

**HIDS**<sub>2016</sub>  
Human IDentification Solutions

**HIDS 2016**  
REGISTRATION  
OPEN

BARCELONA  
10-11 JUN

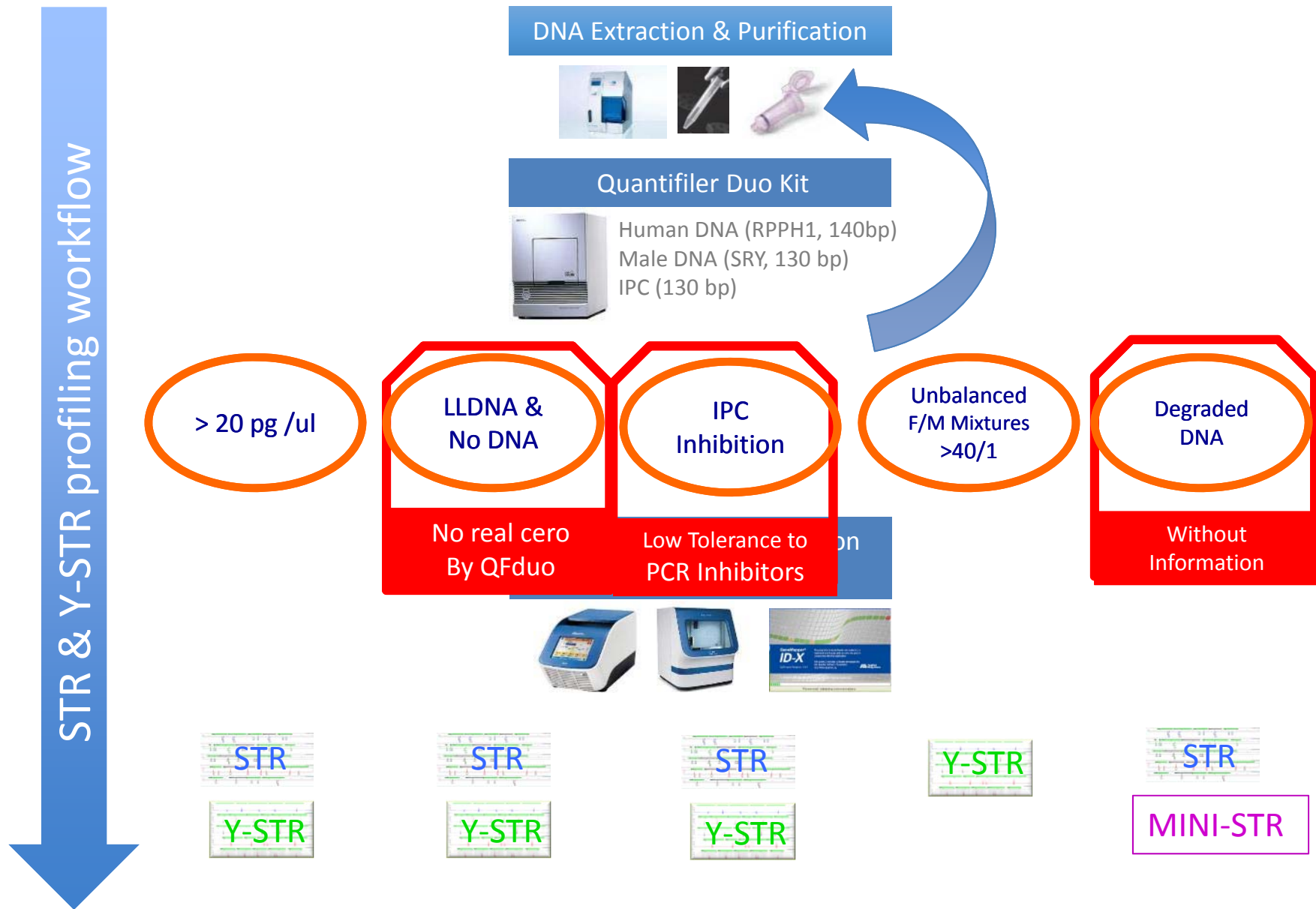
Innovations **and** perspectives

Innovations in quantitative and qualitative forensic assessment of human DNA  
Streamlining STR & Y-STR profiling workflow

*Esperanza Gonzalez, Ana Carrillo, M<sup>a</sup> Carmen Periñan, Esther Navajo, and Antonio Alonso* on behalf of the Biology Service at the National Institute of Toxicology and Forensic Sciences. Madrid Department. Spain.



# Our recent past on DNA Quantification and STR profiling workflow ...

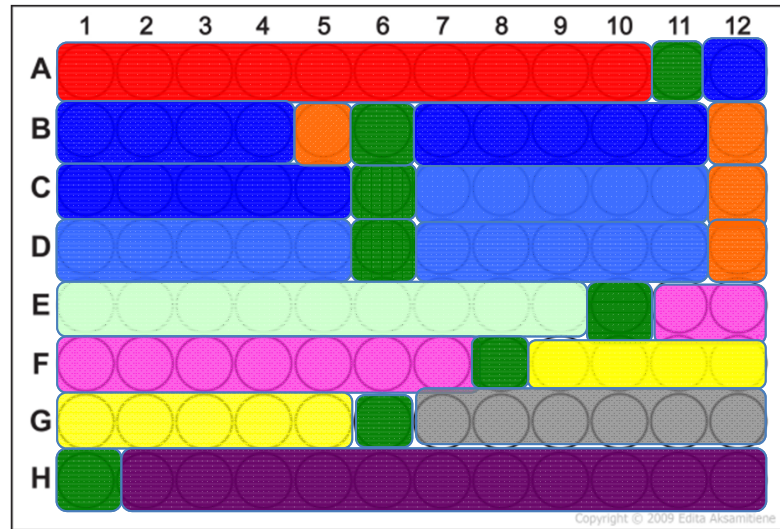


## QFTRIO Targets




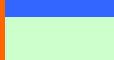
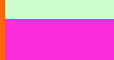





Target	Amplicon Length	Ploidy	Copy number	Dye/Quencher
Human Target, small autosomal	80 bases	Diploid	Multi-copy	VIC Dye MGB quencher
Human Target, large autosomal	214 bases	Diploid	Multi-copy	ABY Dye QSY quencher
Human Male Target	75 bases	Haploid	Multi-copy	FAM Dye MGB quencher
Internal PCR Control	130 bases	NA	Synthetic DNA	JUN Dye QSY quencher

- Human DNA
- Male DNA
- DNA Degradation
- PCR Inhibitors
- Higher sensitivity and tolerance to PCR inhibitors

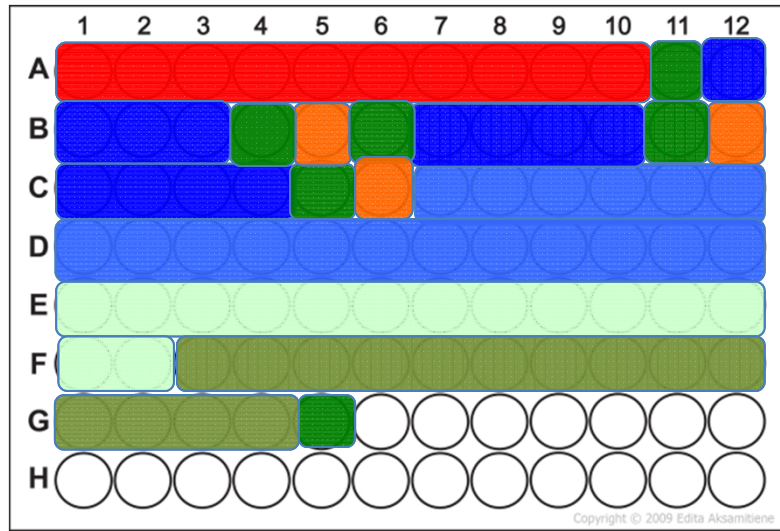
# QFTRIO: Sensitivity, Reproducibility, and Specificity Experiments



X2

		QFTRIO STDs X2 (50,000 5,000 500 50 and 5 pg/μl)
Sensitivity & Reproducibility		5 QFDUO STDs X3 (Last 5 dilutions: 1,850 620 210 68 and 23 pg/μl)
		5 NIST SRM 2372 Dilution series X3 (5,700 570 57 5.7 and 0.57 pg/μl)
		3 Male Genomic DNA dilution series X3 ( 500, 50 and 5 pg/μl)
		3 Female Genomic DNA dilution series X3 ( 500, 50, and 5 pg(μl)
Reproducibility Dynamic range		3 GHEP Proficiency samples X3
		2 High DNA concentrated samples X3 (> 100 ng)
Specificity		21 Bacterial DNAs
		8 Negative PCR Controls
		4 Positive PCR Controls (DNA 007; input 100 pg)

# QFTRIO: F/M Mixtures and Forensic samples

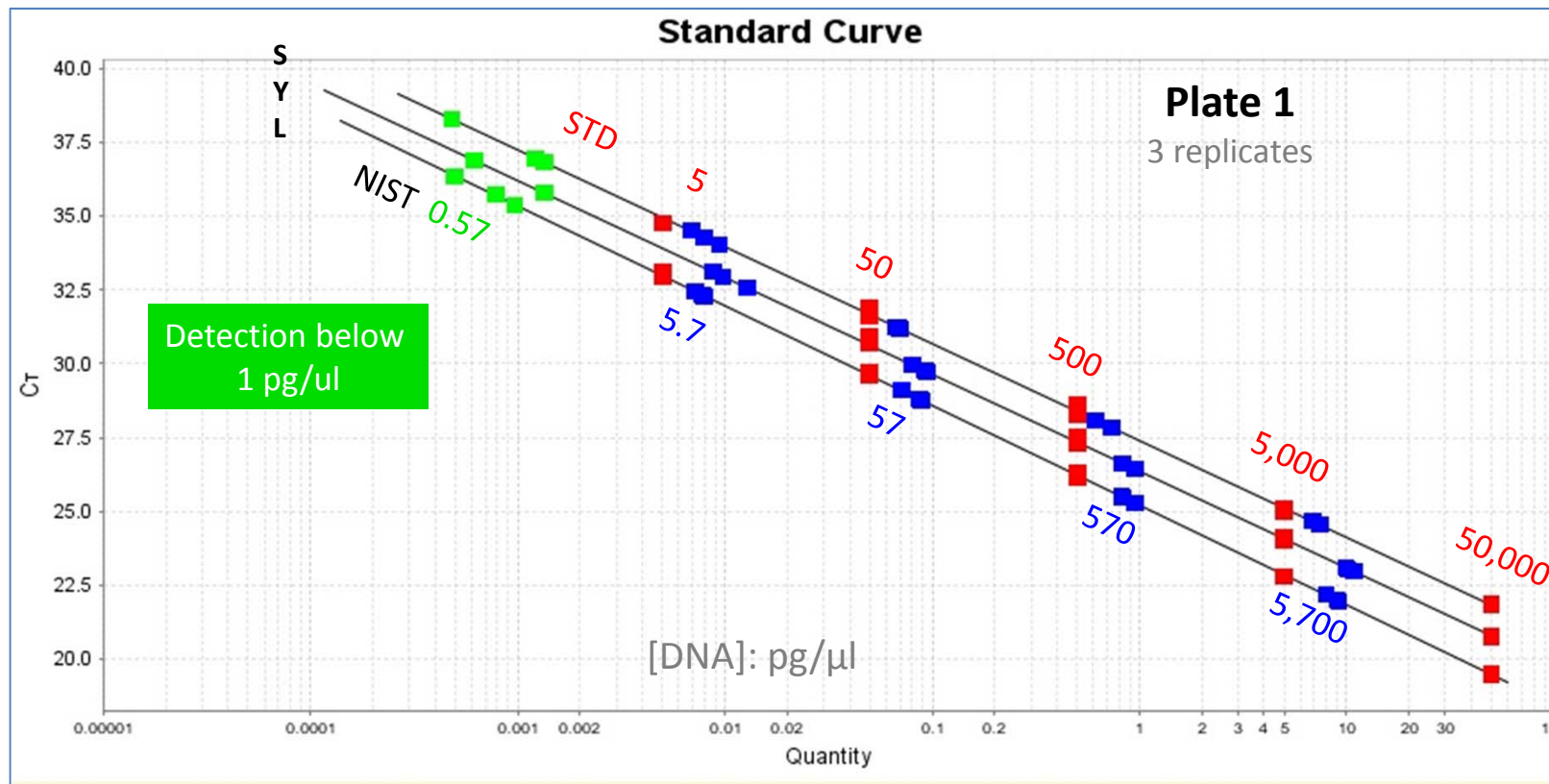


- |                  |   |   |
|------------------|---|---|
| DNA Mixtures     | <span style="color: red;">■</span>        | QFTRIO STDs X2 (50,000, 5,000, 500, 50, and 5 pg)                       |
|                  | <span style="color: blue;">■</span>       | 4 Female/Male DNA Mixtures X3 (100:1 1,000:1 2,000:1 and 4,000:1)       |
|                  | <span style="color: lightblue;">■</span>  | 9 Body Fluid Mixtures X2 (GHEP & GEDNAP samples) (2 and 3 contributors) |
| Forensic samples | <span style="color: lightgreen;">■</span> | 7 Forensic DNA samples X2: Low Level Y-DNA content and Inhibition       |
|                  | <span style="color: olive;">■</span>      | 7 Forensic DNA samples X2: Degraded DNA from bones and FFPE samples     |
|                  | <span style="color: green;">■</span>      | 6 Negative PCR Controls   |
|                  | <span style="color: orange;">■</span>     | 3 Positive PCR Controls (DNA 007; input 100 pg)                         |

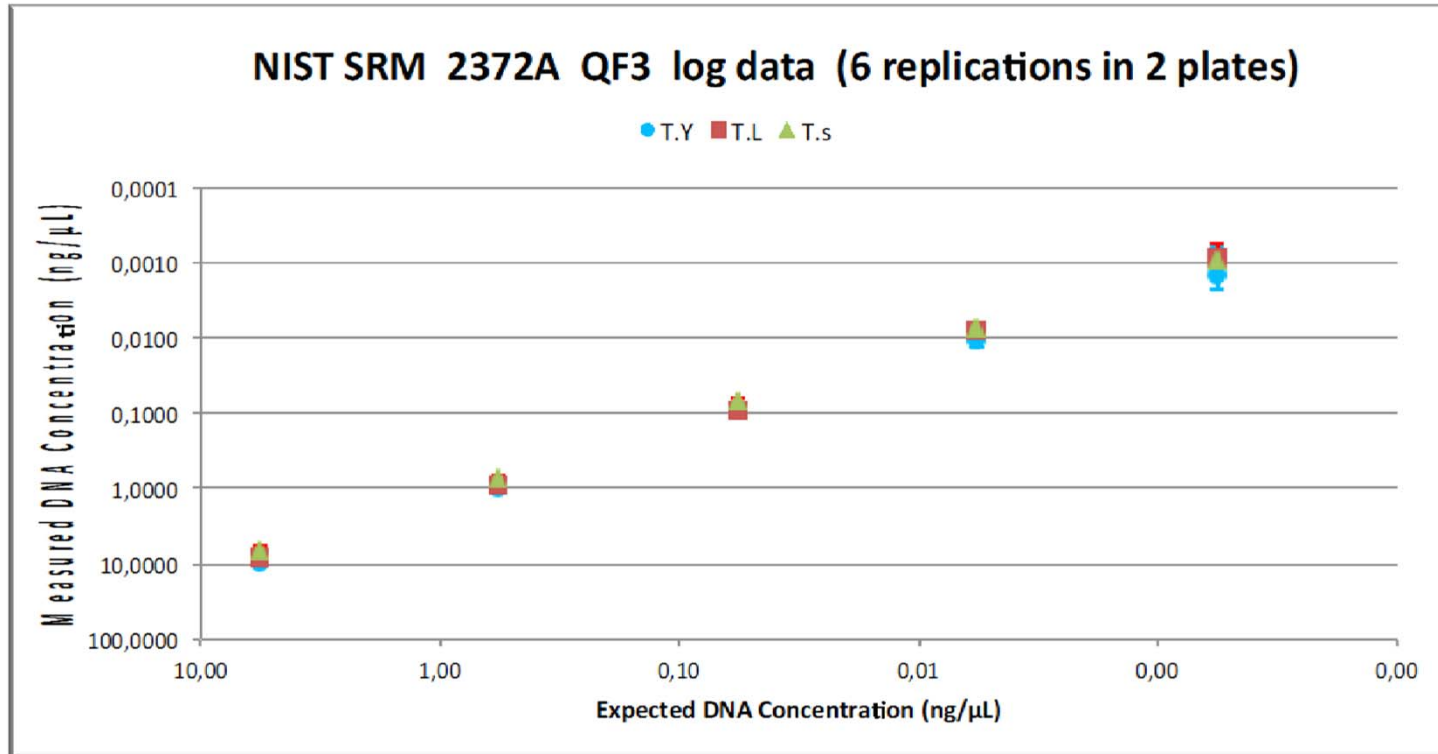
# QFTRIO: NIST SRM 2372A Quantitative Data

## QFTRIO Standard Curves (Small auto, Y, and Large auto)

■ NIST SRM 2372A, ■ QFTRIO Standards



# QFTRIO: NIST SRM 2372A Quantitative Data



# QFTRIO: Sensitivity to detect male DNA from mixtures

Mixture	F/M ratio	Female DNA (9974)	Male DNA (2800)	Human Male Target results (ng/ul)
---------	-----------	-------------------	-----------------	-----------------------------------

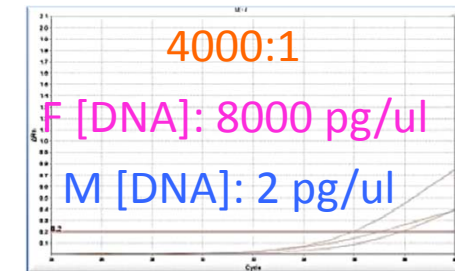
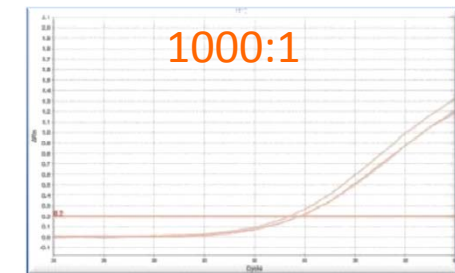
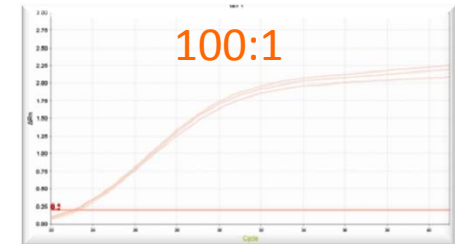
MIX_1-1	100:1	5 ul (10 ng/ul)	5 ul (0.1ng/ul)	0.0703
MIX_1-2		5 ul (10 ng/ul)	5 ul (0.1ng/ul)	0.0772
MIX_1-3		5 ul (10 ng/ul)	5 ul (0.1ng/ul)	0.0812

MIX_2-1	1000:1	5 ul (10 ng/ul)	5 ul (0.01ng/ul)	0.0062
MIX_2-2		5 ul (10 ng/ul)	5 ul (0.01ng/ul)	0.0063
MIX_2-3		5 ul (10 ng/ul)	5 ul (0.01ng/ul)	0.0083

MIX_3-1	2000:1	10 ul (10 ng/ul)	5 ul (0.01ng/ul)	0.005
MIX_3-2		10 ul (10 ng/ul)	5 ul (0.01ng/ul)	0.0048
MIX_3-3		10 ul (10 ng/ul)	5 ul (0.01ng/ul)	0.0052

MIX_4-1	4000:1	10 ul (10 ng/ul)	2.5 ul (0.01ng/ul)	0.0004
MIX_4-2		10 ul (10 ng/ul)	2.5 ul (0.01ng/ul)	0.0007
MIX_4-3		10 ul (10 ng/ul)	2.5 ul (0.01ng/ul)	0.0014

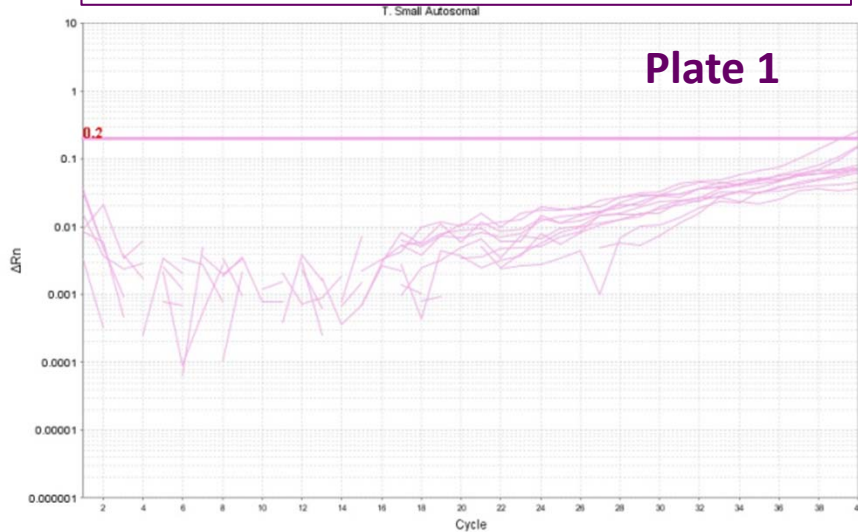
Amplification Plots  
Male DNA Target





# QFTRIO: Specificity against Bacterial DNA

## Amplification Plots Small Autosomal Target



- 01** *Neisseria meningitidis* (2 ng/ul)
- 02** *Streptococcus pneumoniae* (2 ng/ul)
- 03** *Streptococcus agalactiae* (serogrupo B) (2 ng/ul)
- 04** *Staphylococcus aureus* (2 ng/ul)
- 05** *Escherichia coli*, (2 ng/ul)
- 06** *Bordetella pertussis* (10 ng/ul)
- 07** *Bordetella parapertussis* (10 ng/ul)
- 08** *Alcaligenes faecalis* (10 ng/ul)
- 09** *Acinetobacter calcoaceticus* (10 ng/ul)
- 10** *Flavobacterium odoratum* (10 ng/ul)
- 11** *Morganella morganii* (10 ng/ul)
- 12** *Pseudomonas stutzeri* (10 ng/ul)
- 13** *Providencia stuartii* (10 ng/ul)
- 14** *Proteus mirabilis* (10 ng/ul)
- 15** *Citrobacter freundii* (10 ng/ul)
- 16** *Pseudomonas aeruginosa* (10 ng/ul)
- 17** *Salmonella enteritidis* (10 ng/ul)
- 18** *Neisseria lactamica* . (10 ng/ul)
- 19** *Shigella flexneri* (10 ng/ul)
- 20** *Serratia marcencens* (10 ng/ul)
- 21** *Haemophilus Influenzae* (serotipo b)(2 ng/ul)

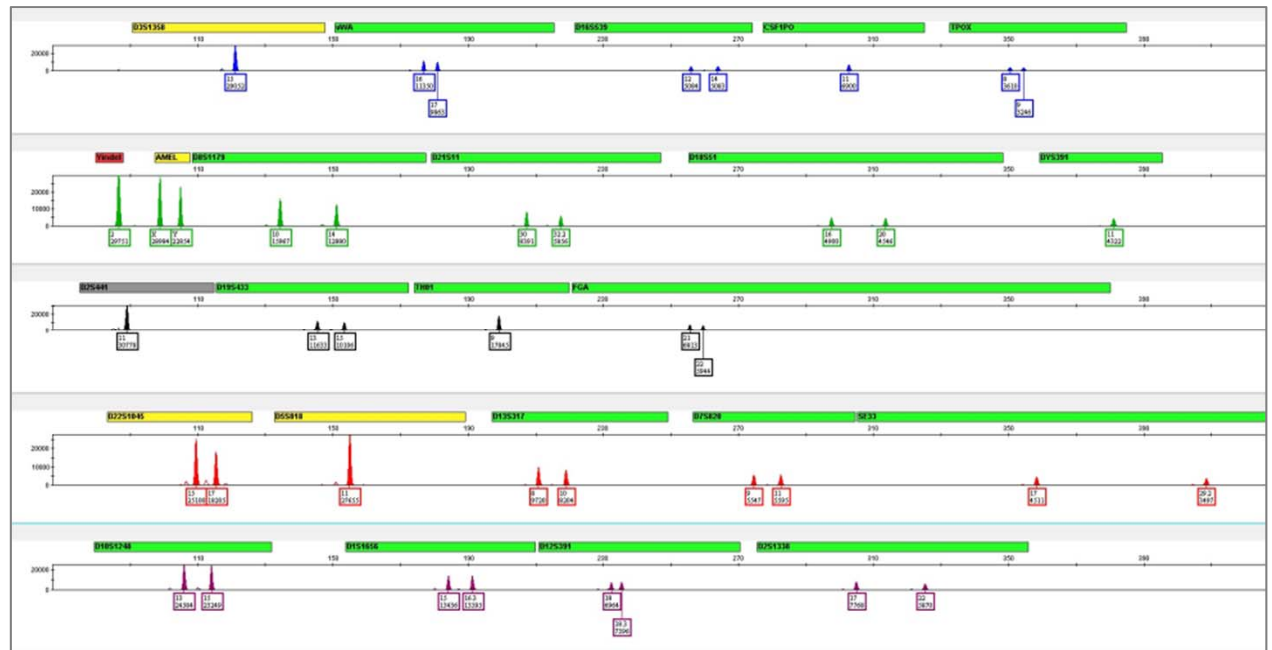
# QFTRIO: Forensic Samples with PCR inhibitors

## Bloodstain on leather

QFDUO (ng/ul)	
Human DNA	N.D.
Male DNA	N.D.
IPC	N.D.

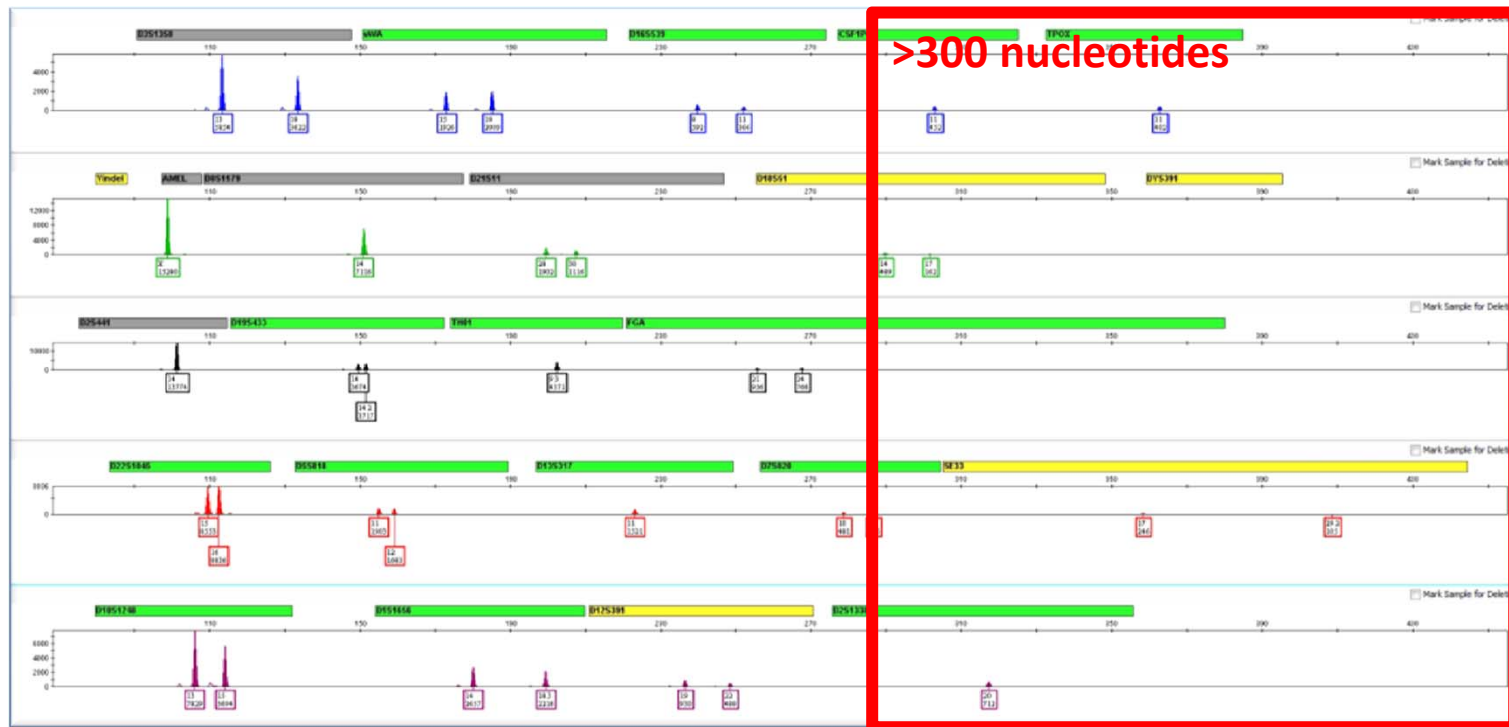
QFTRIO (ng/ul)	
Large Auto DNA	N.D.
Small Auto DNA	2.2778
Male DNA	1.6993
IPC	N.D.

## Globalfiler Full Profile



# QFTRIO: Forensic Samples with Degraded DNA

QFTRIO (ng/ul)		
Femur	Large Auto DNA	<b>0.0438</b>
	Small Auto DNA	<b>0.3596</b>
	Male DNA	<b>N.D.</b>
	IPC	<b>27.260</b>
	DI	<b>8.2173</b>



Globalfiler

# QFTRIO: Forensic Samples with Degraded DNA

QFTRIO (ng/ul)		
Femur	Large Auto DNA	0.0061
	Small Auto DNA	0.3011
	Male DNA	N.D.
	IPC	27.333
	DI	49.098



Globalfiler

# QFTRIO: Forensic Samples with Degraded DNA

QFTRIO (ng/ul)		
FFPE	Large Auto DNA	0.0002
	Small Auto DNA	0.0169
	Male DNA	0.0141
	IPC	27.206
	DI	<b>102.10</b>



Globalfiler

## QFTRIO-Y: A Sensitive method for Male DNA screening: Comparison with Y-STR profiling sensitivity

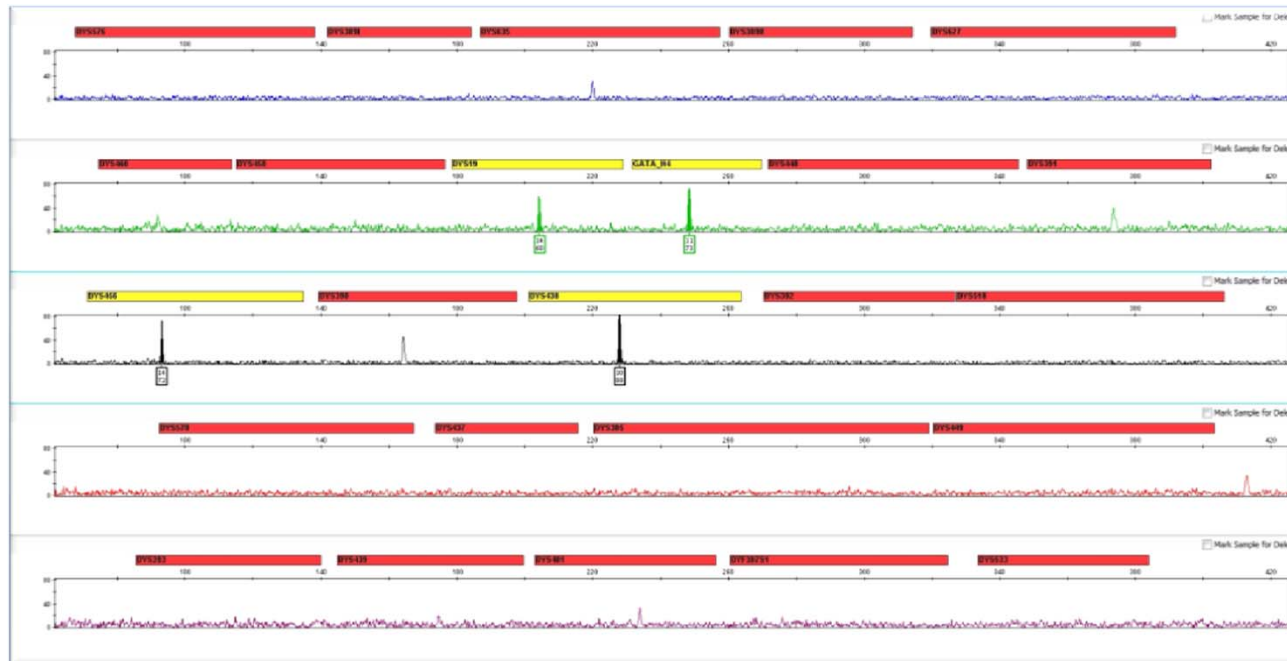
245 Forensic DNA samples with male DNA concentration below 10 pg / $\mu$ l (QFTRIO Y-Target) and also analysed by Y-STR-PCR (YFilerPlus) and CE detection were reviewed to compare and calibrate the detection limits of each method

The samples belong to criminal cases in which **the investigation of male cell remains** is requested. Body samples (vaginal swabs, pubis swabs, perineal swabs, anal swabs, oral swabs, fingernails,..) were mainly analyzed but also several biological stains on different substrates

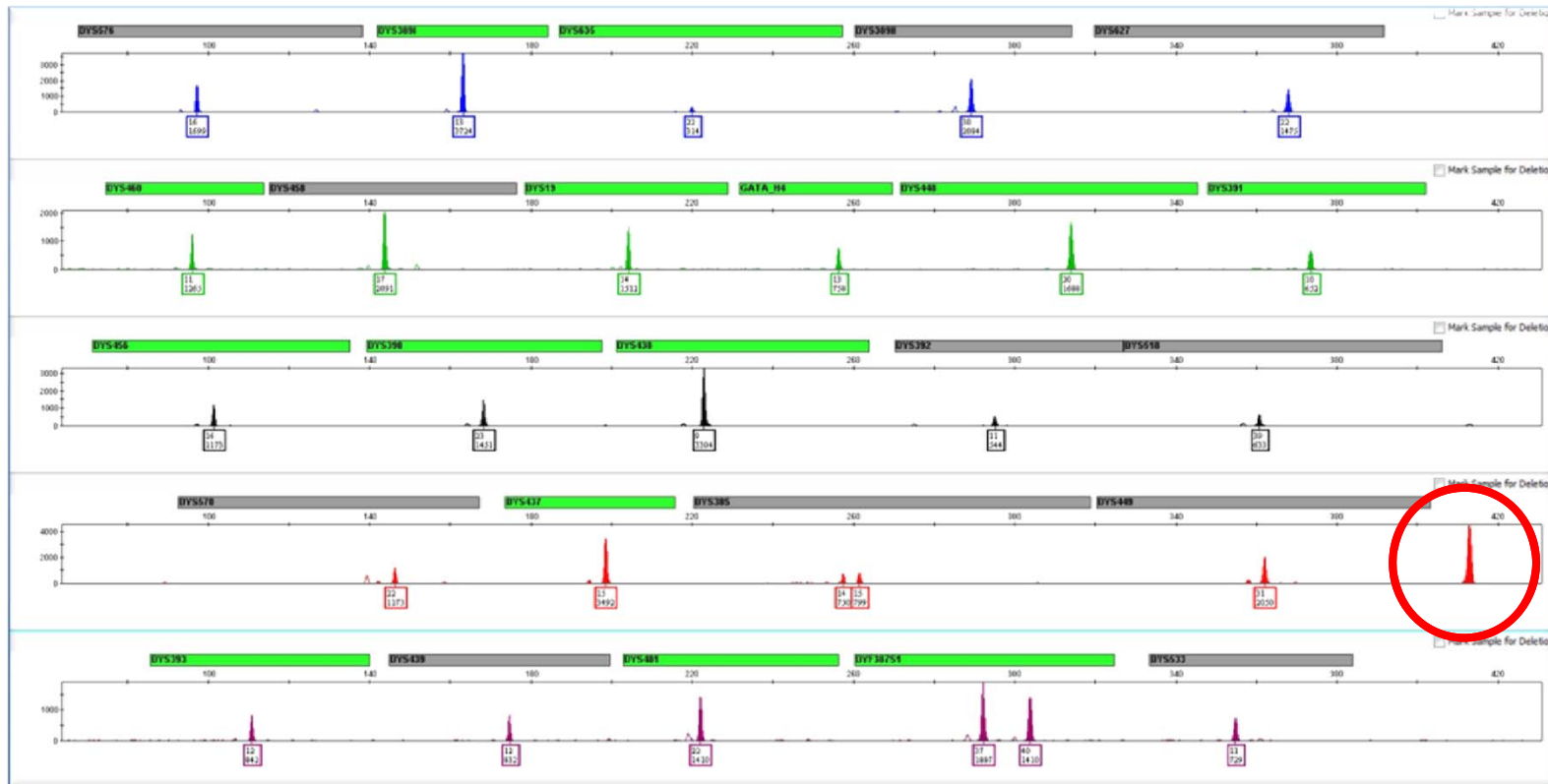
DNA samples	Male DNA Concentration	Y-STR Profiling success
<b>164</b>	<b>&lt; 1 pg/<math>\mu</math>l</b>	<b>N.D. or Inconclusive results</b>
<b>81</b>	<b>10 -1 pg/<math>\mu</math>l</b>	<b>59 % ( mainly partial and/or low signal Y-STR profiles)</b>

# Samples with Qftrio-Y results under 1 pg/ $\mu$ l rendered Negative (or Inconclusive) Y-STR profiling results

QFTRIO (ng/ $\mu$ l)		
Vaginal swab	Large Auto DNA	1.0476
	Small Auto DNA	1.0871
	Male DNA	0.0007
	IPC	27.2851
	DI	1.0377



# 59 % of Samples with Qftrio-Y results between 10- 1 pg/μl rendered Y-STR profiles (mainly partials) Endocervix Swab



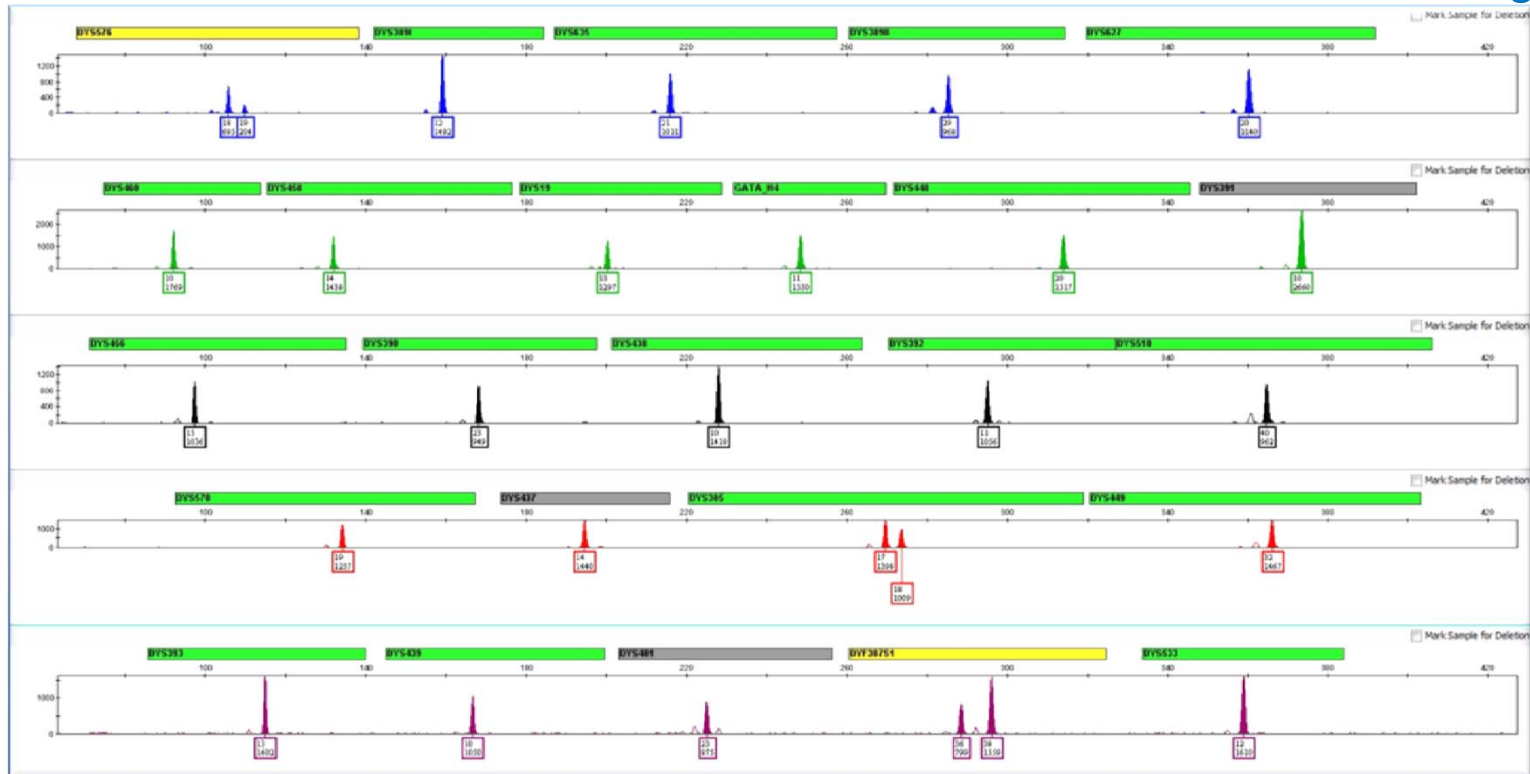
QFTRIO (ng/ul)	
Large Auto DNA	37.0246
Small Auto DNA	37.4354
Male DNA	0.010
IPC	29.1755
DI	1.0111

F:M DNA ratio: 3700 : 1

Yfiler Plus DNA input (10μl)	
Male DNA	100 pg
Female DNA	370 ng



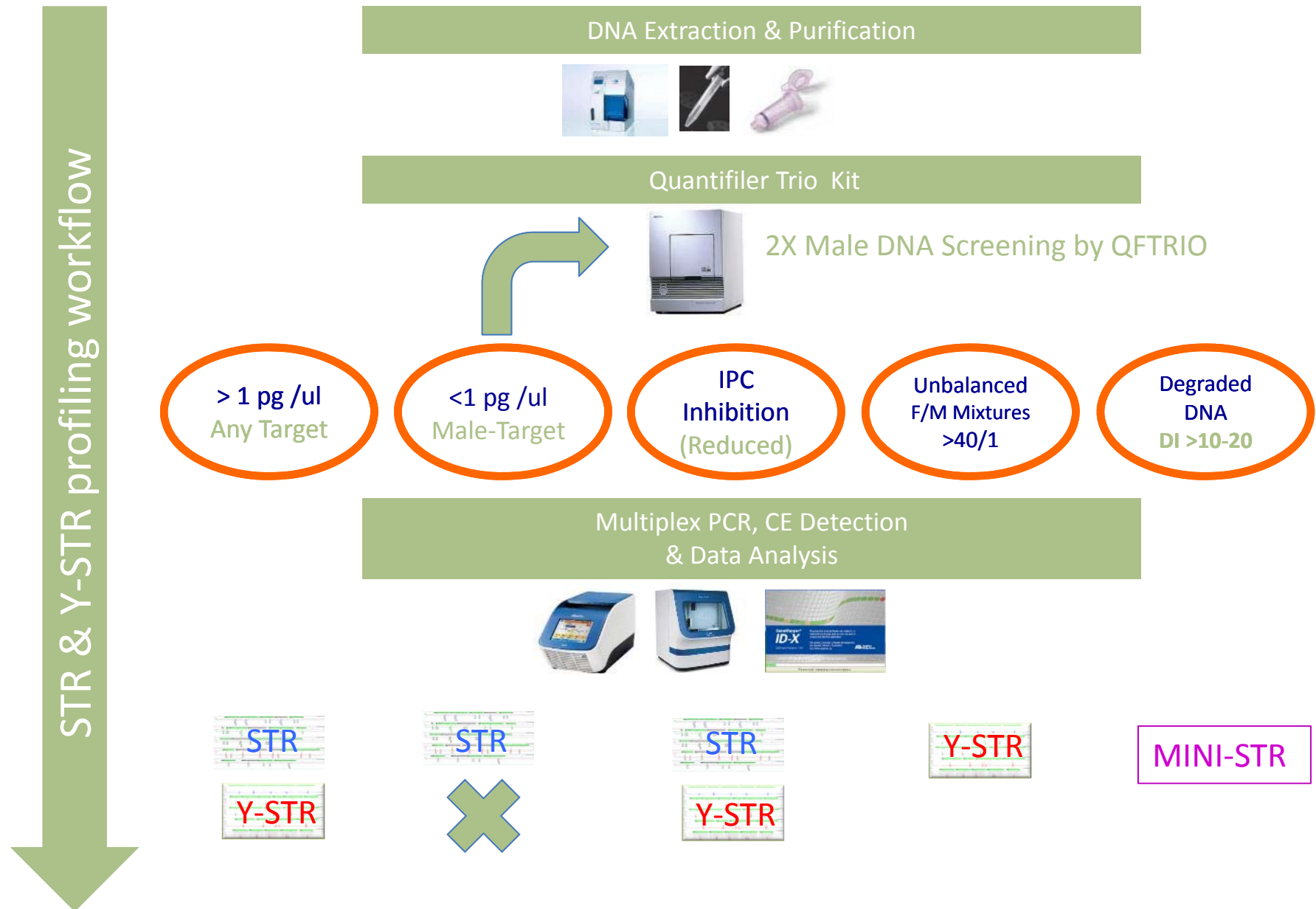
# 59 % of Samples with Qftrio-Y results between 10- 1 pg/ $\mu$ l rendered Y-STR profiles (mainly partials) Neck Swabbing



QFTRIO (ng/ $\mu$ l)	
Large Auto DNA	0.0146
Small Auto DNA	0.0208
Male DNA	0.0067
IPC	27.6831
DI	1.4193

Yfiler Plus DNA input (10 $\mu$ l)	
Male DNA	67 pg

# Our present on Human DNA Quantification and STR profiling workflow



# Acknowledgements

Design, Experiments and Data Analysis

Esperanza González

Ana Carrillo

M<sup>a</sup> Carmen Periñan

Esther Navajo

Antonio Alonso

STR and Y-STR EPGs

Pablo Martin

Figure in slide 7

Pedro Barrio

Special thanks also to the staff of our microbiology laboratory for providing the microbial genomic DNA samples used in the specificity study.



INTCF-Madrid  
SERVICIO DE  
BIOLOGIA

When used for purposes other than Human Identification the instruments and software modules cited are for Research Use Only. Not for use in diagnostic procedures.

Speaker was provided travel and hotel support by Thermo Fisher Scientific for this presentation, but no remuneration