



Development of a forensic DNA quantification assay using digital PCR



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Disclosures

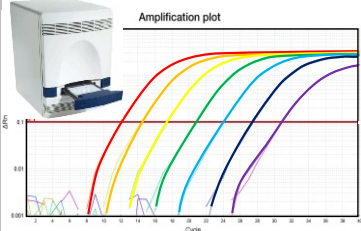
This work is the result of a forthcoming Collaborative Research and Development Agreement (CRADA) between the Federal Bureau of Investigation Federal Laboratory and QIAGEN Sciences Inc.

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Quantification of nuclear and mito DNA

- Currently, two separate protocols in the DNA Casework Unit of the FBI Laboratory



Double

- Standard curves
- Instrument Time
- Reagent Costs
- Quality Control

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dPCR

- Fundamentally still PCR
- Thousands of ~0.8nL endpoint reactions

The diagram illustrates the dPCR workflow in four stages: 1. Sample & Assay: A pipette dispenses a sample into a microfluidic chip. 2. Partition: The sample is divided into thousands of small droplets. 3. PCR: Each droplet undergoes a PCR reaction. 4. Count: The resulting fluorescence is measured, and a bar graph shows the distribution of droplets with and without target sequences.

- Signals (+/-) counted and concentration determined by Poisson statistics

Photo: Qiagen Integrated RNA Technologies

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QIAcuity instrument and plates

- Nanowell plates
- QIAcuity dPCR instrument

The image shows three nanowell plates with different colored wells (blue, green, red) and the QIAcuity dPCR instrument, which is a white, boxy device with a screen and a sample tray.

Photo: Qiagen

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A combined mitochondrial and nuclear assay

Pentaplex Assay

The diagram shows a Pentaplex Assay with five channels. The channels are labeled: Mito Small (green), Mito Large (yellow), Inhibition (orange), Nuclear (red), and Y-Chrom (pink). The y-axis is labeled RFU (Relative Fluorescence Units). The assay shows a distribution of signals for each channel, with a red box highlighting the Mito Large and Nuclear channels.

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Initial development

- Mito Short
 - Mito quantification
 - 105bp
 - FAM
- Mito Long
 - Mito degradation
 - 316bp
 - VIC
- IPC
 - Assay inhibition
 - 65bp
 - NED

Development of a triplex mtDNA qPCR assay to assess quantification, degradation, inhibition, and amplification target copy numbers
 Mark F. Kavlick*
 *Correspondence and Requests for Materials: Research Unit, Laboratory Division, National Agency of Biotechnology, 2501 Investigator Parkway, Quantico, VA 22135, United States

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Singleplex comparison to qPCR

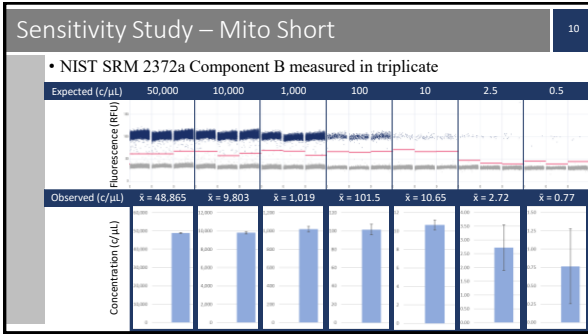
	QiAcuity	7500		
	Copy/ μ L RXN	Ct Value	Copy/ μ L Sample	Copy/ μ L RXN
FAM - Mito Small	46.8	27.88	971.5	48.6
VIC - Mito Large	43.5	30.8	~900	~45
NED - IPC	53	27.95	~1000	~50

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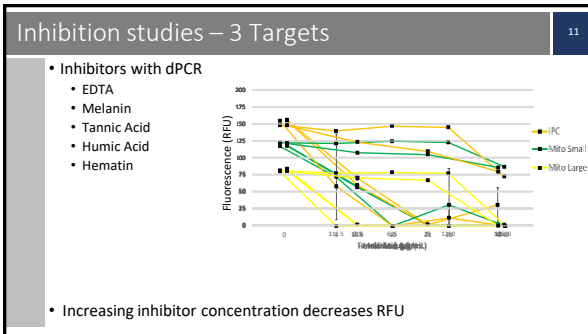
Optimized mitochondrial triplex on dPCR

- Forward and reverse primers and probe concentrations were optimized for each target
- Optimized triplex conditions resulted in RFU separations seen below

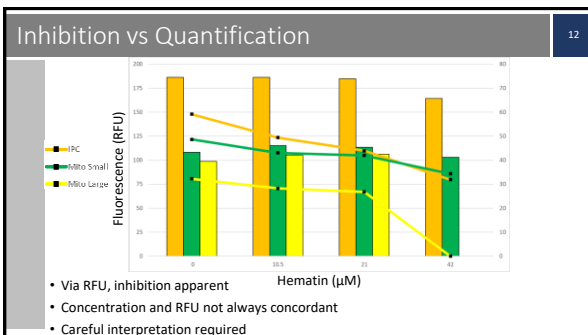
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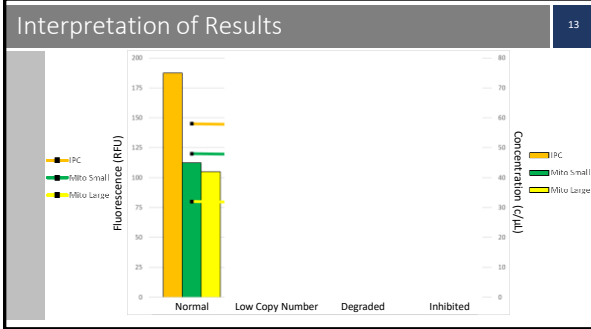
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Conclusions

- Mitochondrial triplex on dPCR
- Highly sensitive down to 0.5 c/μL
- Sensitive to 5 of 5 inhibitors tested
- Digital PCR works for forensics and clinical

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Future Directions

- The ultimate goal is a pentaplex assay
 - Add small Nuclear and Y targets
 - Difference between nuclear and mito copy number
 - Saturation of mito assay on instrument
- Developmental validation
- Test on casework type samples
 - Hair
 - Bone


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Acknowledgements 16

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References 18

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