

Multiplex quantitative PCR for detection of *Ehrlichia canis*, *Babesia canis* and canine ACTB gene

Ofer Peleg¹, Gad Baneth², and Shimon Harrus²

¹GenAphora, ²School of Veterinary Medicine, Faculty of Agriculture, Hebrew University of
Jerusalem



GenAphora

- GenAphora
- *Ehrlichia canis*
- *Babesia canis*
- Singleplex Experiments
- Multiplex Experiments



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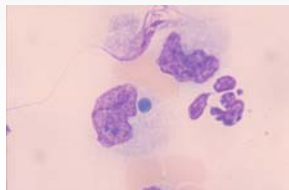
- GenAphora was founded on the basis of computational biology
- The Singleplexer of GenAphora is PERL software that searching for the most suitable and complex primers & probes for qPCR
- The Multiplexer of GenAphora is a PERL software that searching for the best primers & probe combinations for multiples qPCR
- Currently, construction of singleplex and multiplex assays for pathogens detection in veterinary medicine



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Ehrlichia canis

- Small gram-negative bacteria
- Infecting monocytes
- Clusters called morulae – hard to visualize
- Disease prevalent worldwide
- The rickettsia can persist in infected dogs for months & years
- Variable clinical signs
- May be fatal

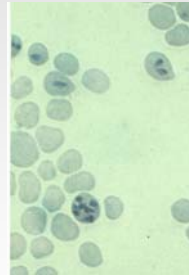


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Babesia species

- *Babesia* spp. are protozoa infecting erythrocytes, causing hemolysis > resulting in anemia
- *B. canis canis*, *B. canis vogeli*, *B. canis rossi* and *B. gibsoni* are commonly known organisms that infect dogs



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Common Vector

- Both *E. canis* & *B. canis vogeli* are transmitted by *R. sanguineus*
- Ticks found throughout Asia, Africa, Europe, the Middle East, and North America



Rhipicephalus sanguineus
The brown dog-tick



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Aim

Construction of a multiplex qPCR for
E. canis & *B. canis vogeli*
(ACTB as a reference gene)

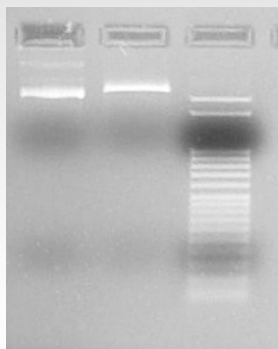


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M & M

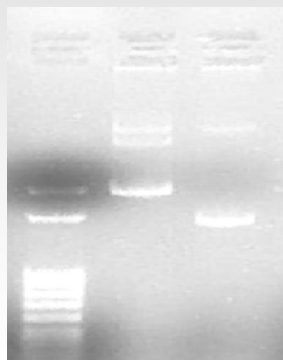
Plasmid Construction

Pst I Cut



pGMT pGMT-16S Ladder

Circular

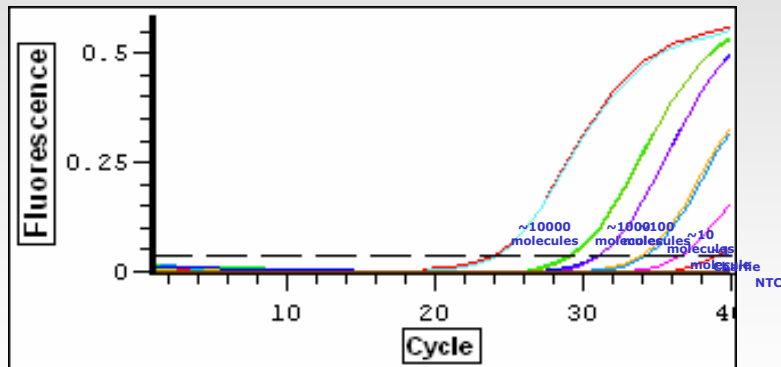


Ladder pGMT-16S pGMT



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E. canis Assay Sensitivity (dual labeled probe)

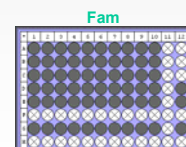
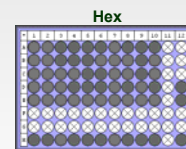
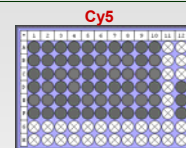
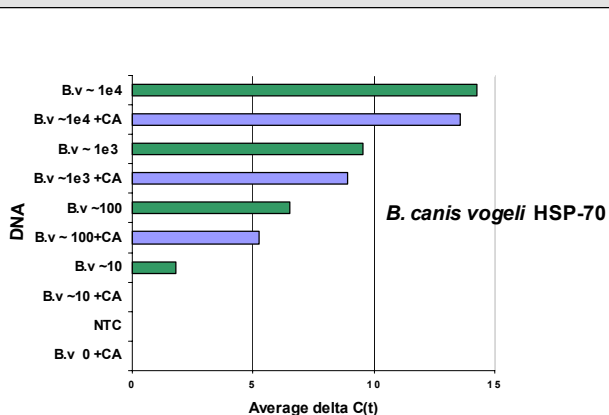


• Same performed with *Babesia canis vogeli*



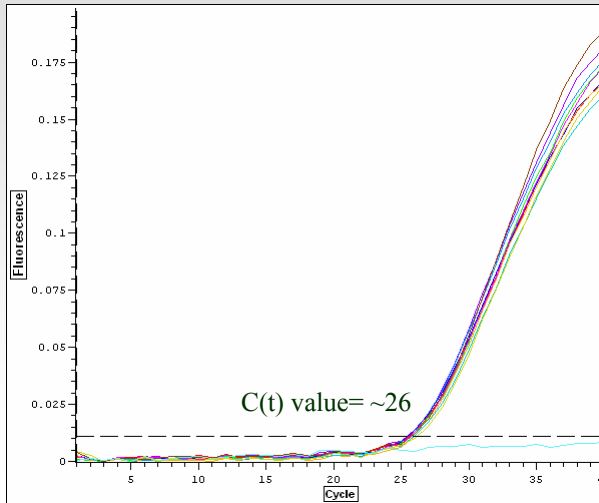
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Duplex: *Babesia canis vogeli* hsp70 (Hex) with and without 10^4 /ml canine β actin (Cy5)

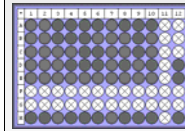


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Triplex: $\sim 10^4$ /ml *B. canis vogeli* hsp70 gene (Hex) in elevated concentrations of *E. canis* 16s RNA gene (Fam) and 10^4 copies/ml canine β actin gene (Cy5)

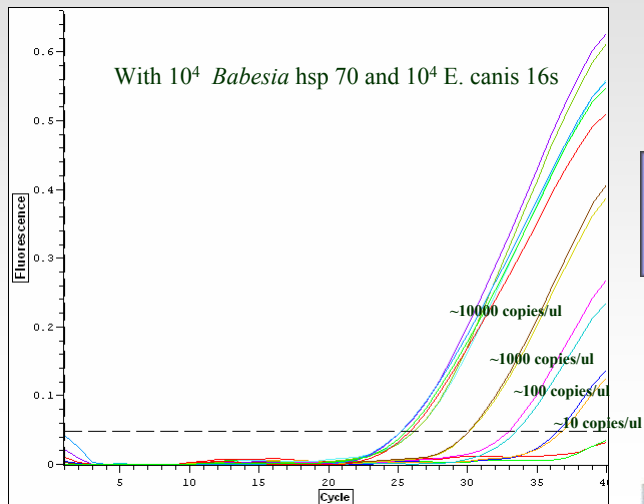


Hex

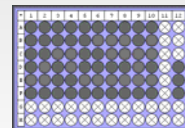


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Elevated concentrations of canine β actin gene (Cy5) with or without *B. canis vogeli* hsp70 gene & *E. canis* 16s RNA gene

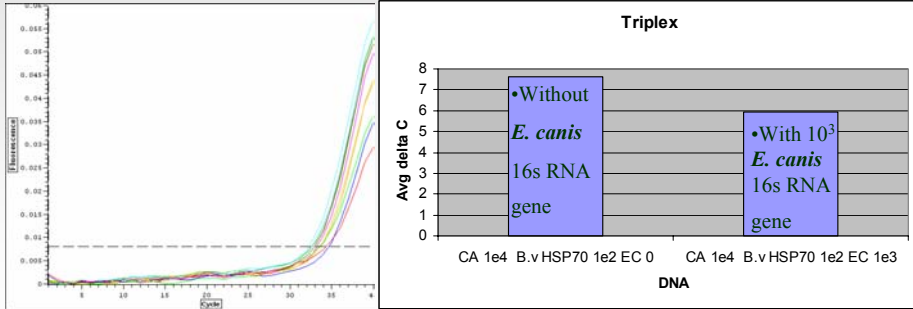


Cy5



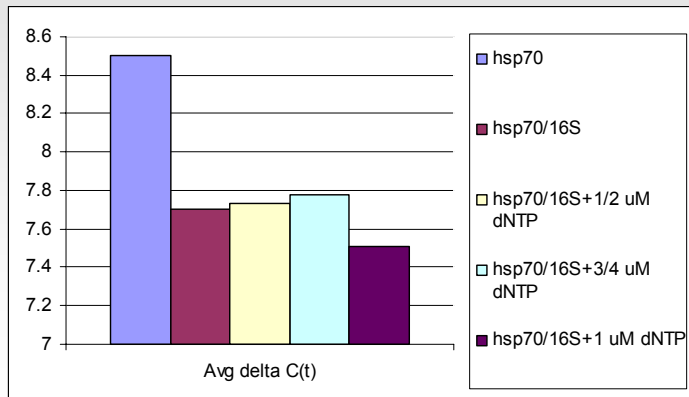
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Triplex: $\sim 10^2$ copies/ μ l *B. canis vogeli* hsp70 gene in elevated concentrations of *E. canis* 16s RNA gene and 10^4 /ml canine β actin gene



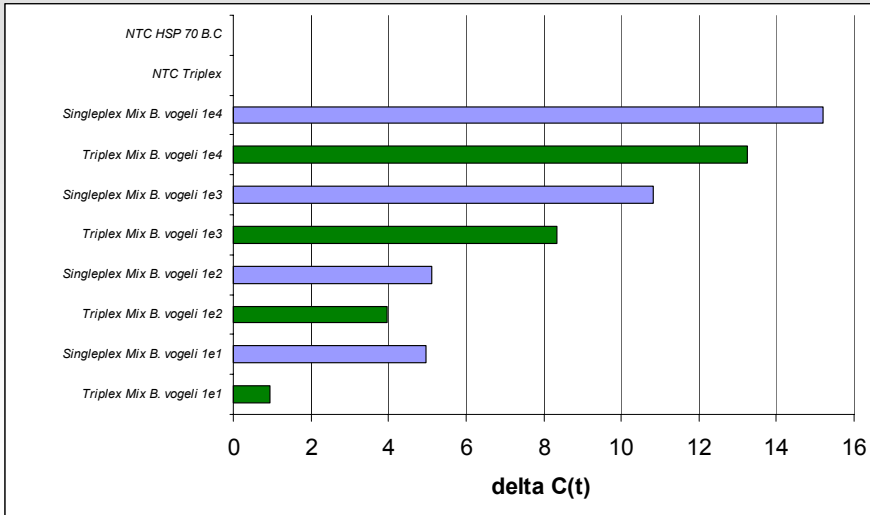
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dNTPs titration



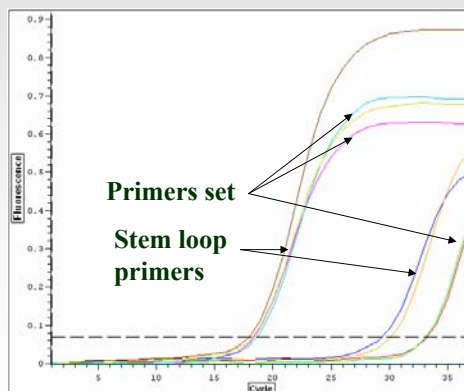
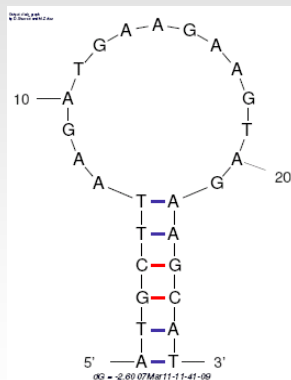
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Singleplex using triplex primer-probe sets



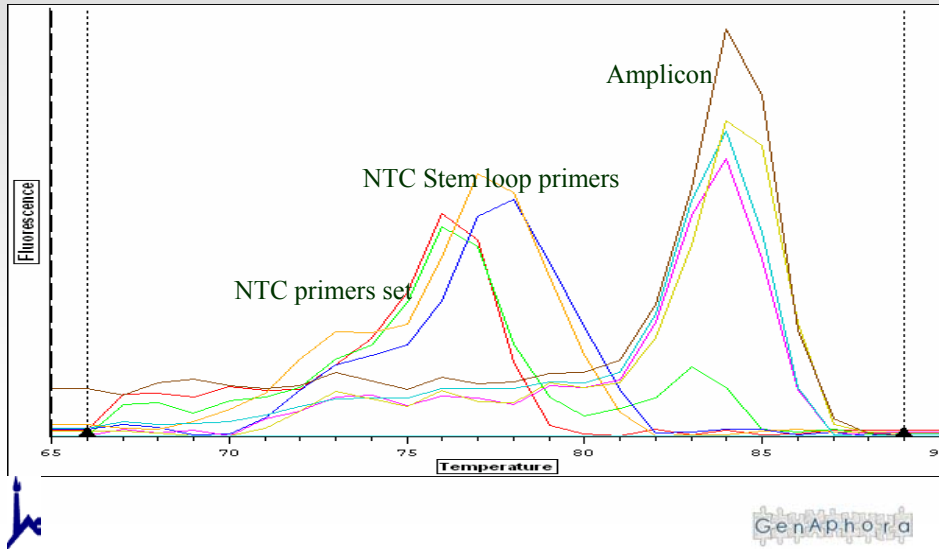
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B. canis vogeli hsp70 stem loop primers

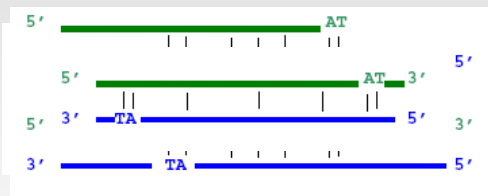


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B. canis vogeli hsp70 stem loop primers melting curve



Primers sequencing



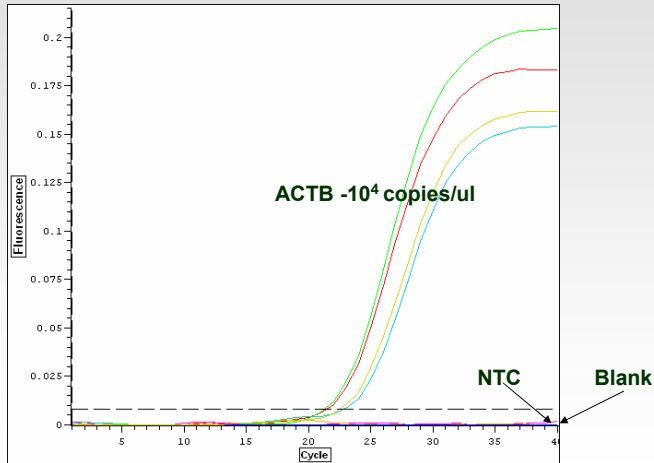
Hot start enzymes?

- These primer-dimers cannot be formed in 61C
- The order of introducing ingredients into the reaction mix tube effects the C(t) value of the NTC
- Therefore, the enzyme is probably not 100% hot start.
- Improving t_m calculation in the primer design script can reduce primer-dimers.

Children, I say plainly, watch out of taking the hot start feature for granted!



Improved primer design primer dimers are not inevitable



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summary

- Triplex qPCR assay was successfully constructed
- Triplex reduces the assay sensitivity by about one order of magnitude
- The cause for this reduction is not system resources overuse
- Primer-dimers is probably the major cause
- We manage to overcome primer-dimers by improving primer design
- Improved multiplex is underway



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