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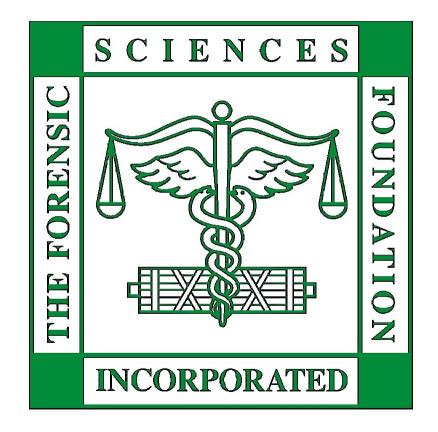
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## On combining microRNA analysis with DNA profiling in a single stream process

Donny van der Meer MSc Supervisor: Dr Graham Williams University of HUDDERSFIELD Inspiring tomorrow's professionals

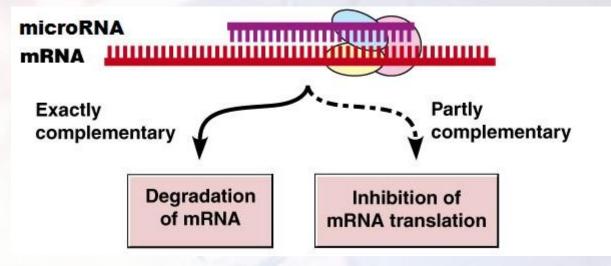


### FSF Emerging Forensic Scientist Award Oral Presentation

What are microRNAs and why are we interested in them?

Small (~22nt) non-coding RNAs Regulate mRNA expression

Advantages for forensic science Stable High expression levels Sensitive and specific detection Co-extracted with DNA





MicroRNAs can be used for body fluid identification

More than 2500 microRNAs in humans Tissue specific expression patterns

Previously identified markers Blood: miR-16a, miR-142 and miR-451a Saliva: miR-203a and miR-205 Semen: miR-10a and miR-135a Vaginal material: miR-1260b Control: SNORD44

Hanson, E. K. et al. (2009) Anal Biochem, 387(2), 303–14. Zubakov, D. et al. (2010) Int J Leg Med, 124(3), 217–26 Park, J.-L. et al. (2014) Electrophoresis 35(21-22), 3062-8

#### Improve current methodology with our novel method

#### Current

- microRNAs: RT-qPCR
  - Separate reaction per microRNA

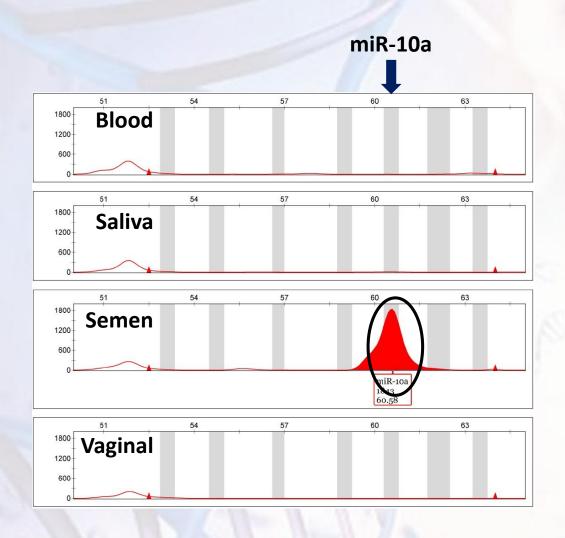
#### Our novel method

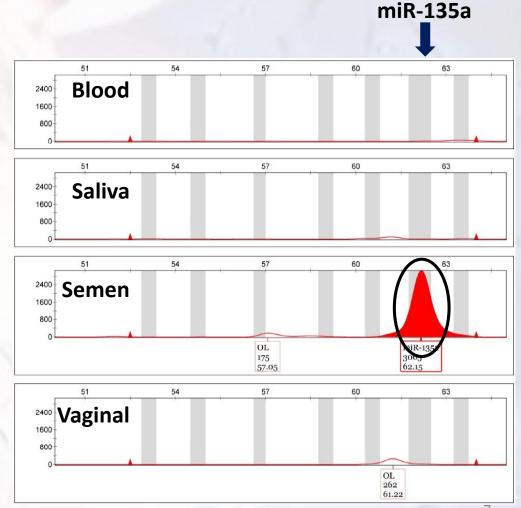
- Analyse microRNAs with capillary electrophoresis (CE)
  - Multiplex microRNAs in single test
  - Possibility to combine microRNA analysis with DNA profiling

#### Methods and materials

- 5 samples of 4 tissue types
  - Blood, saliva, semen and vaginal material
- DNA extraction
- Normalised to 0.5ng/μl human DNA
- Tested for 9 markers
- Multiplex stem-loop reverse transcription
- ROX-labelled primers

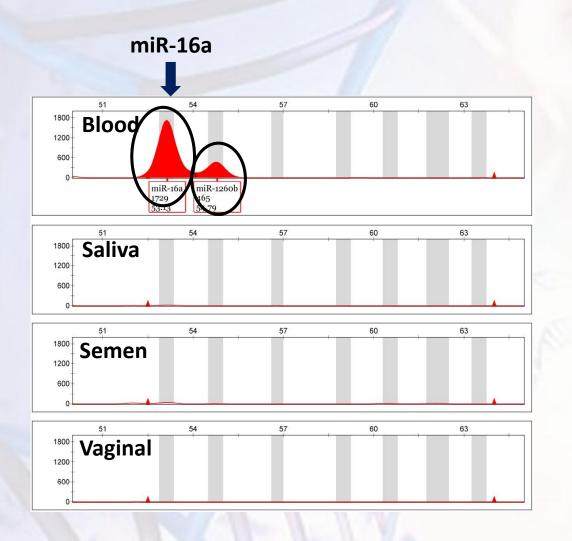
#### miR-10a and miR-135a are exclusively detected in semen

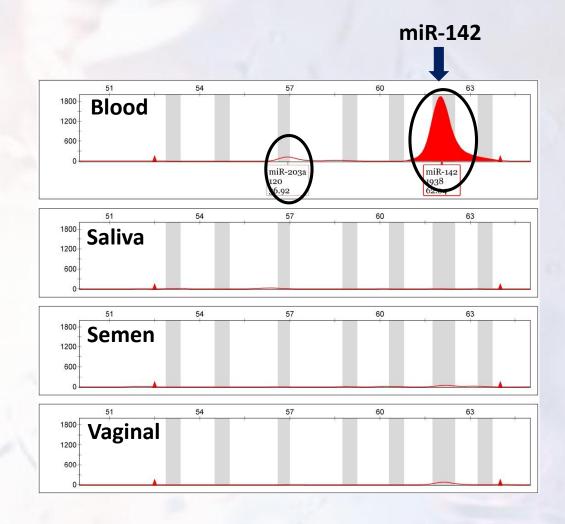




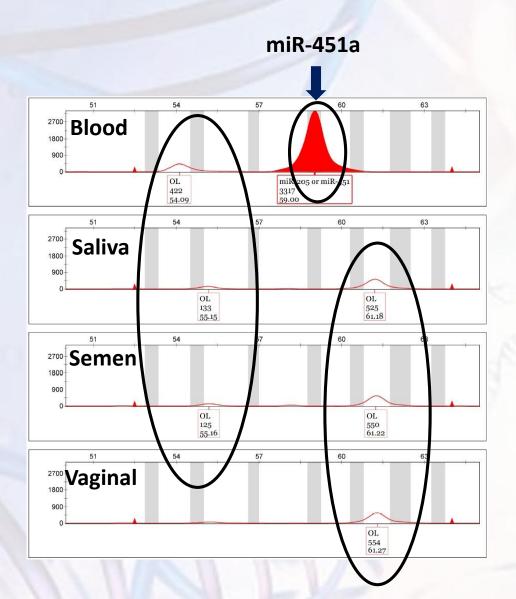
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### miR-16a and miR-142 are exclusively detected in blood





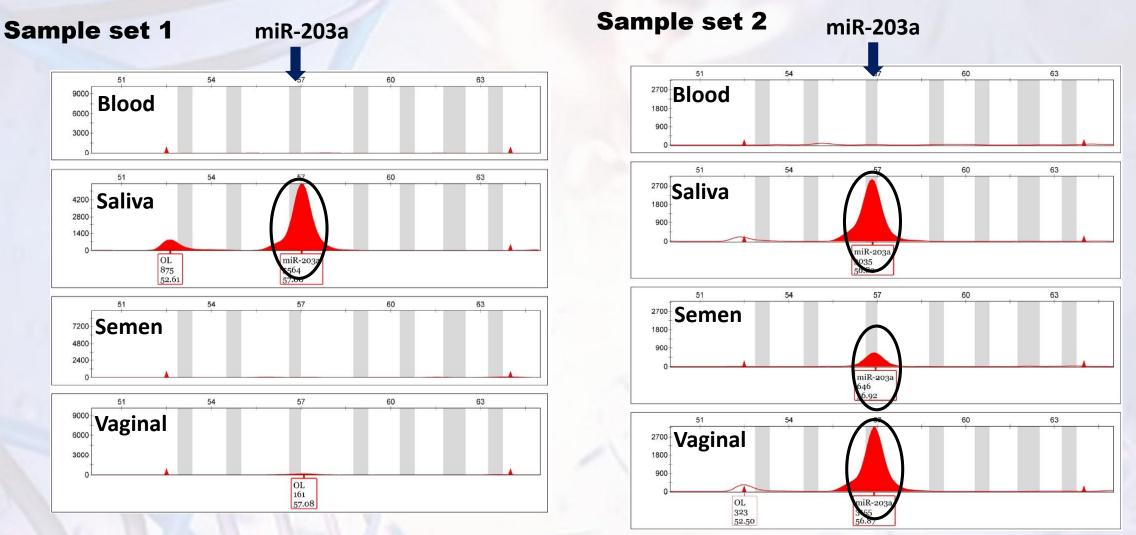
## miR-451a is exclusively detected in blood



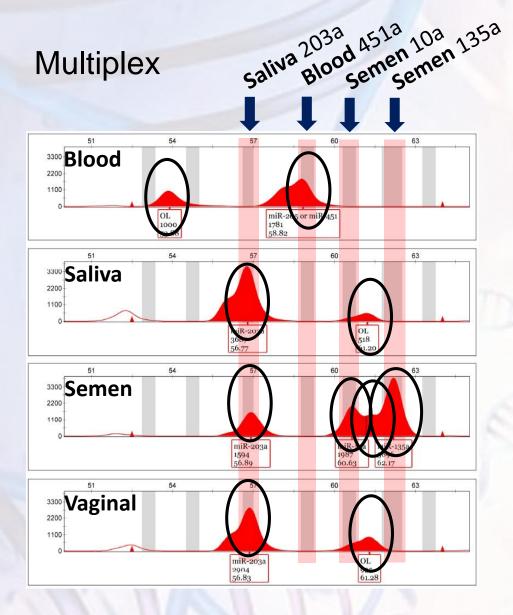
Much lower peaks of by-products found in all tissues

## miR-203a is mainly detected in saliva

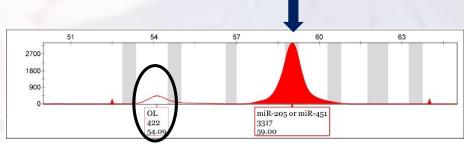
#### **Expressed in epithelial cells**



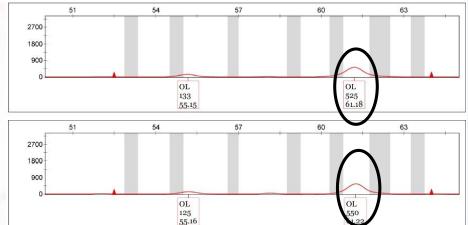
#### Multiplexing multiple markers yields expected results

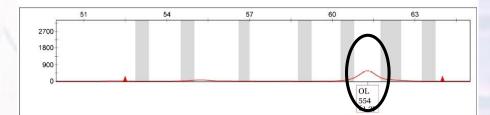


Singleplex



miR-451a

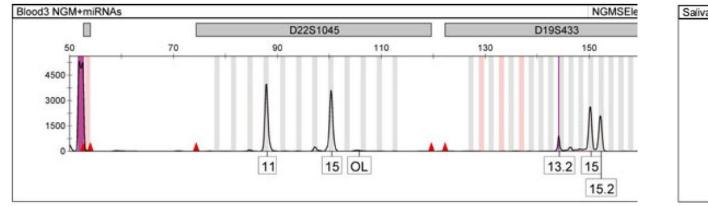


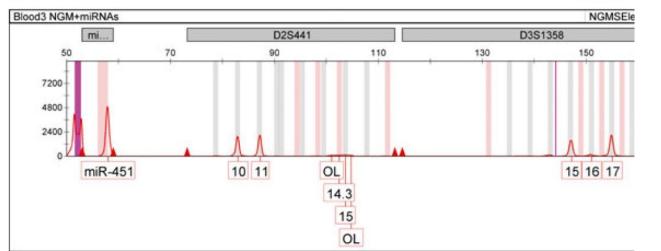


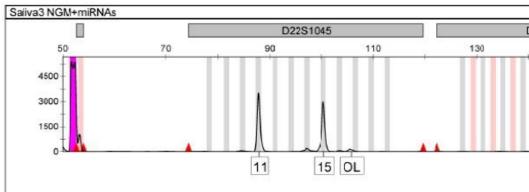
### Multiplex with STR markers

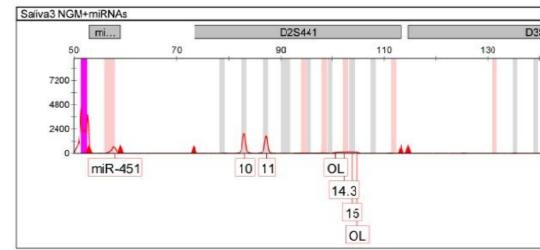
#### Blood

#### Saliva

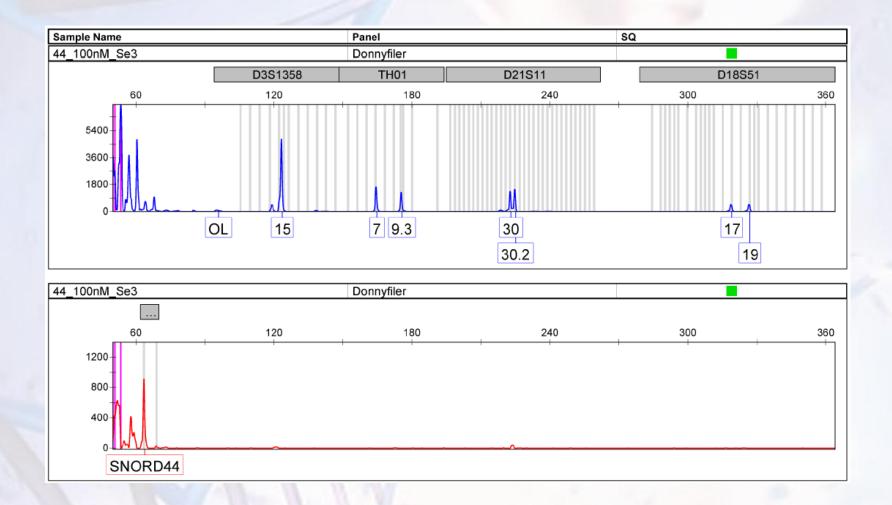








## Multiplex with STR markers



#### Conclusion

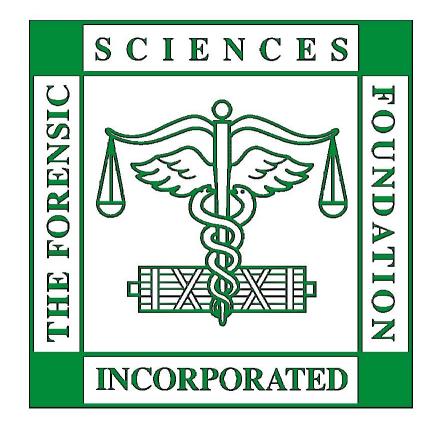
- Analysing microRNAs with CE is viable
- Potential for future single confirmatory test
- Combining microRNA analysis with DNA profiling is technically feasible

#### Future work

- Reduce non-specific amplification
- Physically separate markers
  - Increase product length
- Optimise multiplex reaction
- Combination with DNA profiling

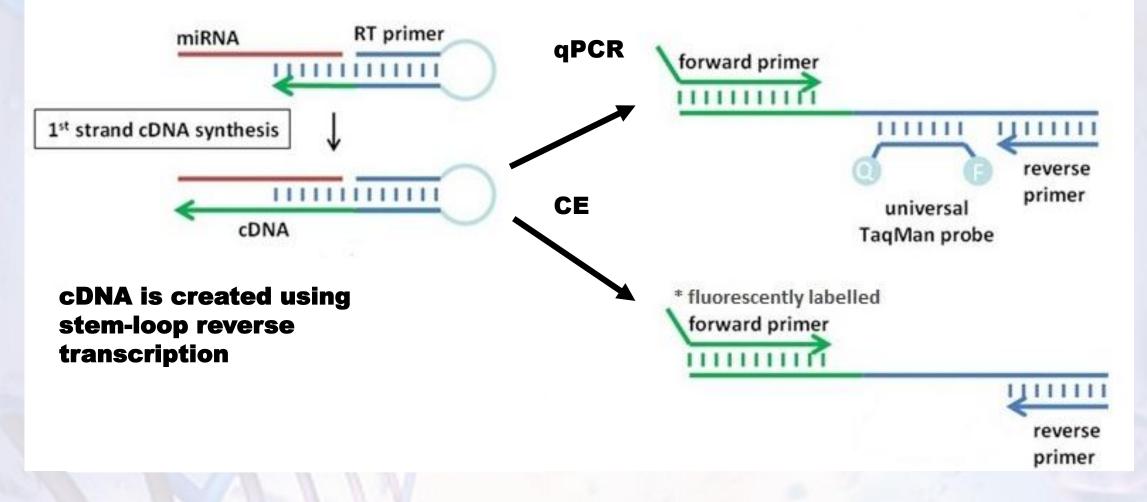
## Thank you

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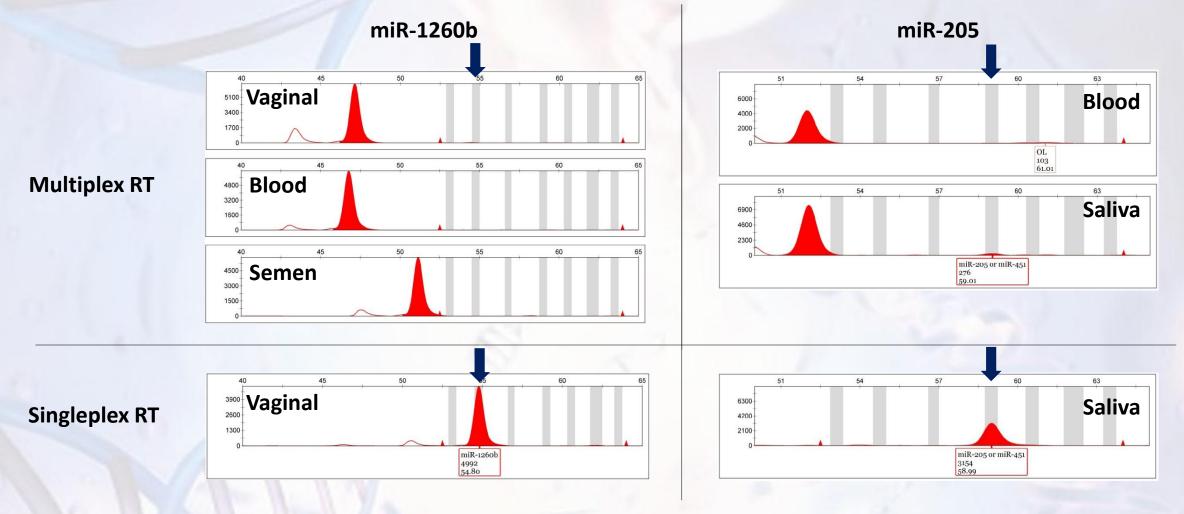
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#### Our workflow



Chen, C. et al. (2005) Nucl Acids Res, 33(20); Jung, U. et al. (2013) RNA, 19, 1864-73

# miR-1260b and miR-205 fail due to multiplexing reverse transcription



## Multiplexing multiple markers yields expected results

