

Current AI-related activities at Riems

FRIEDRICH-LOEFFLER-INSTITUT

FLI

Bundesforschungsinstitut für Tiergesundheit
Federal Research Institute for Animal Health

O.I.E. and National Reference Laboratory
for Avian Influenza

Amplification**Transcription****Extraction****Armored RNA**
**(modified recombinant
MS2 phage particles)**

Armored RNA for avian influenza virus

Universal IC

Eurasian control

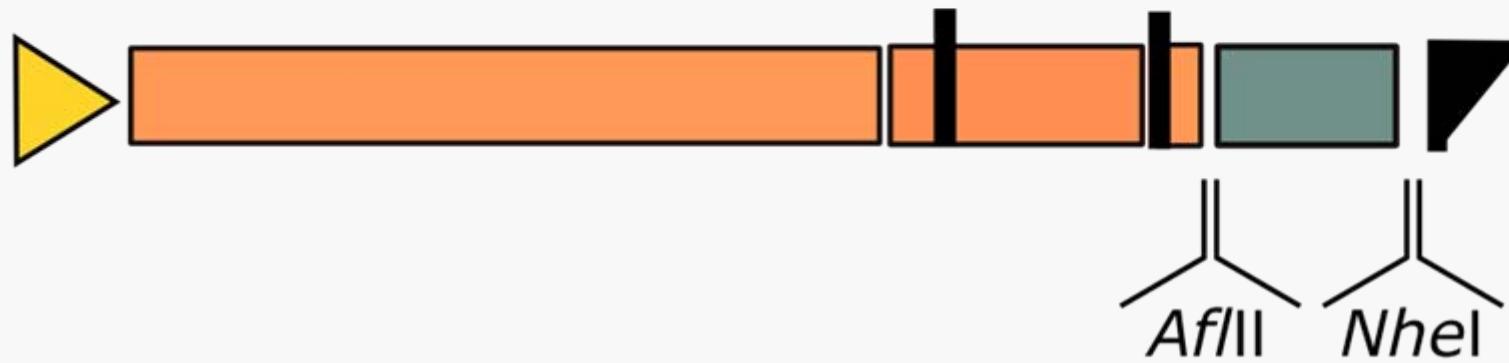
American control



IC

EGFP gene fragment

pT7 MS2 (*mat-coat-repf*) 6H pac IC T7t



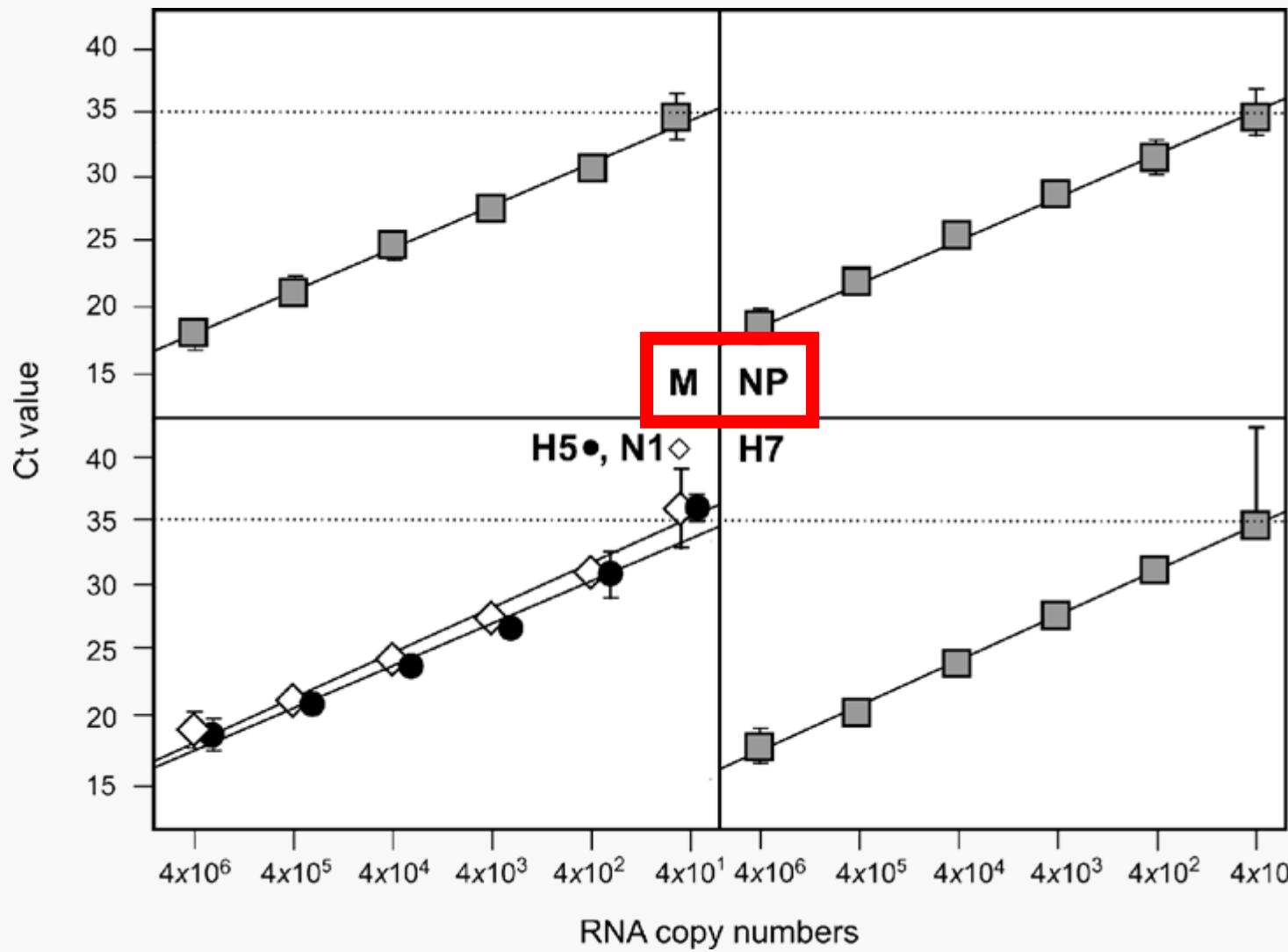
Eurasian control

EU**New**

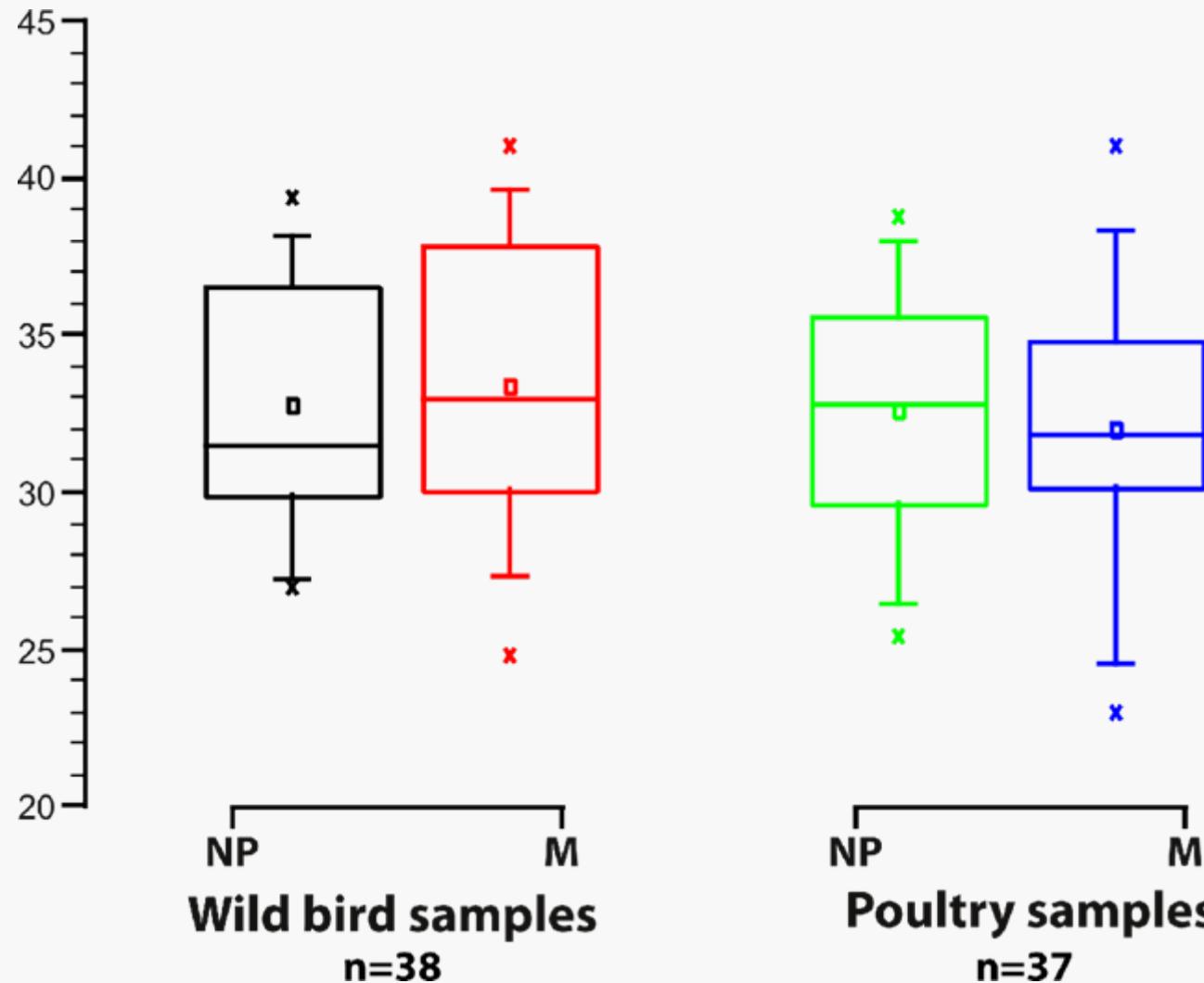
pT7 MS2 (*mat-coat-repf*) 6H pac H5 H7 M NP N1 T7t



Eurasian control – run-off RNA



Features of the new „universal“ NP rRT-PCR



Features of the new „universal“ NP rRT-PCR

	Kappa	Range	Weighted	Criterion
WB	0,91	0,818-1,009	0,89	very good
P	0,87	0,777-0,972	0,87	very good
All	0,91	0,84-0,969	0,89	very good

WB=68

		M		Inhib
		+	-	
NP	+	38	0	1
	-	0	27	1
		0	1	0

P=98

		M		Inhib
		+	-	
NP	+	37	3	0
	-	2	55	1
		0	0	0

All=166

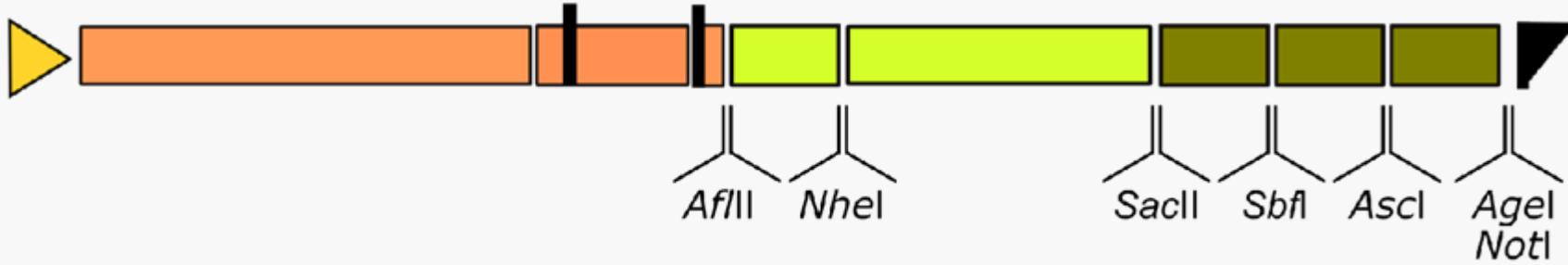
		M		Inhib
		+	-	
NP	+	75	3	1
	-	2	82	1
		0	1	0



American control

US

pT7 MS2 (*mat-coat-repf*) 6H pac H5 H7-1/H7-2 M NP N1 T7t



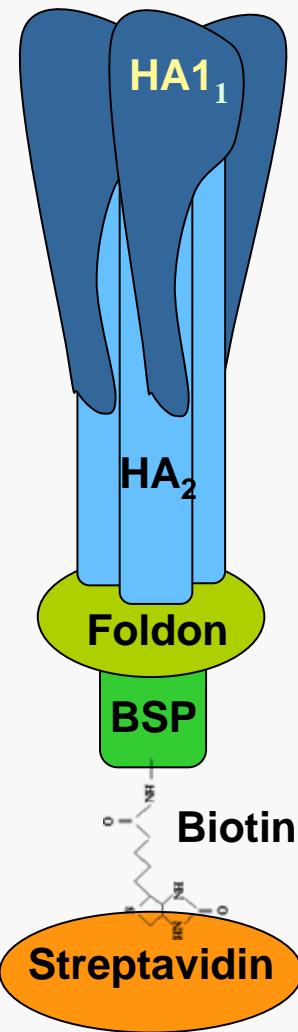
“Production” of recombinant MS2 phages failed so far

In vitro transcribed RNA ready for distribution to OFFLU reference labs (ring trial)

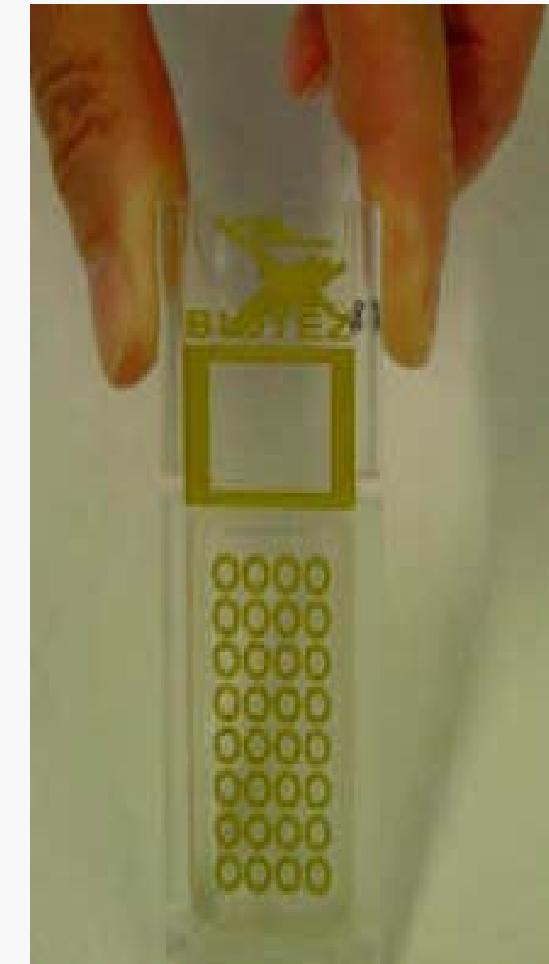
Open for continuing curation and adaptation to changing primer/probe/target recommendations



On-site, subtype-specific serology



- Recombinant, baculovirus-expressed, *in vivo* biotinylated antigen (H5, H7, H9, NP)
- Macrochip reaction system
- Competition ELISA format
- DAS ELISA format



Reverse genetics group (Juergen Stech)

Recombinant vaccines group (Jutta Veits)

Pathogenicity group (Martin Beer)

**Surveillance/epidemiology/data base group
(Christoph Staubach, Timm Harder)**

**Genome sequencing group (Elke Starick,
Dirk Hooper)**



**Thanks,
and best regards from the island!**



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