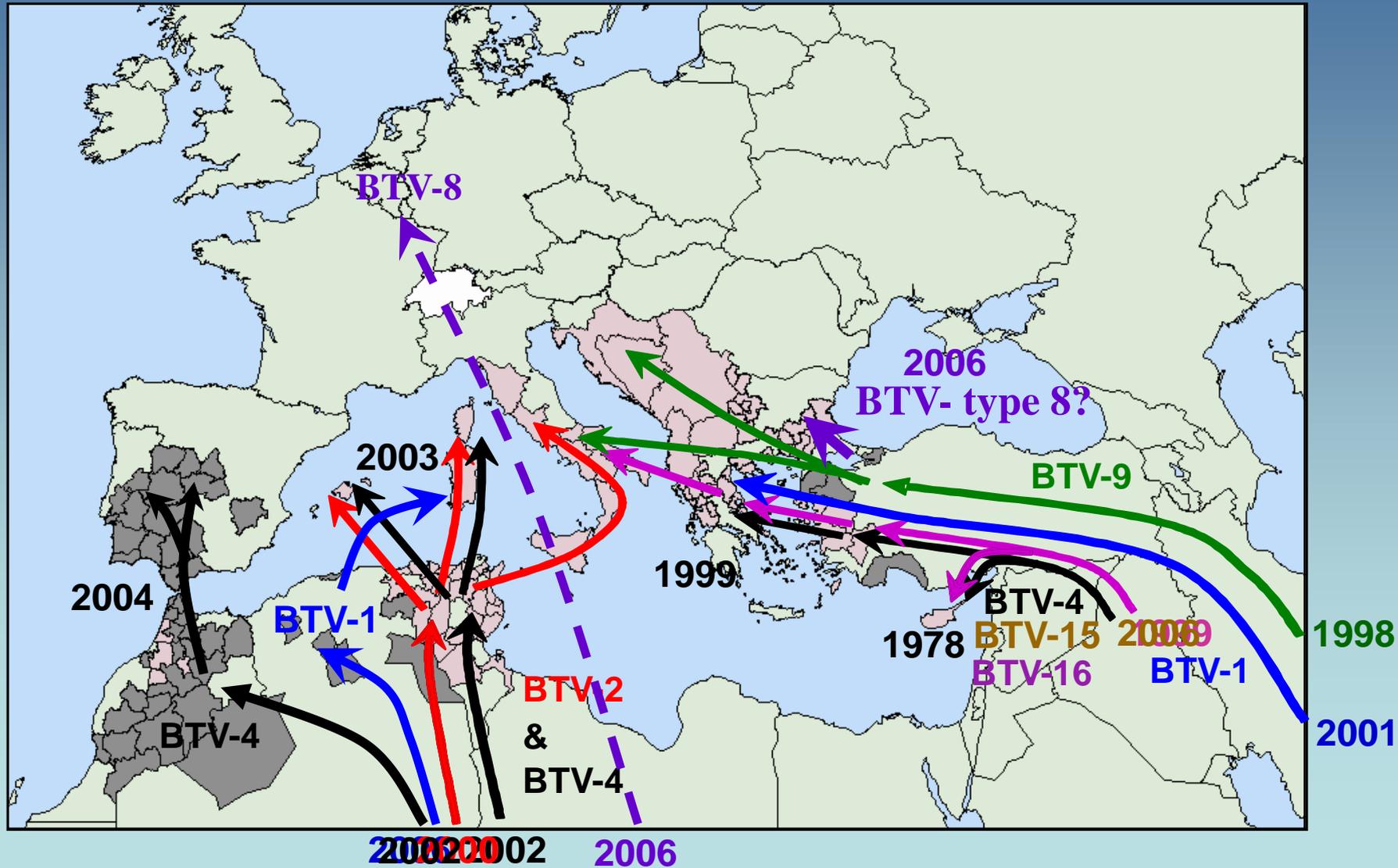


MedReoNet:  
Molecular Epidemiology Work Package

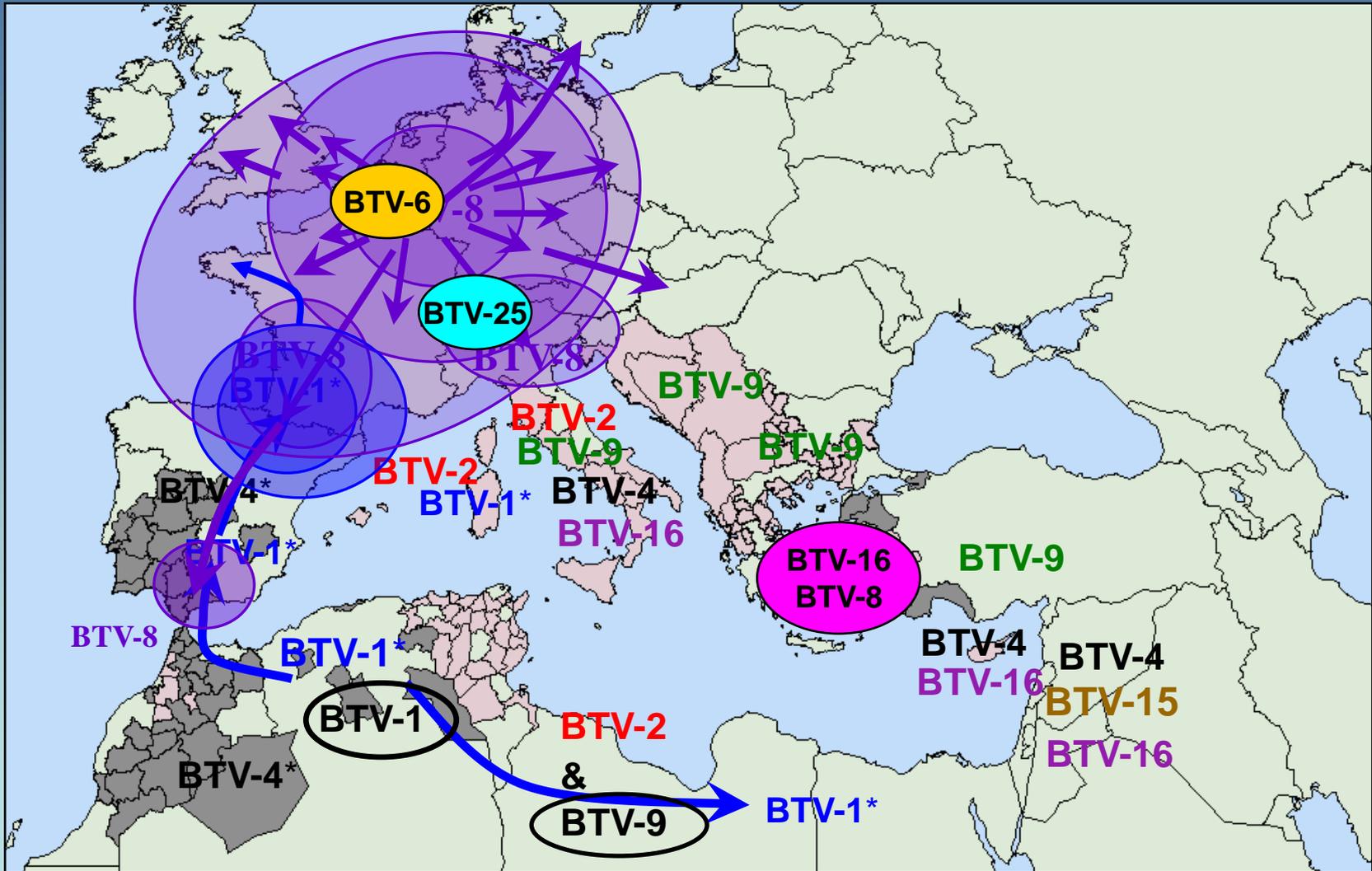
**Update for 2008**

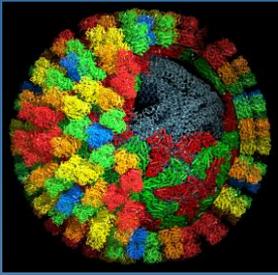
# The 10 introductions of different BTV strains into Europe and north Africa 1998 to end of 2006



# 12 different Introductions of BTV into Europe since 1998

~~May 2008~~





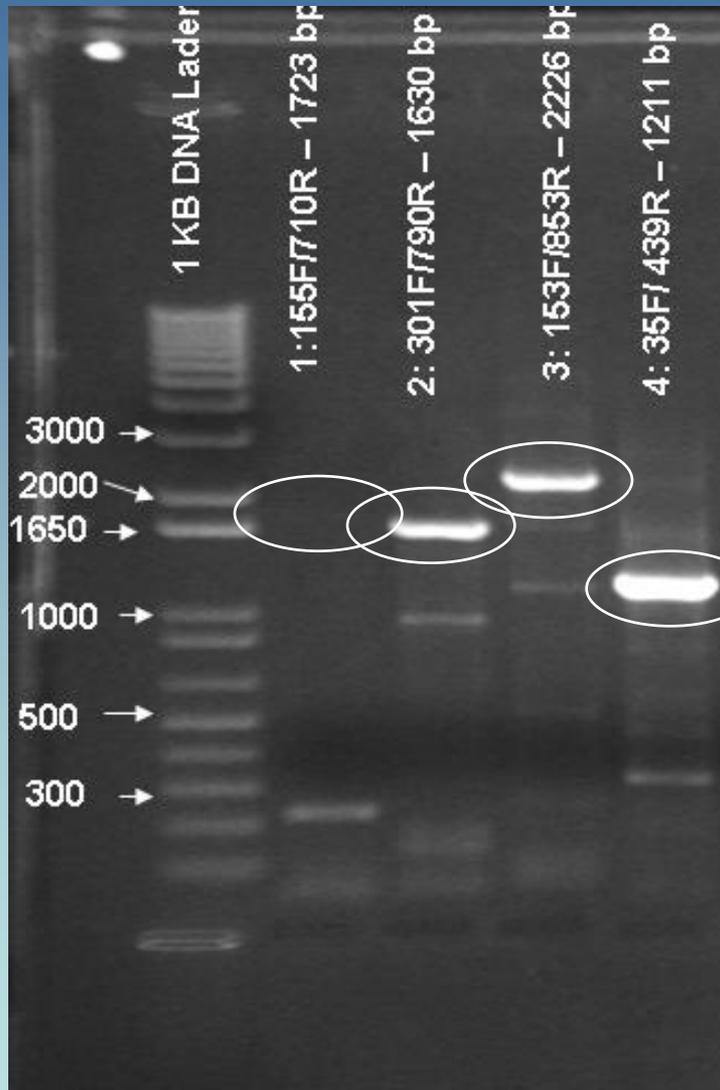
# New BTV strains for 2008?

The Netherlands (October )



# RT-PCR and agarose gel analysis of a KC1 isolate using segment 2 specific primers against BTV- 6

([Sample 2-X08021376 – Heeten- NET2008/04](#),

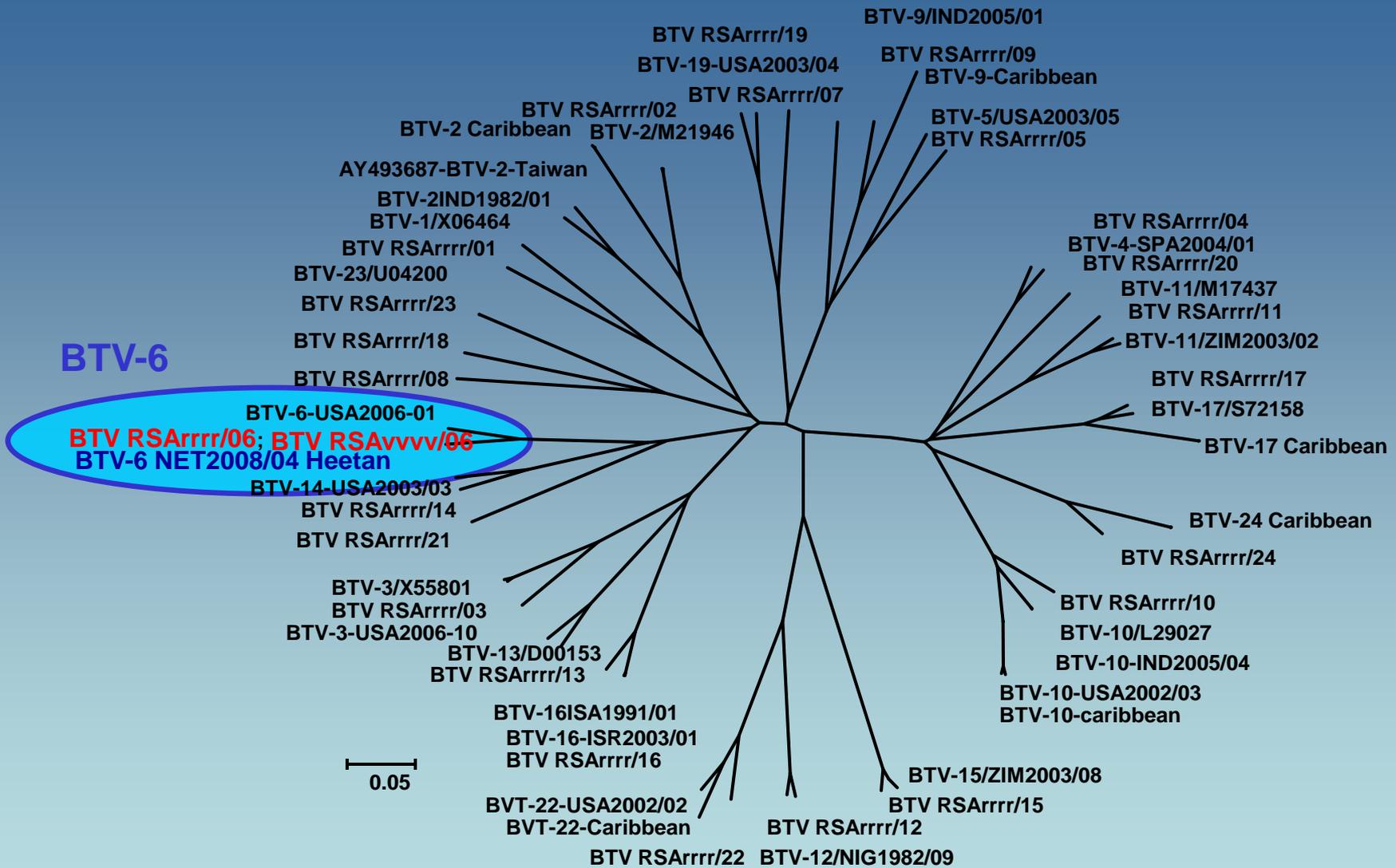


Lane 1: 1723 bp product with type 6 primer pair: BTV-6/2/p155F and BTV-6/2/p710R

Lane 2: 1630 bp product with type 6 primer pair BTV-6/2/p301F and BTV-6/2/p790R

Lane 3: 2226 bp product with type 6 primer pair BTV-6/2/p153F and BTV-6/2/p853R

Lane 4: 1211 bp product with type 6 primer pair BTV-6/2/p35F and BTV-6/2/p439R



**Phylogenetic analysis of full length Seg-2 of BTV-6 Netherlands 2008**

## Similarity between BTV-6 (NET2008/04) and other selected BTV strains, in each genome segment

<b>BTV-6 (Netherlands)</b> Genome segment Number	% similarity with <u>BTV-6 reference</u> strain (RSArrrr/06)	% similarity with <u>BTV-8 Netherlands</u> strain (NET2008)	% similarity with <u>BTV-2 vaccine</u> RSAvvv2/01
1			
2	→ 99.7%		
3			
4			
5	→ 99.9%		
6			
7		→ 99.9%	
8	→ 99.6 %		
9	→ 99.6 %		
10	→ 93.3%		→ 98.4%

\* Data for the other segments of the BTV-6 vaccine are being generated in South Africa :  
BTV-2 was used as a component of the Israeli multivalent vaccine

# **New BTV strains for 2008?**

**Switzerland**

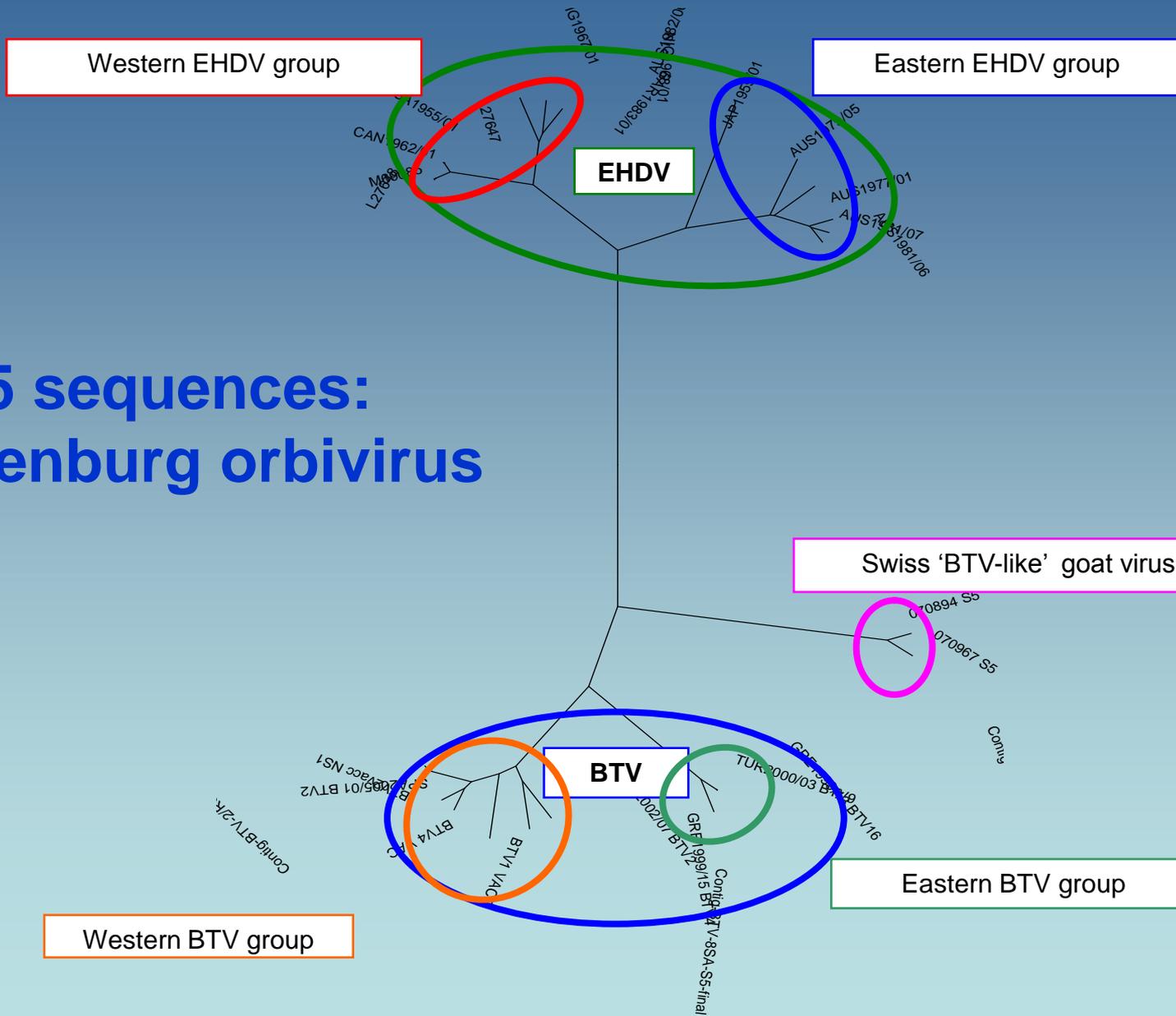
# New BTV strains for 2008?

Toggenburg orbivirus, Cases in goats in Switzerland

No  
clinical  
signs of  
infection



# Seg-5 sequences: Toggenburg orbivirus



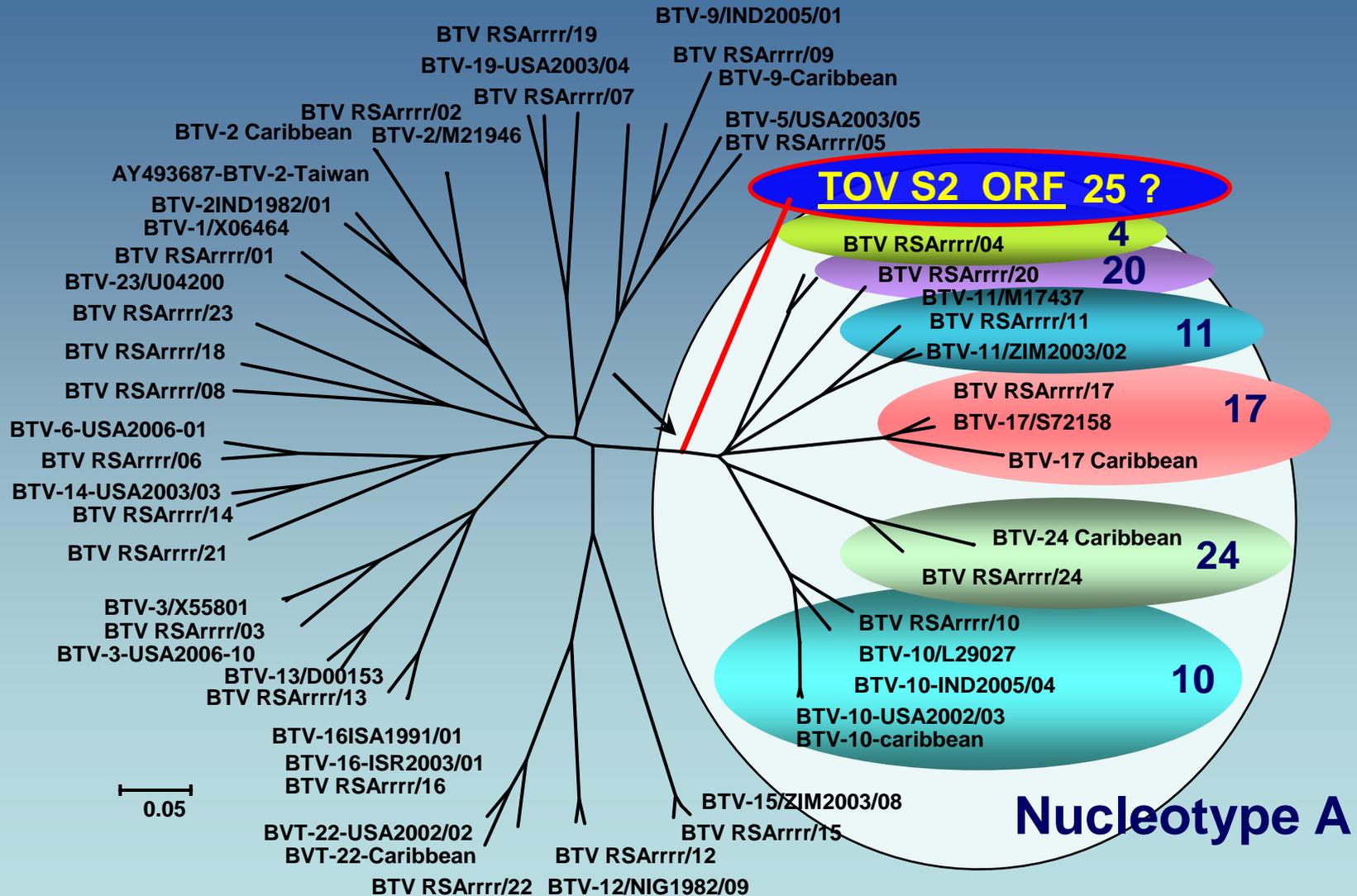
BTV	A119/08 1 CAE0016	A119/08 2 CAE0020	A119/08 3 CAE0034	A119/08 4 CAE0040	A119/08 5 CAE0068	A119/08 6 CAE0089	A119/08 7 CAE0097
1	-	-	-	-	-	-	-
2	-	-	-	-	-	-	-
3	1/10	-	-	1/10	-	-	1/20
4	1/10	-	-	-	-	-	-
5	-	-	-	-	-	-	-
6	-	-	-	-	-	-	-
7	-	-	-	-	-	-	-
8	-	-	-	-	-	-	-
9	-	-	-	-	-	-	-
10	-	-	-	-	-	-	-
11	-	-	-	-	-	-	1/10
12	-	-	-	-	-	-	-
13	-	-	-	-	-	-	-
14	-	-	-	-	-	-	-
15	-	-	-	-	-	-	-
16	-	-	-	-	-	-	-
17	-	-	-	-	-	-	-
18	-	-	-	-	-	-	-
19	-	-	-	-	-	-	-
20	1/30	1/10	-	1/20	-	-	1/15
21	-	-	-	-	-	-	-
22	-	-	-	-	1/10	-	1/10
23	1/10	-	-	1/15	-	-	-
24	1/20	1/10	-	1/10	1/10	-	1/15

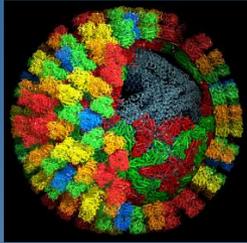
**Serum Neutralisation tests for seven animals infected with the Toggenburg orbivirus, using reference strains of BTV1-24.**

**Post infected animals would normally be expected to show neutralisation titres of at least 1/256**

# Phylogenetic analysis of full length

**Seg-2 of BTV serotypes** *But will TOV reassort with other BTVs ?*





**Hofmann et al (2008) Genetic Characterization of  
Toggenburg Orbivirus, A New Bluetongue Virus, from  
Goats, Switzerland.**

**Emerg Infect Dis, doi 10.3201/eid1412.080818; in press.**



# Latest News

- BTV-1 identified in Algeria
- BTV – 9 identified in Libya (and typed)
- BTV in Israel (untyped – samples requested)
- BTV-16 identified in Greece
- BTV-8 identified in Greece

# Activities in 2008

- Publish full genome sequence analysis of BTV-8 from Netherlands (Virology **377**, 308-318)
- Identification of reassortant BTV-16 /BTV-2 in Italy (Virus Res. **137**:56-63.)
- Identify BTV-6 (vaccine strain) in The Netherlands (first time in Europe)
- Identify reassortment between strains of BTV-6 and 8 in the Netherlands
- BTV in Libya (typed but awaiting official-OIE announcement)
- BTV-1 in Algeria
- BTV-16 in Greece
- BTV-8 in Greece (for first time)
- BTV-25 identified in Switzerland
- Set up Relais system for BTV
- Sequence full genome of 100 BTV strains from around the world, as reference, to help identify reassortment events.
- Identified several reassortment events in eastern Mediterranean field strains