" file to initialize ABI 7500 Fast System. Start the run by clicking the "Start" button.
- 1. After the reaction completed successfully, the results can be analyzed, viewed and reported (see Figure 2 for an example).

Analysis setting:

- i. Manual Ct
- ii. Threshold: 0.05 (w/ROX)

iii. Manual Baseline

Start (cycle): 3 End (cycle): 15

m. Interpretation of Results

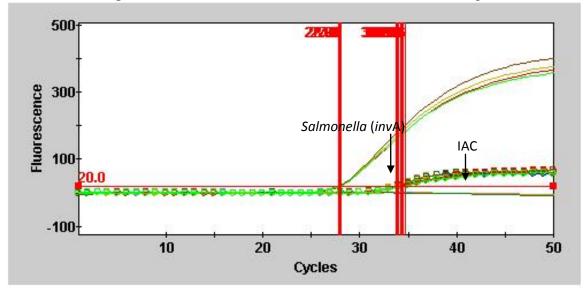
If both *Salmonella* target (invA gene) in the FAM channel and InC in the Cy5 channels are positive with Ct values, the isolate is a *Salmonella* spp. If only the Cy5 (InC) channel is positive with a Ct value and undetected (undetermined) in FAM channel, the isolate is a non-*Salmonella* spp. For each sample, check if it has a normal amplification plot as shown in Figure 2.

Figure 1. Example of results output from Smart Cycler II

lesults Table	Site ID	Protocol	Sample ID	Sample Type	Notes	Status	FAM Std/Res	FAM Ct	Cy5 Std/Res	Cy5 Ct
nalysis Settings	A1	Salmonella	Neg-1	UNKN		OK	NEG	0.00	POS	33.85
Protocols	A2	Salmonella	Neg-2	UNKN		OK	NEG	0.00	POS	33.94
JTM of Melt	A3	Salmonella	Neg-3	UNKN		OK	NEG	0.00	POS	33.73
emperature	A4	Salmonella	Neg-4	UNKN		OK	NEG	0.00	POS	33.90
ntercalate	A5	Salmonella	SLM 8324-1	UNKN		OK	POS	27.88	POS	34.35
felt	A6	Salmonella	SLM 8324-2	UNKN		OK	POS	27.80	POS	34.44
AM	A7	Salmonella	SLM 8324-3	UNKN		OK	POS	28.07	POS	34.26
>y3	A8	Salmonella	SLM 8324-4	UNKN		OK	POS	27.95	POS	34.66
exas Red										
>y5										
Standard										
AM-Cv5										

1A. Results table view for Salmonella enterica, ATCC 8324 and negative control.

1B. Results: Graphical view for Salmonella enterica ATCC 8324 and negative control.



Site	I eq	hren
ono	1008	orner

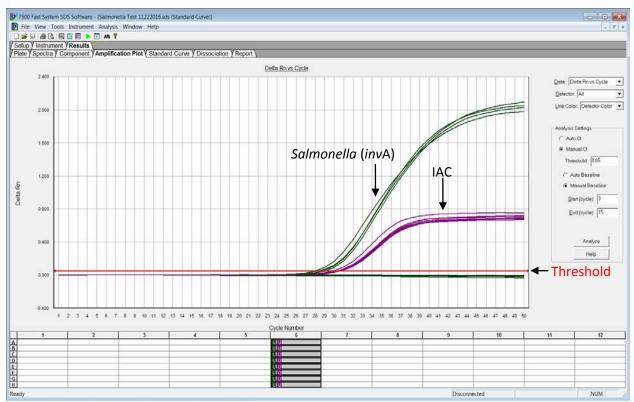
Site ID	Sample ID	FAM Ct	Cy5 Ct	Protocol
A1 📃	Neg-1	0.00	33.85	Sahmonella
A2 🔳	Neg-2	0.00	33.94	Sahmonella
A3 📕	Neg-3	0.00	33.73	Sahmonella
A4 🔳	Neg-4	0.00	33.90	Sahmonella
AS 📕	SLM 8324-1	27.88	34.35	Salmonella
A6 📕	SLM 8324-2	27.80	34.44	Sahmonella
A7 📕	SLM 8324-3	28.07	34.26	Sahmonella
A8 📕	SLM 8324-4	27.95	34.66	Sahnonella

Data Type	Line Type
Primary Curve	

Channel	Symbol
FAM	None
Суб	000

Figure 2. Example of results output from ABI 7500 Fast

2A. Amplification Plot Salmonella (FAM) and IAC (Cy5)



Amplification Plot

Data: Delta Rn vs. Cycle

Detector: All,

Line Color: "Well color" or "Detector Color"

Analysis Settings

Manual Ct

Threshold: 0.05 (w/ROX)

Manual Baseline

Start (cycle): 3

End (cycle): 15

2B. Plate

		0S Software - [Salmone Instrument Analysis		(Standard Curve)]								
Setup / Instrument / Results Plate (Spectra / Component / Amplification Plot / Standard Curve / Dissociation / Report												
	1	2	3	4	5	6	1	8	9	10	11	12
						Neg-1 Undet 29.78						
						Neg-2 Undet J 30.46						
0						Neg-3 Undet. 1 30.36						
						Neg-4 1 Undet. 1 30.36						
						SLM #8324-1 28.66 30.64						
						SLM #8324-2 29.04 30.63						
						SLM #8324-3 28 33 30 44	-					
						SLM #8324-4 17 28.01 17 30.25						
ady			l						Disconne	and		NUM

3C. Report

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strument YResul	ts		and works to									
				StdDev Ct	Quantity	Mean Qty	StdDev Qty	Filtered	Tm	User Defined #1	User Defined #2	User Defined #
											-	
Neg-3												
Neg-4		Unknown										
Neg-4		Unknown										
SLM #8324-1	Salmonella	Unknown										
	InC	Unknown										
SLM #8324-2	Salmonella	Unknown	29.0366									
SLM #8324-2	InC	Unknown	30.629									
SLM #8324-3	Salmonella	Unknown	28.3283								-	
SLM #8324-3	InC	Unknown	30.435									
SLM #8324-4	Salmonella	Unknown	28.0126									
SLM #8324-4	InC	Unknown	30.2465									
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	Sample Name Sample Name Neg-1 Neg-2 Neg-3 Neg-3 SLM #8324-1 SLM #8324-1 SLM #8324-2 SLM #8324-2 SLM #8324-2 SLM #8324-3 SLM #8324-3 SLM #8324-3 SLM #8324-3 SLM #8324-3	Ormponent Y Amplificati Sample Name Detector Sample Name Detector Neg-1 InC Neg-2 InC Neg-3 InC Neg-3 InC Neg-3 InC Neg-3 InC Neg-3 InC Neg-4 InC Summerita Sammerita NM 8324-1 InC SUM 8324-1 InC SUM 8324-2 InC SUM 8324-3 InC SUM 8324-3 InC SUM 8324-3 InC SUM 8324-3 InC SUM 8324-4 InC	Ormponent Y Annglification Pix / Y Standa Sample Name Detector Task Neg-1 IC Uchnown Neg-2 IC Uchnown Neg-3 IC Uchnown Neg-3 IC Uchnown Neg-3 IC Uchnown Neg-3 IC Uchnown Neg-4 IC Uninown Neg-4 IC Uninown Neg-4 IC Uninown NUM 8224-1 IC Uninown NUM 8224-2 IC Uninown NUM 8224-2 IC Uninown NUM 8224-2 IC Uninown NUM 8224-2 IC Uninown NUM 8224-3 IC Uninown NUM 8224-3 IC Uninown NUM 8224-3 IC Uninown NUM 8224-4 InC Uninown NUM 8224-4 InC Uninown NUM 8224-4 InC Uninown	Octoportent / Amplification Plot / Standard Curve / Dissoci Sample Name Detector Fask Ct Neg-1 Salmonella Unknown Udeknown Udeknown	Intervent Yanghilication Pix/ Ystandard Curve Y Dissociation Report Sample Name Detector Task Ct Statbev Ct Neg-1 IC Uchanoun Udata Versite Statbev Ct Neg-1 IC Uchanoun Udata Versite Statbev Ct Neg-2 IC Uchanoun Udata Versite Versite Versite Neg-3 IC Uchanoun Udata Versite V	Sample Rame Ortgottom Pick / Standard Curve // Dissociation /Report Sample Rame Detector Tek Ct StatDet Rame Neg-1 Salinosella Unknown Undet Outentity Neg-1 InC Unknown 10 det Neg-1 Neg-2 Salinosella Unknown 10 det Neg-1 Neg-2 Salinosella Unknown 10 det Neg-1 Neg-2 Salinosella Unknown 10 det Neg-3 Neg-3 Salinosella Unknown 10 det Neg-3 Neg-3 Salinosella Unknown 10 det Neg-4 Neg-4 Salinosella Unknown 10 det Neg-4 Neg-4 Salinosella Unknown 10 det Neg-4 NM 8324-1 Salinosella Unknown 20 3565 Salinosella Salinosella	Oragonetty Oragin Other Status Other Status Other Status Mean Cay Neg-1 Salmoetlar Unincome Longet Status Mean Cay Neg-1 Salmoetlar Unincome Longet Status Mean Cay Neg-1 Salmoetlar Unincome Longet Status Mean Cay Neg-2 Salmoetlar Unincome Longet Salmoetlar Salmoetlar Neg-2 Salmoetlar Unincome Doddt Neg-3 Salmoetlar Unincome Neg-3 Salmoetlar Unincome Doddt Neg-4 Neg-4 Neg-4 Neg-3 Salmoetlar Unincome Dodt Neg-4 Neg-4 Neg-4 Neg-3 Salmoetlar Unincome 20 5056 Salmoetlar Salmoetlar Salmoetlar Salmoetlar Salmoetlar Unincome 20 32 35 Salmoetlar Salmo	Sample Rame Detector Y Sample Rame Detector Y Sample Rame Under Version Subserversion Mean Mean	Sample Rame Detector Y Amplification Pice Y Standard Curve Y Dissociation Report Neg-1 Salmonella Udetown Udetown Udetown Neg-1 Report Salmonella Udetown Filtered Neg-1 RC Udetown Udetown Udetown Vectown Neg-1 Report Neg-1 Report Neg-1 Report Neg-1 Report Neg-2 Neg-2	Sample Rame Detector Y Standard Curvey / Dissociation / Report / Sample Rame Detector Task Ct StatDer Ch Mean Ch StatDer Ch Filtered Tm Neg-1 Salmonella Udexom 129.716.4 Image /	Sample Rame Detector Y Standard Curve Y Dissociation / Report / Sample Rame Detector Tak Ct Submer X Gaussity Mean Oxy Statuse Rame Tm User Defined P1 Neg-1 Sample Rame Udexom Udexom Udex Tm User Defined P1 Neg-1 Sc Udexom Udexom Udex Tm User Defined P1 Neg-2 Sample Rame Udexom Udexom Udexom Udexom Tm User Defined P1 Neg-2 Sample Rame Udexom Udexom Udexom Udexom Udexom Tm Neg-1 Neg-3 Sample Rame Udexom Udexom Udexom Udexom Tm Neg-1 Tm Neg-1	Sample Rane Detector Yanglification Piol Standper Max User Defined #1 User Defined #2 Neg-1 Salmonkia Udetown 29.7754

InC: internal amplification control (IAC)