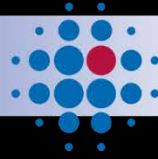


1910–2010



100 JAHRE

FRIEDRICH-LOEFFLER-INSTITUT

FLI

Bundesforschungsinstitut für Tiergesundheit  
Federal Research Institute for Animal Health



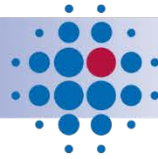
# **22<sup>nd</sup> Annual Workshop**

## **of the National Reference Laboratory for Fish Diseases**

**May 30<sup>th</sup> – 31<sup>st</sup> 2018**

**DTU, Lyngby, Denmark**

(presentation May 31<sup>st</sup> 11:20 – 11:40)

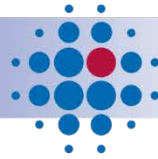


# New research on KHV and KHVD

Dr. Dr. habil. Sven M. Bergmann

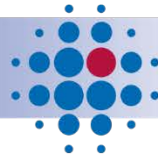
OIE Reference Laboratory for KHVD  
German NRL for KHVD

Certified Veterinary Specialist (consultant) for Aquaculture and for Fisheries

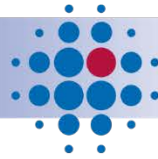


## Research update on:

- 1. KHV diagnostics**
- 2. Treatment against KHVD**
- 3. Confirmation of KHV**



# 1. KHV diagnostics



CiD (EU) 2015/1554 (Sept. 11<sup>th</sup> 2015)  
valid from April 1<sup>st</sup> 2017

Manual of Diagnostic Tests for  
Aquatic Animals 2017, ch 2.3.7.

**EU**

**only to molecular assays**

**OIE  
recommendation**

## 1. KHV qPCR

Gilad et al. 2004

(Bergmann et al. 2010)

- pooling
- health monitoring max. 2 fish

## 2. KHV conventional PCR

Bercovier et al. 2005

(CEFAS / FLI + nested+ seq.)

Engelsma et al. 2013 (+ nested + seq.)

## 1. KHV conventional PCR

Bercovier et al. 2005

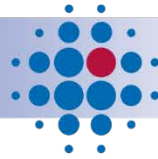
Yuasa et al. 2005

(both + seq.)

## 2. KHV qPCR for surv. only

Gilad et al. 2004

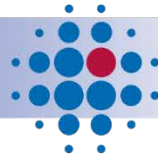
- pooling
  - KHVD max. 2 fish
  - health monitoring max. 5 fish



# Pathogenesis and sample collection

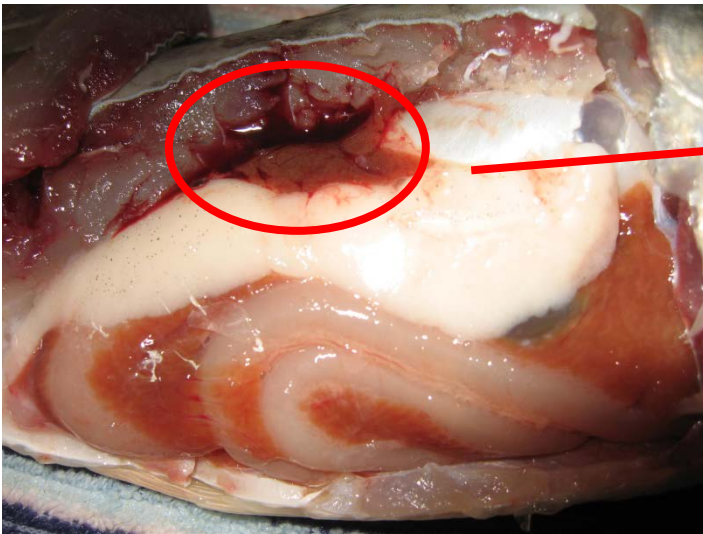
## Early pathogenesis: 0 – 3 dpi

- to 1<sup>st</sup> dpi:**
- KHV enters through gill and gut tissues but not via skin (no mRNA)
  - after 2 hpi in gill and gut cells (weak mRNA)
  - after 4 hpi in leucocytes (weak mRNA – first capsid assembly)
  - after 6 hpi in all inner organs (weak mRNA)
- from 1<sup>st</sup> dpi** – explosive replication in kidney and liver
- spread by leucocytes
  - released via gut
- from 3<sup>rd</sup> dpi**
- occurrence in skin, gut and gills again in leucocytes but also epithelial cells (gills, perhaps gut)
  - virus release via gut and gill



# carp kidney samples and KHV

## ISH carp kidney



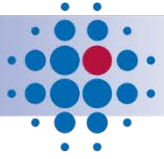
25 mg

PCR -

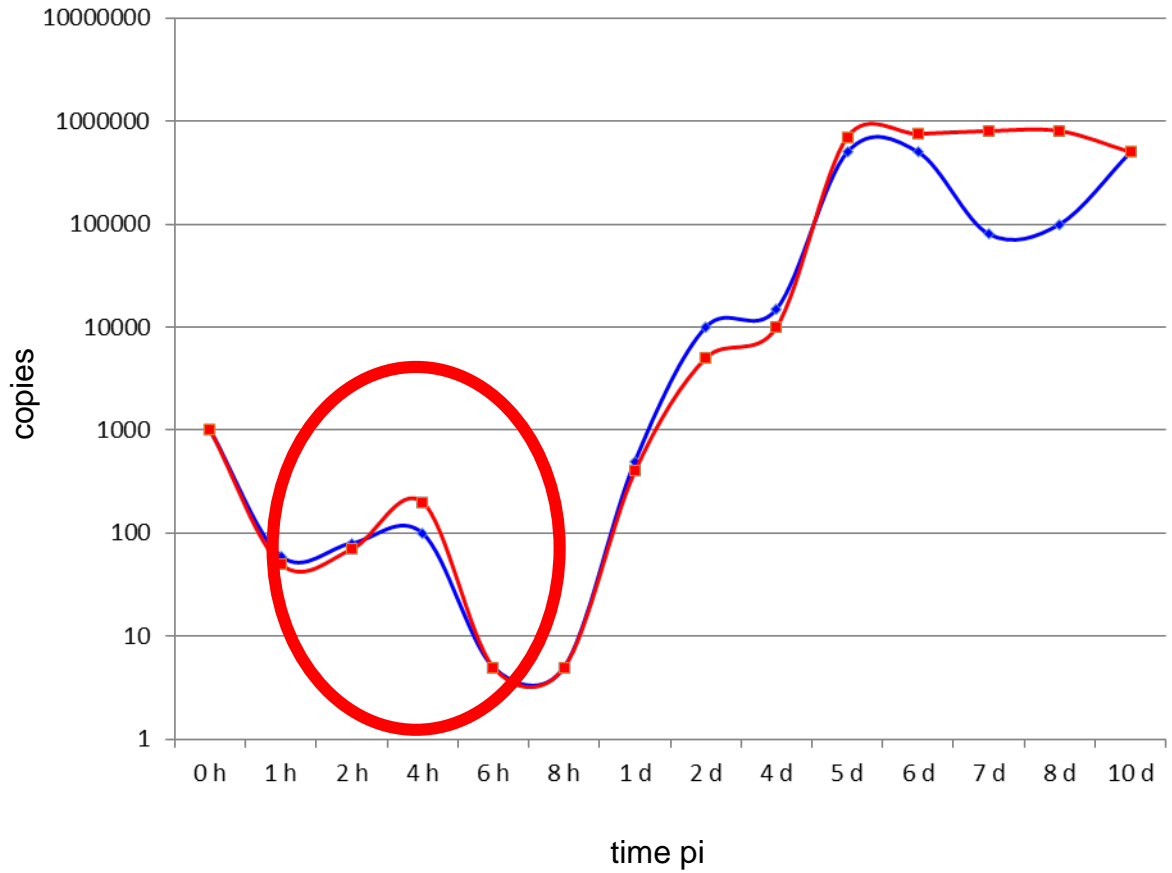
PCR +

low sensitivity of virus detection methods due to low concentrations

**+ sensitivity of the assay**  
(enzyme etc.)



### „outside“ tissues (peracute KHVD)



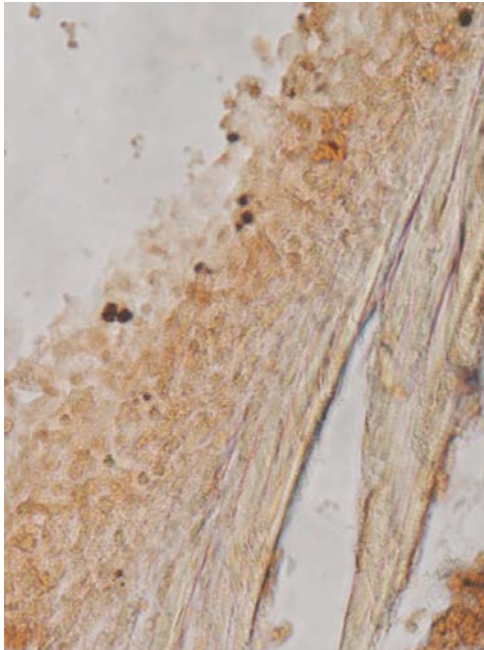
mRNA  
gill  
skin  
no mRNA



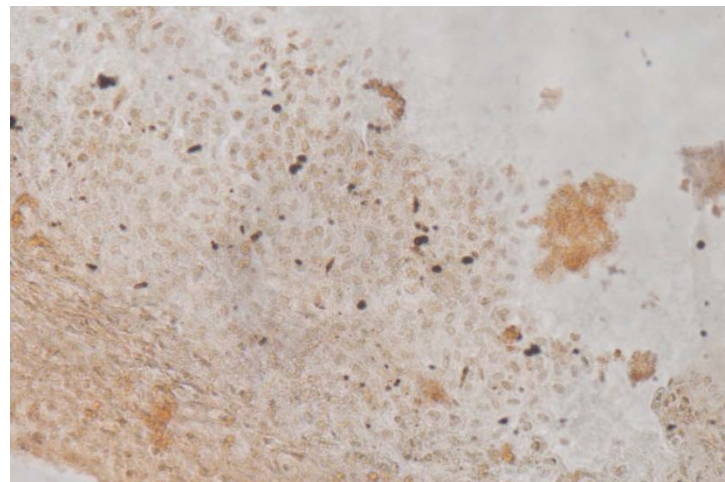
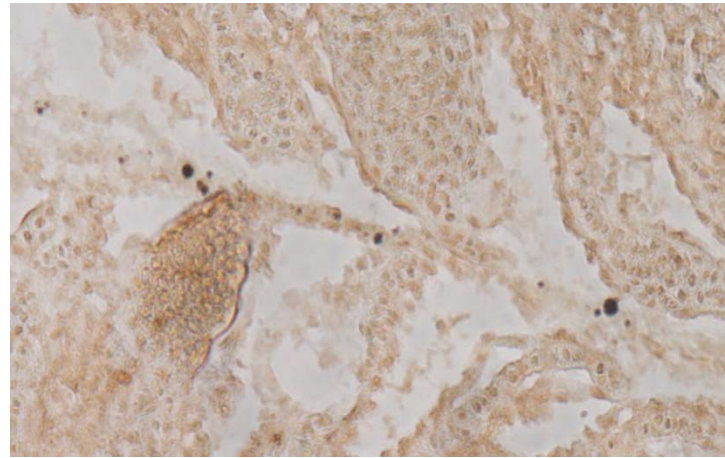
## tissues 1 - 2 h pi

DNA bearing cells in the mucus

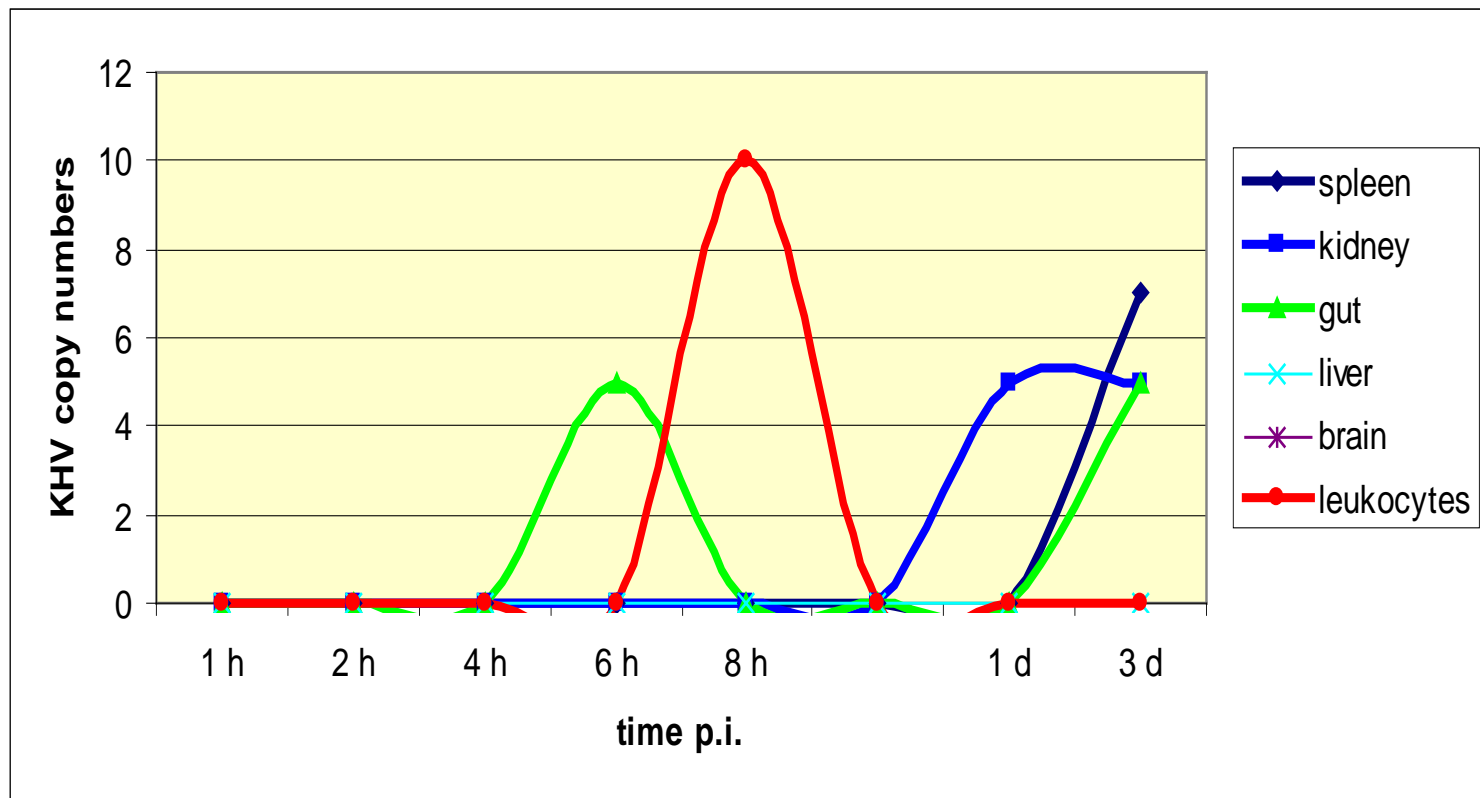
gill

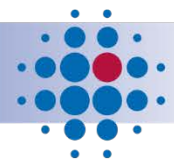


attached from outside

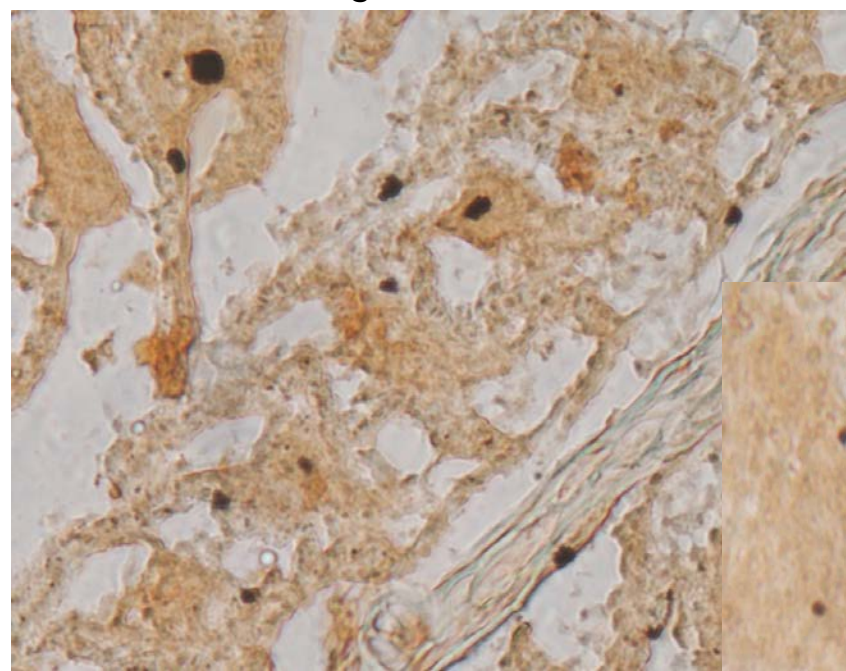


skin  
in the mucus only

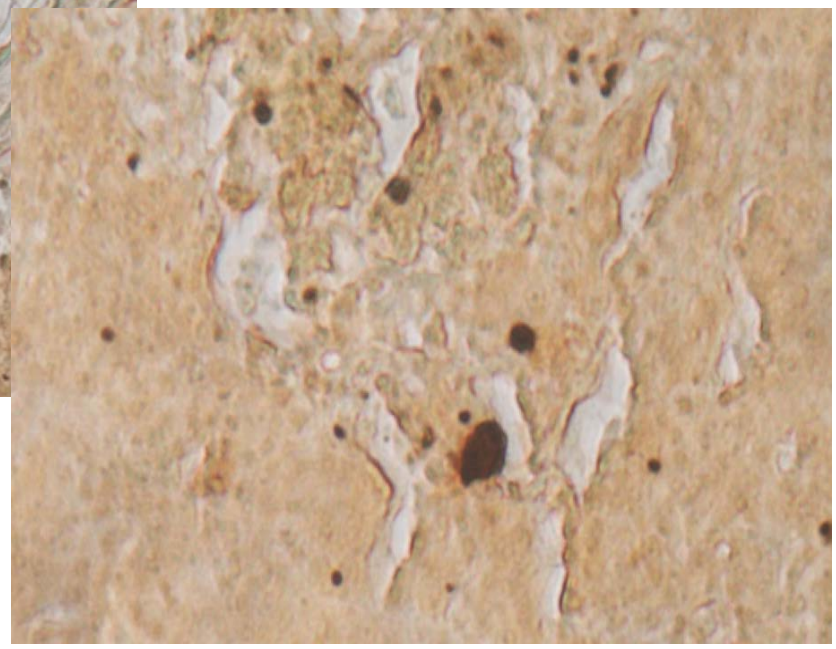
**„inside“ tissues (peracute KHVD)**

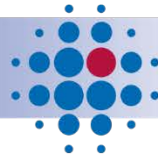


gill

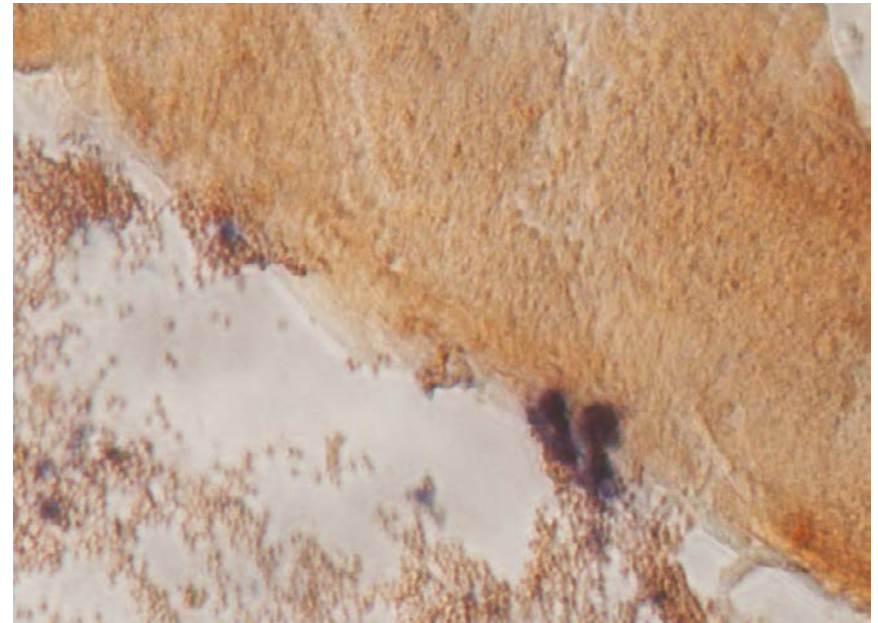
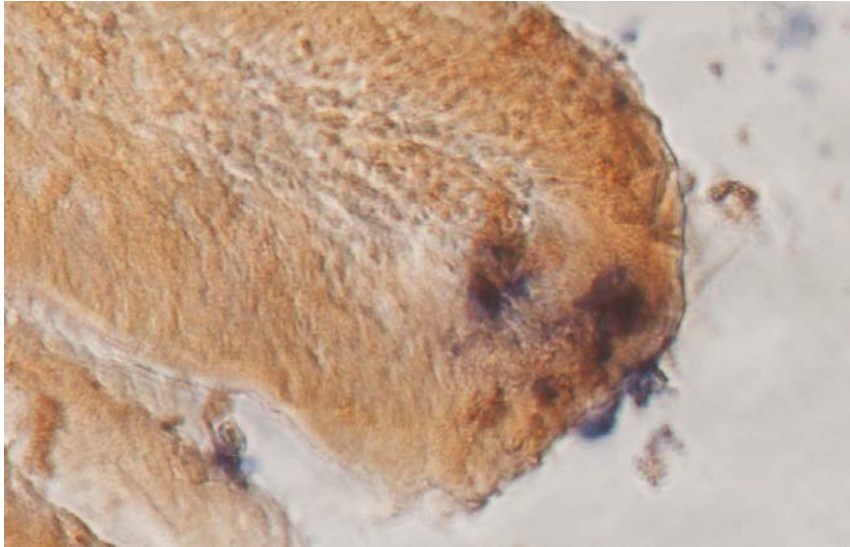


spleen, blood vessels

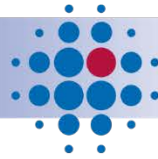




## gut tissue 4 h pi

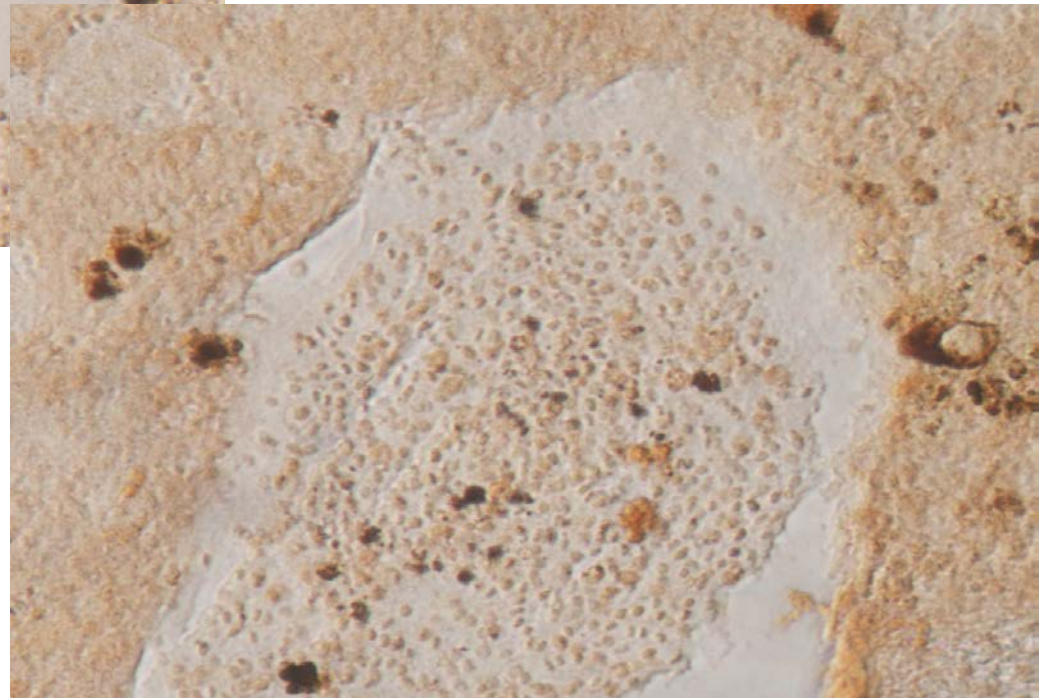
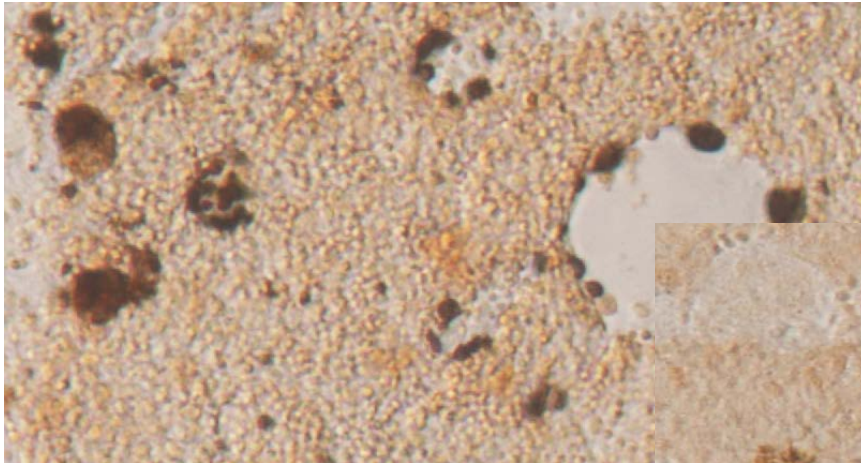


**KHV released at the first time!**



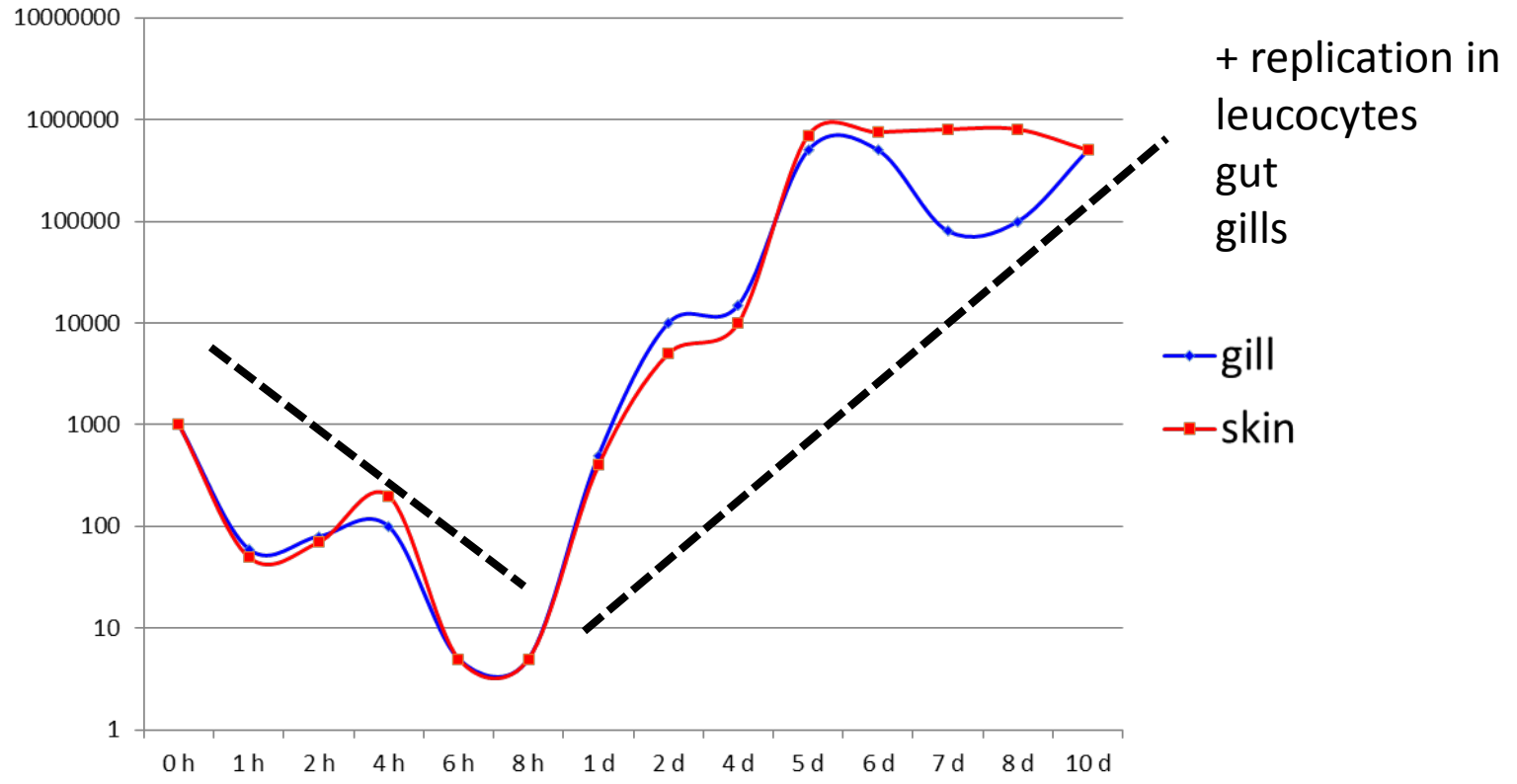
## tissues 4 - 8 h pi

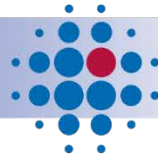
kidney (vessel associated)



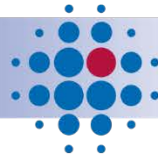


# „outside“ tissues



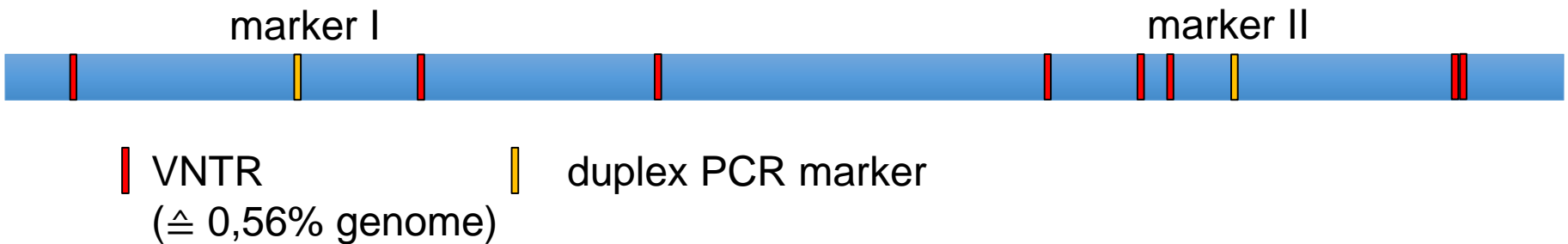


# KHV divertisity

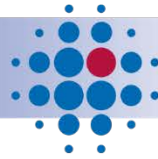


# KHV diversity / variation

- necessary to detect different types of KHV for vaccine development
- molecular tracing for KHV all over the world

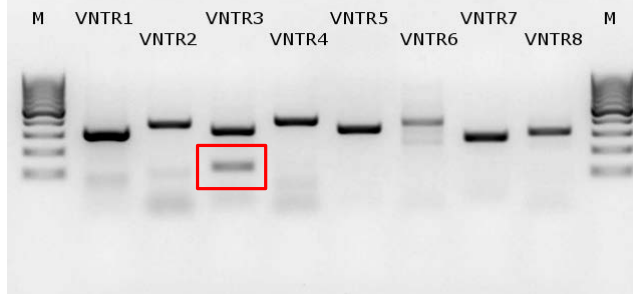




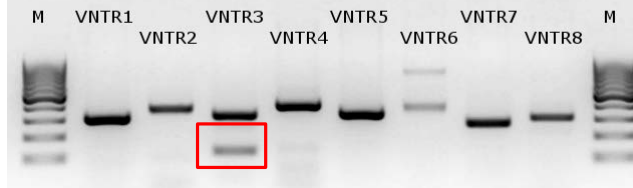


# VNTR PCRs in praxi

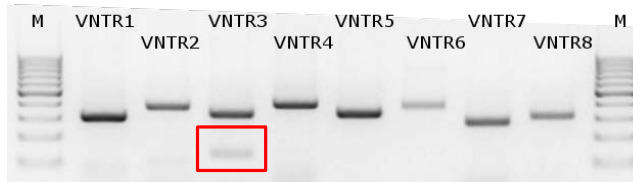
KHV-T 100 Passagen; Passage 25; neue VNTR Primer; 15x60°C 15x55°C



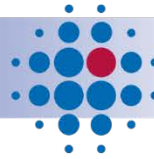
KHV-T 100 Passagen; Passage 51; neue VNTR Primer; 15x60°C 15x55°C



KHV-T 100 Passagen; Passage 78; neue VNTR Primer; 15x60°C 15x55°C



1. new primers
2. diff. annealing temp.
3. differences to CyHV-1 /2 in VNTR3

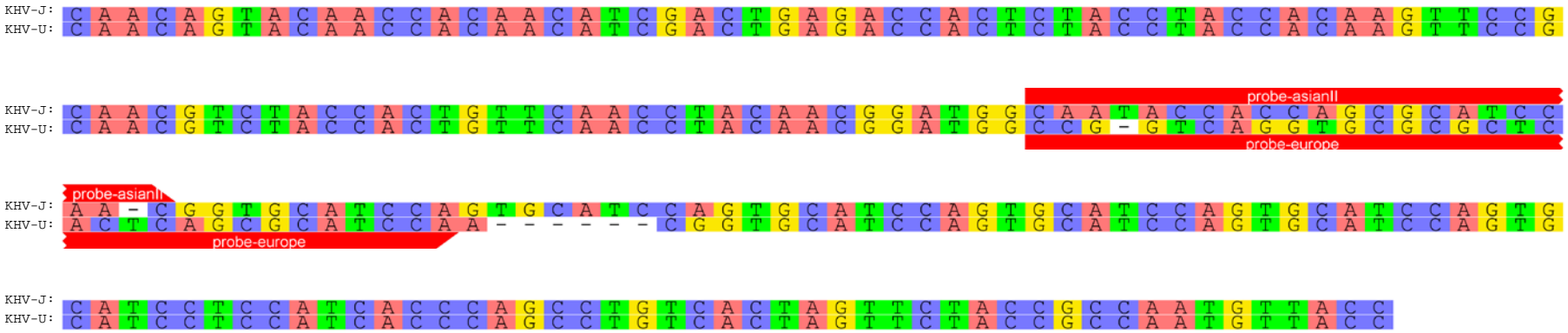


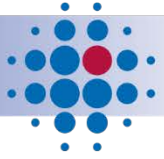
# MLVA applied to KHV passages (100) onto CCB at 20°C



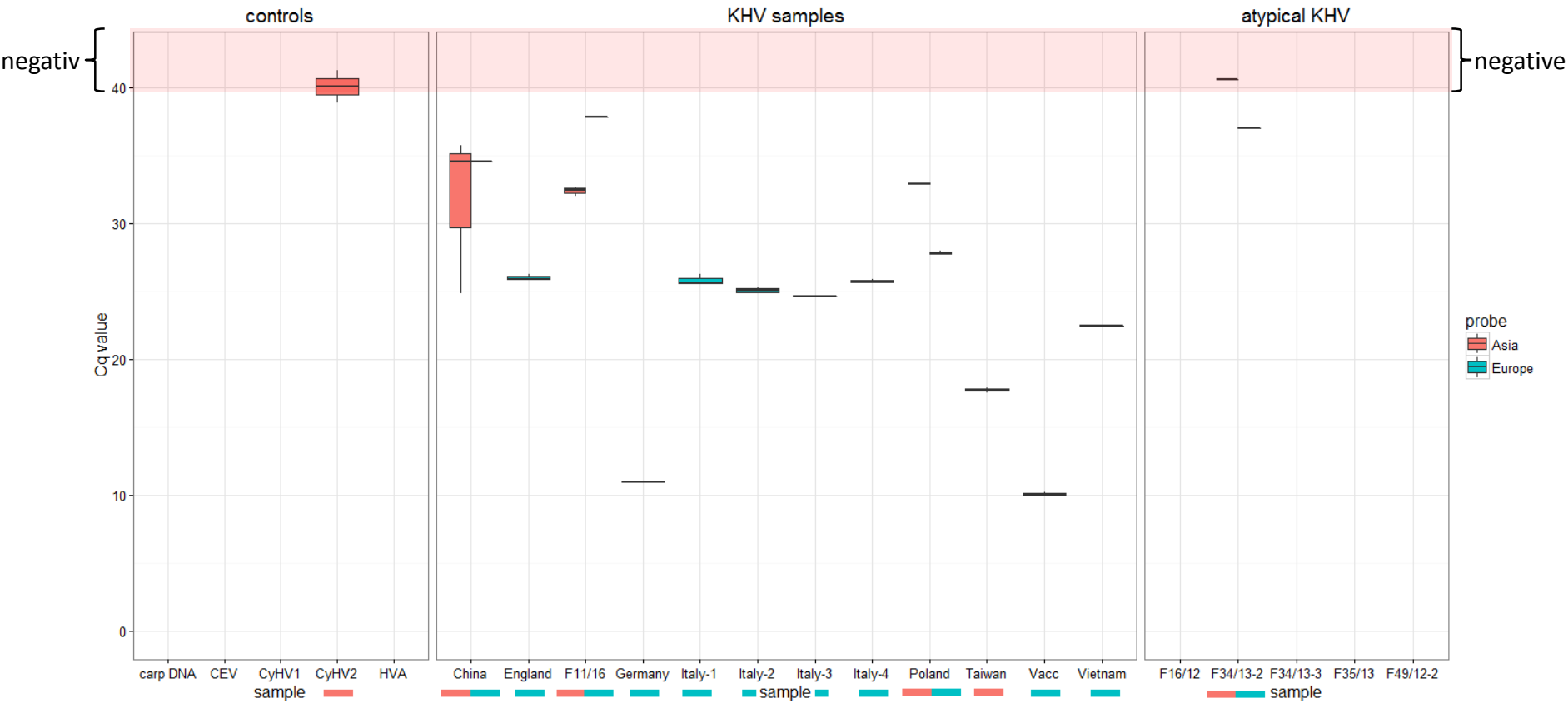


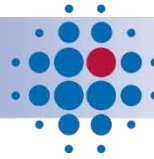
# Development on a new TaqMan qPCR based on VNTR3



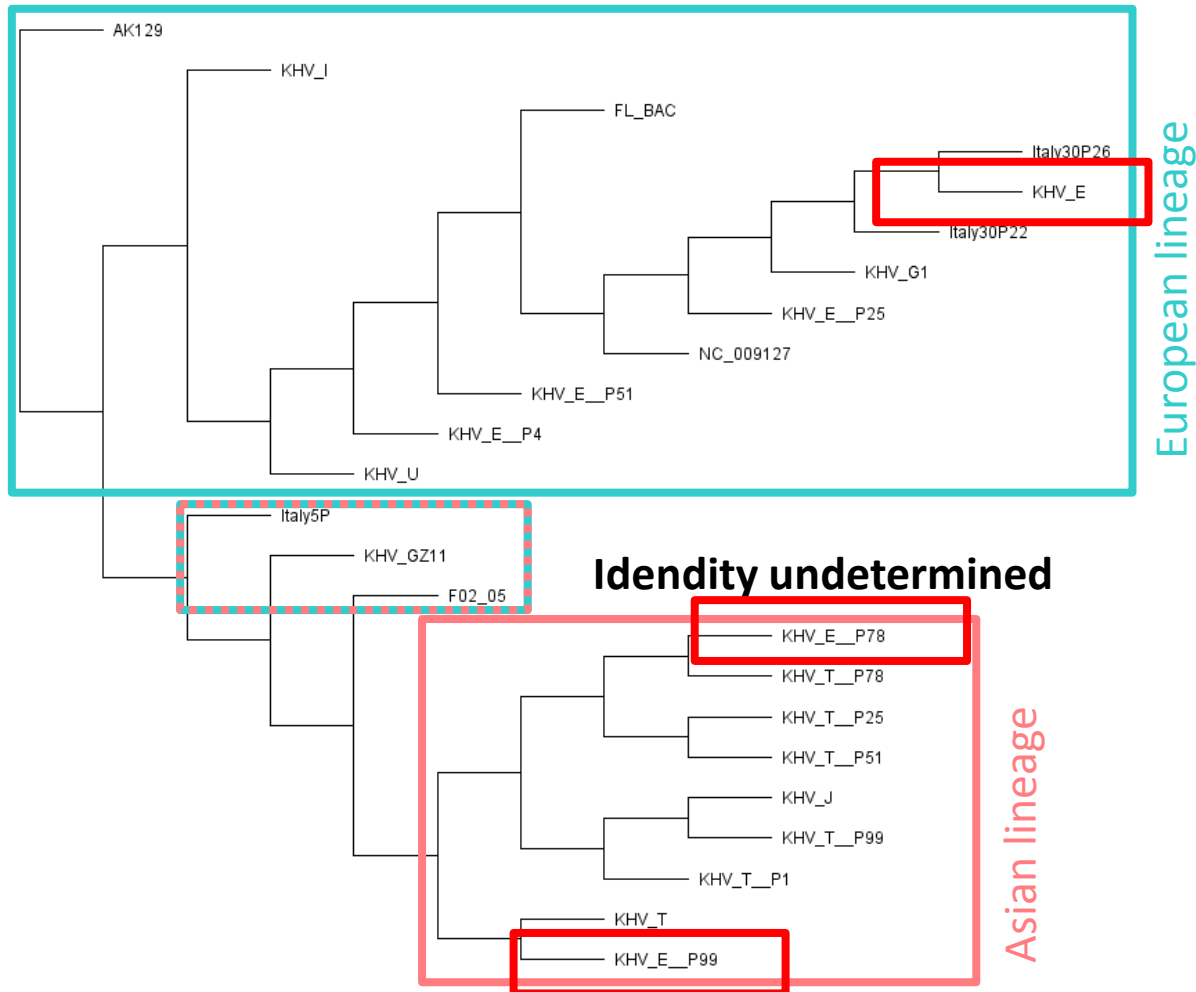


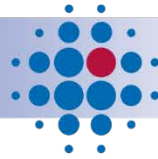
# Some results with the new VNTR3 qPCRs





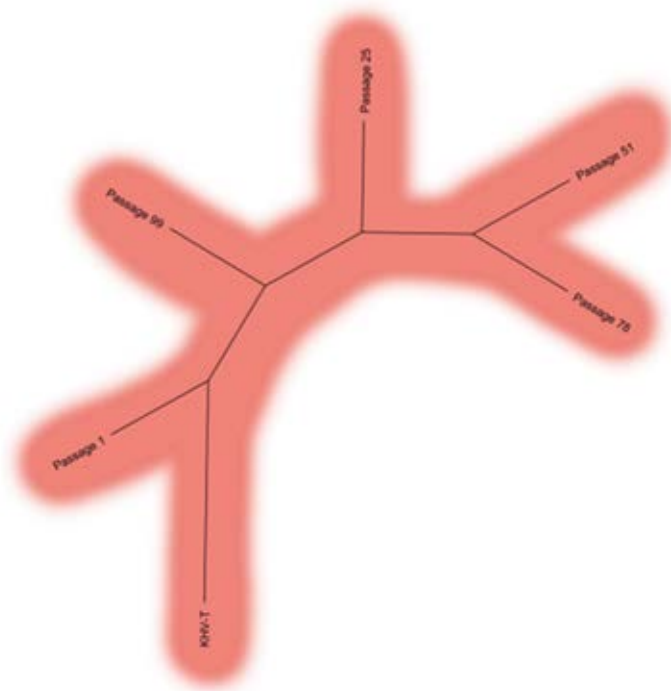
# Sequence analysis using 8 VNTRs





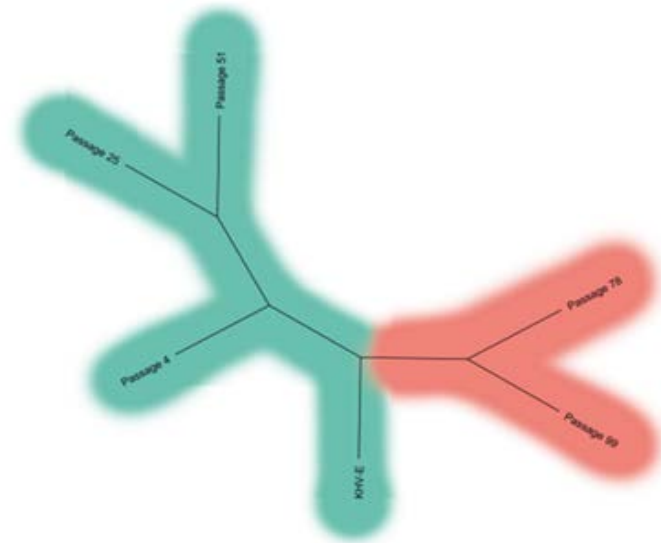
# Variable numbers of tandem repeats qPCR (VNTR-qPCR) with sequence analysis of 8 VNTRs (Klafack et al. 2017)

KHV-T

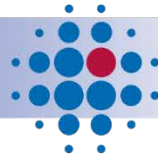


Asian lineage

KHV-E

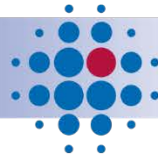


European lineage - Asian lineage



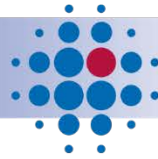
# Recommendation for the diagnostics

1. Use of at least **two molecular assays** (diagnostics + monitoring)
  - qPCR + VNTR3 qPCR
  - qPCR + TK (+ nested + sequence analysis)
  - qPCR + PAN (+ nested + sequence analysis)
  
2. **Sequence analysis** of the fragments
  
3. **Sample collection** (from 5 dpi)
  - Organs - gill swabs or gill biopsate (non-lethal)
  - leucocytes (non-lethal)
  - gill and kidney (lethal)
  
4. **Pooling**
  - early infection – single samples (gills)
  - from day 5 pi – two fish (gills)
  - latent infection – stress induction (gills)  
(24 – 48 h) (30 -60% pos.)



## 2. Treatment against KHVD



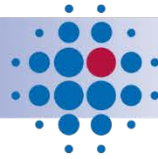


## 2.1. Hygienic measuremets

- „clean“ parents generation (cat. I)
- always clean hatchery
- disinfection (biosecurity)

## 2.2. Immunoprophylaxis (immunization)

- 2.2.1. inactivated vaccine from (a high virulent) KHV
- 2.2.2. attenuated live vaccine
- 2.2.3. genteically engeneered vaccine
- 2.2.4. immunostimulation with virus eradication (alternative)



## 2.2.1. Inactivated vaccine (heat, BEI)

KHV-I ( $10^4$  TCID<sub>50</sub>/ml)

KHV-T ( $10^6$  TCID<sub>50</sub>/ml)



CCB cells

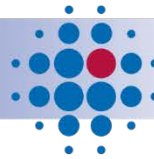
20°C

26°C

field test  
10.000 carp ip

67.8 % PSR

78.8 % PSR



## 2.2.2. attenuated live vaccine



### Germany FLI

#### AK 129 (KHV-I)

#### Att Vacc G (KHV-T)

- passages onto CCB cells
- 129 – 140 x at 28°C

- passages onto CCB cells
- 0 – 100 x at 20°C

-  $10^{2.85}$  TCID<sub>50</sub> / 10 µl

-  $10^{4-6}$  TCID<sub>50</sub> / ml

**results:** survival rate pch  
(before 92% survival after ip)

**results:** survival rate pch  
(P78, P 51, P 99)

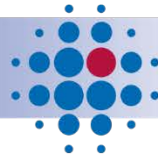
ip	=	100 %
immersion	=	65 %
orally	=	43 %

P 78=	100%	80%
P 51=	10%	-
P 99=	20%	-

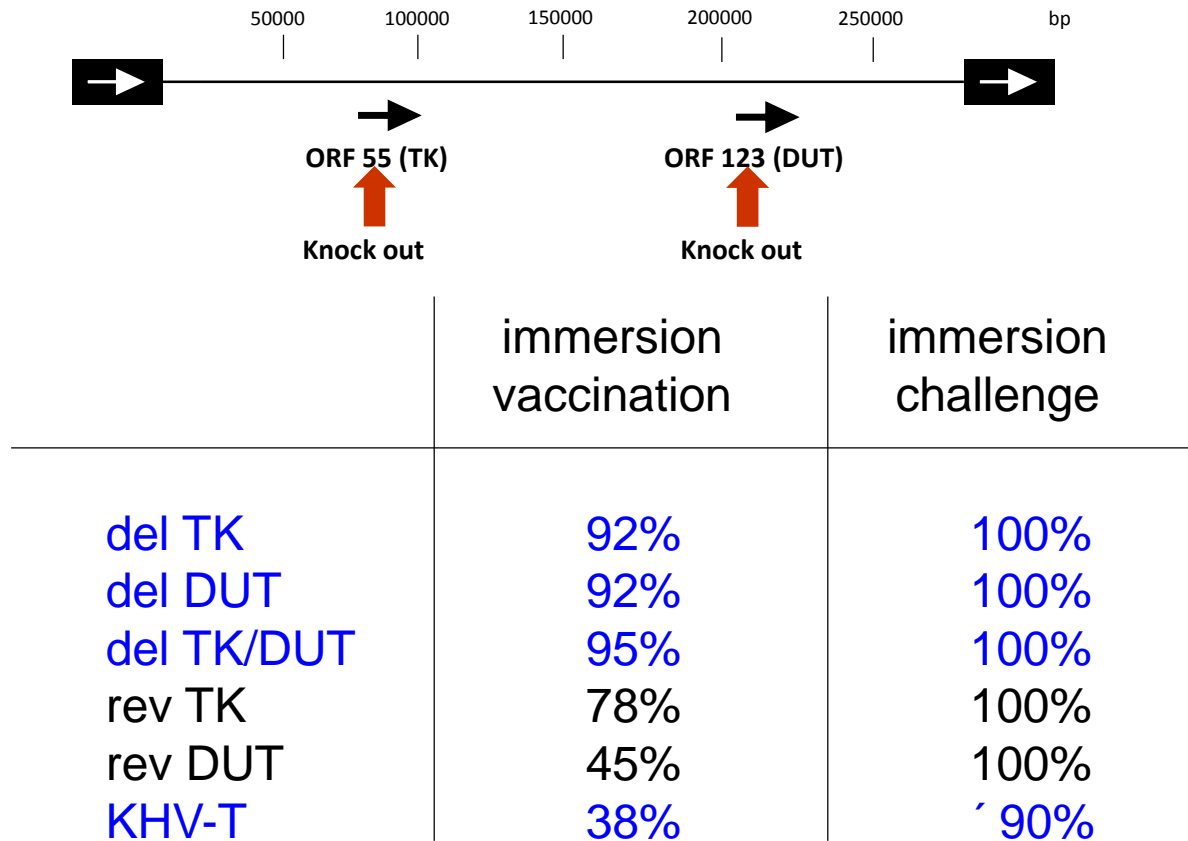
KV 3 (Israel, KoVax Ltd.)

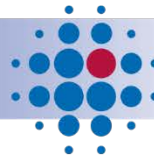
Cavoy (Canada, Novartis)





## 2.2.3. Genetically engineered vaccine (3 variants)



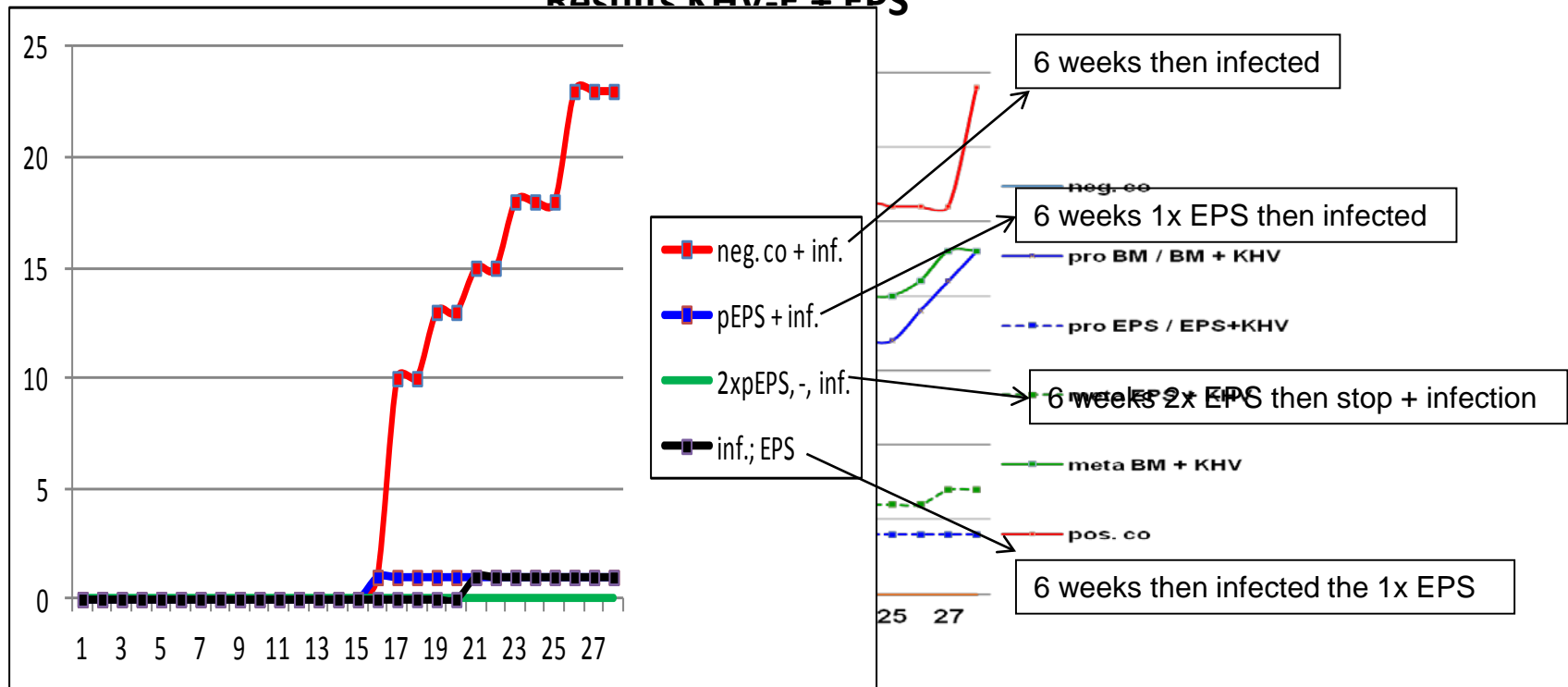


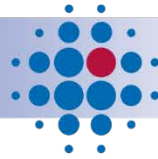
## 2.2.4. immunostimulation with virus eradication

### Exopolysaccharide (EPS) from marine algae

#### Results KHV-T + EPS

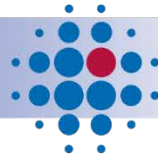
#### Results KHV-E + EPS





## Conclusion for combat

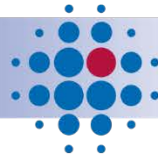
Vaccination and alternatives are possible and can be used very successfully.



### 3. Confirmation of KHV

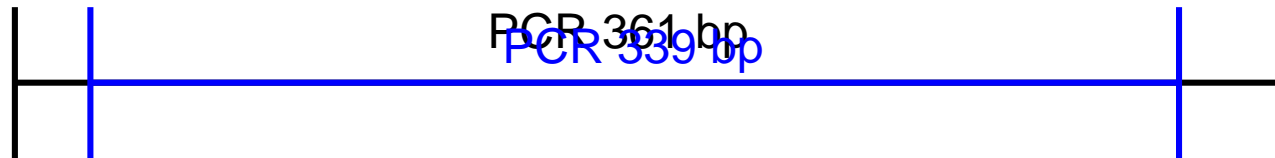


**PCR and nested PCR (Engeslma et a. 2013)**



# PCR according Engelsma et al. 2013

(PAN-CyHV PCR, ORF 79, viral DNA polymerase)



## Recognizing :

CyHV-1 (capo), CyHV-2 (GHV), CyHV-3 (KHV) and AngHV-1 (HVA)

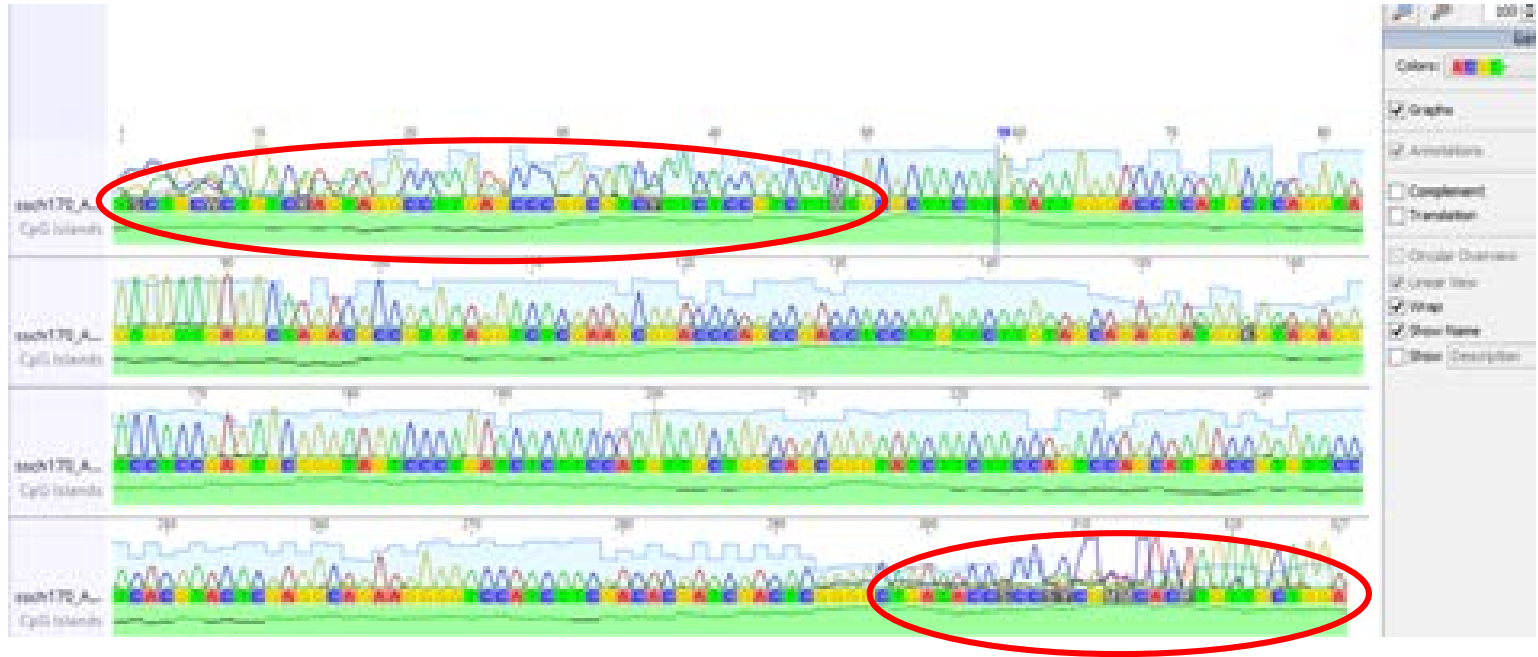
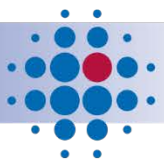
## Sequence analysis of fragments:

1. Which one? PCR or nested PCR or both?
2. Which part of the ORF 79?

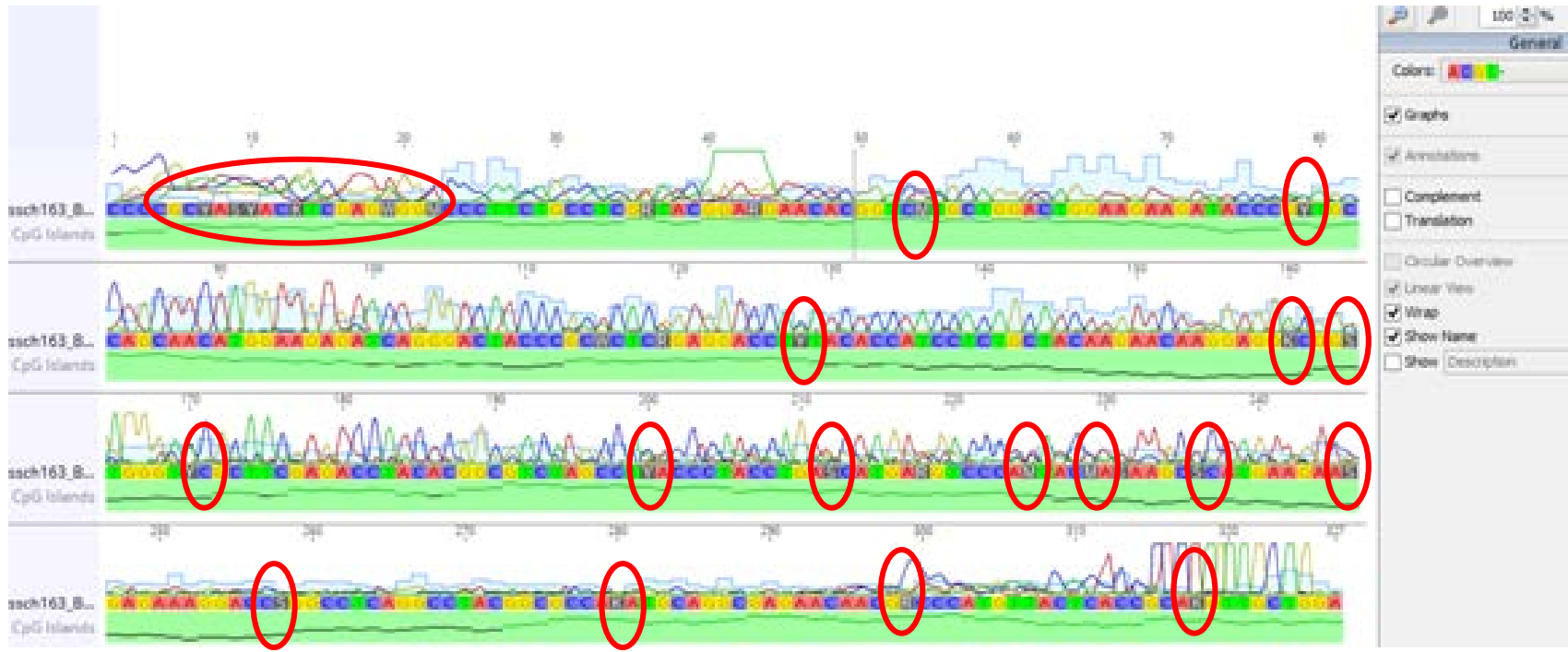
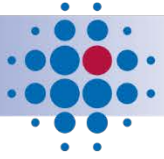
## Recent legislation:

98% similarity of consensus sequences (3) = KHV  
> 98% = no KHV (atypical reacting KHV)

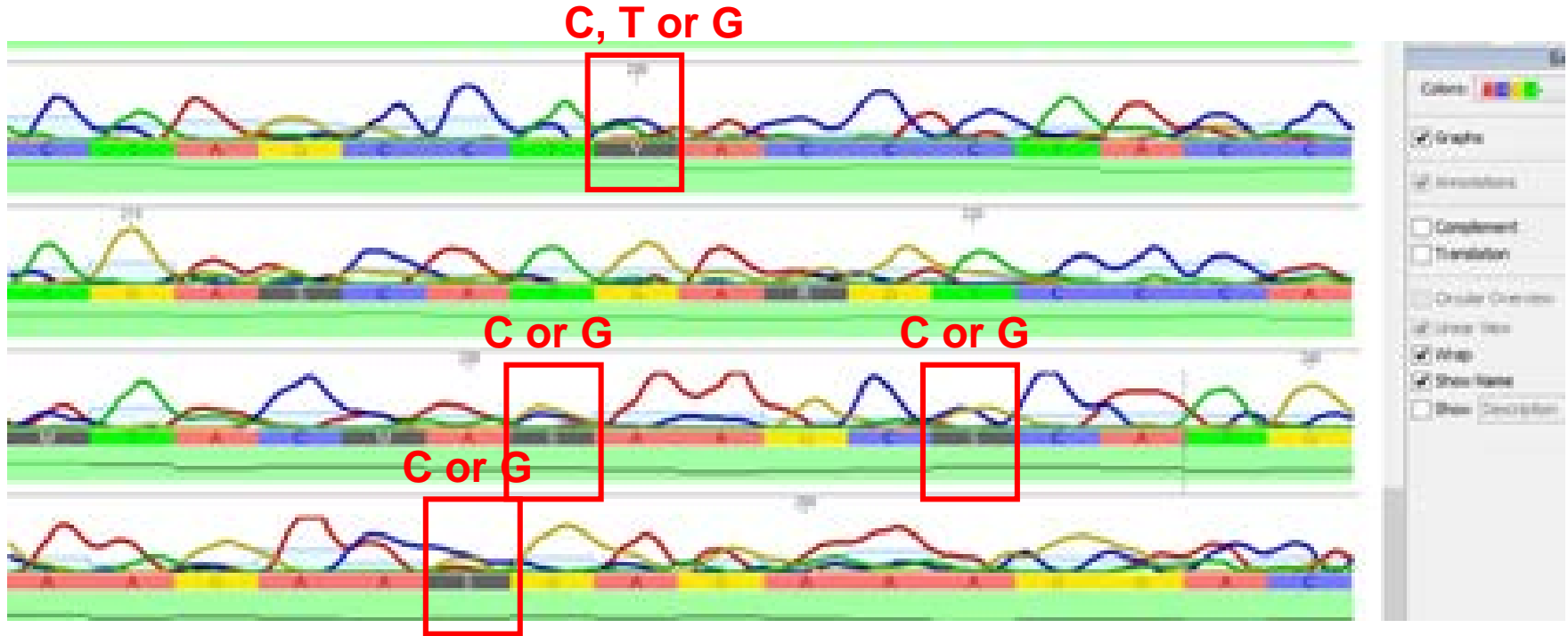
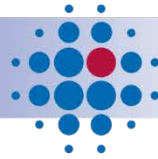




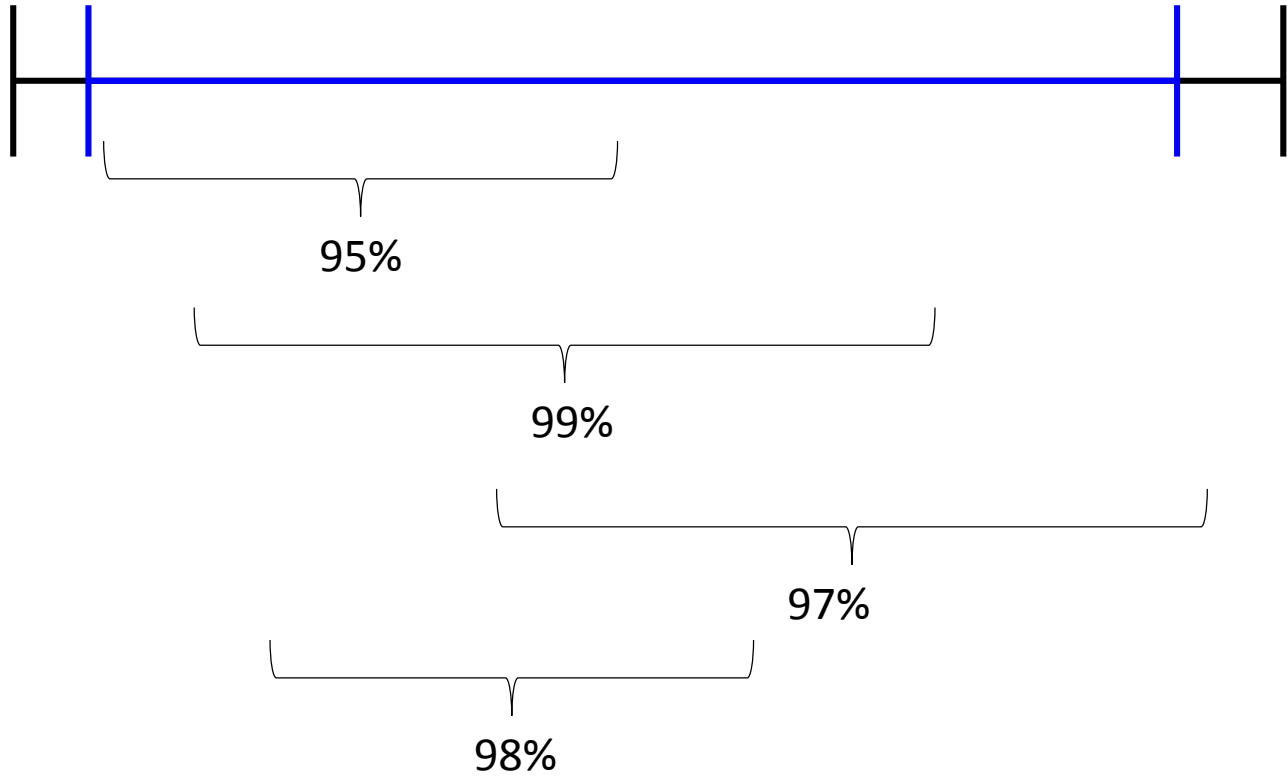
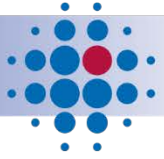
„normal“ nested PCR fragment



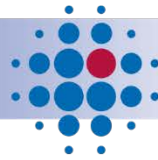
„difficult“ nested PCR fragment



- KHV variants mix
- NGS confirmed by JC Avarre and Bergmann.



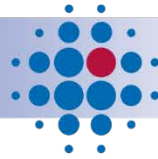
What is right and what is wrong?



# Investigation on atypical reacting and normal reacting KHV (PAN CyHV PCR, % similarity)

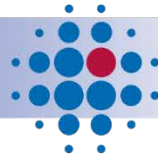
		340-280 bp	150-160 bp	atyp.	Tk/n	ca	GHV	HVA
<b>atyp.</b>	<b>2</b>	<b>96</b>	<b>100</b>	<b>97</b>	<b>-/+</b>	<b>86</b>	<b>81</b>	<b>79</b>
	<b>4</b>	<b>95</b>	<b>99</b>	<b>96</b>	<b>-/+</b>	<b>86</b>	<b>80</b>	<b>78</b>
	<b>5*</b>	<b>96</b>	<b>99</b>	<b>97</b>	<b>-/-</b>	<b>85</b>	<b>78</b>	<b>72</b>
	<b>7*</b>	<b>97,7</b>	<b>99</b>	<b>96</b>	<b>-/-</b>	<b>82</b>	<b>77</b>	<b>71</b>
<b>typ.</b>	<b>8</b>	<b>98</b>	<b>100</b>	<b>98</b>	<b>+/+</b>	<b>84</b>	<b>78</b>	<b>74</b>
	<b>10</b>	<b>100</b>	<b>100</b>	<b>97</b>	<b>+/+</b>	<b>83</b>	<b>78</b>	<b>73</b>

\* isolates onto CCB cells but from the organs



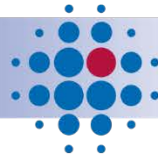
# Investigation KHV isolates at different temperatures (PAN CyHV PCR, % similarity from 10<sup>th</sup> passage)

	15°C	20°C	25°C
KHV-T	100	100	100
KHV-E	96	100	97



## Conclusion for the confirmative PCR

1. value 98% similarities is very, very artificial
2. if necessary 95% would fit to KHV variants (with mortality!)  
95 -100% = KHV  
>90 % = different herpesviruses
3. other and /or confirmative qPCRs? (qPCR VNTR3 or lin. qPCR?)  
(specific to KHV not to other aquatic herpesviruses)



**Thank you very, very much!**



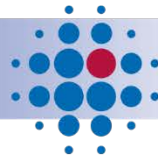
Lars Schröder



Working group Dr. Walter Fuchs

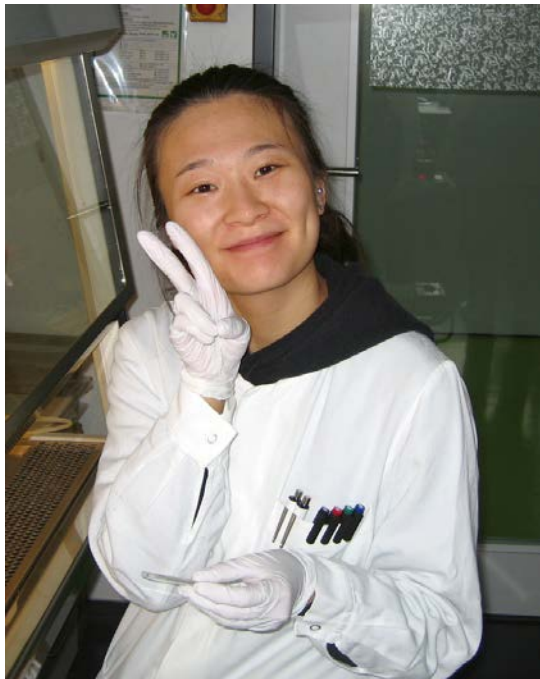
Prof. Dr. Dr. h.c. Thomas C. Mettenleiter





**JC Avarre**

**Thank you very, very much!**



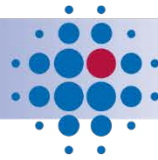
**Yeonhwa Jin**



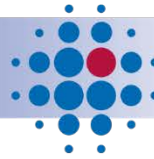
**Sandro Klafack**

**Kyo Hyun Kim**

**Irina Werner**



**Prof. Wang Qing**  
**Dr. Wang Yingying**  
**Dr. Zeng Weiwei**  
**Li Yingying**  
**Prof. Kielpinska**  
**Natalia**



**Thank you for  
your attention!**