

# The Emergence of Exotic Bluetongue

## the importance of vaccination and future threats

**Uwe Mueller-Doblies, Narender Maan, Kyriaki Nomikou, Sushila Maan, and Peter Mertens**

**Biosecurity Group and Vector Borne Diseases Programme,  
Institute for Animal Health , Pirbright UK.**

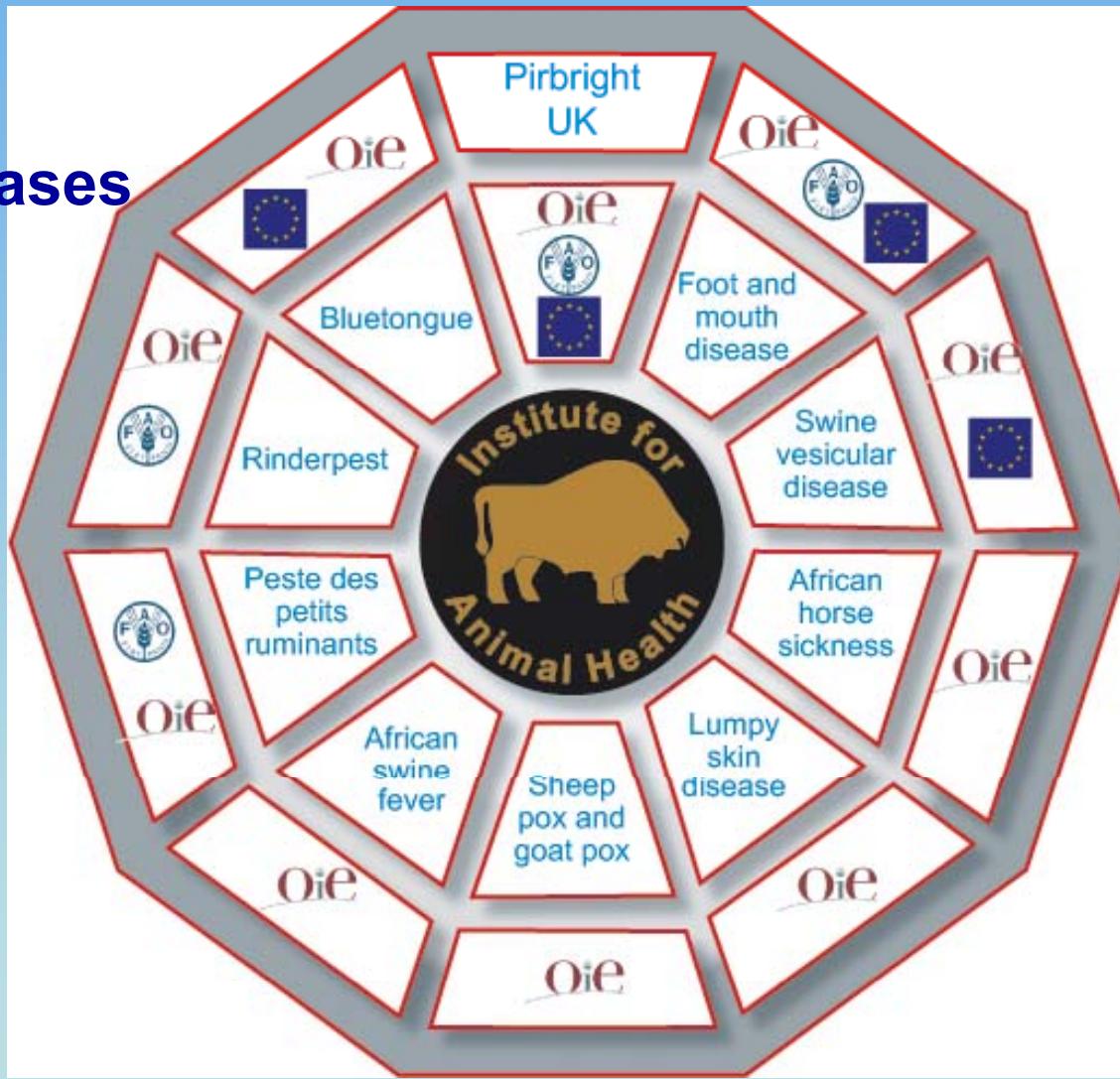


**USDA ARS 1<sup>st</sup> International Biosafety & Biocontainment Symposium**

# The reference labs at IAH-Pirbright

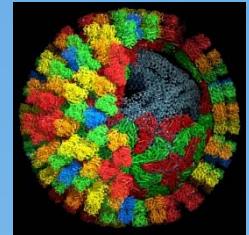
## Reference labs in Vector-borne Diseases Programme

- Bluetongue
- African Horse Sickness
- Lumpy Skin Disease
- Sheep and Goat Pox
- African Swine Fever





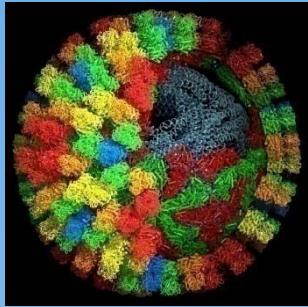
# Overview



- **Bluetongue Disease**
- **Blue Tongue Disease Viruses**
- **Dynamic of recent emergence in Europe**
- **Aspects of Emergence**
  - climatic changes
  - vector competency
  - live attenuated vaccines
  - recombination
- **Biosafety Issues**
-

# Overview

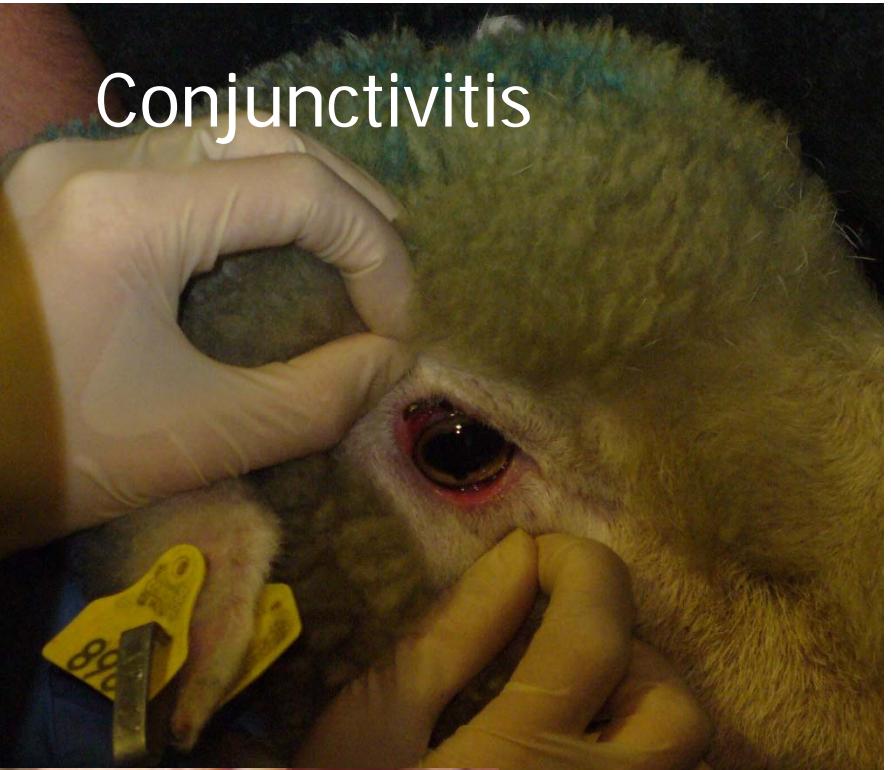
- Bluetongue Disease
- Blue Tongue Disease Viruses
- Dynamic of recent emergence in Europe
- Aspects of Emergence
- Biosafety Issues



# **Bluetongue: Clinical signs and disease in Europe**

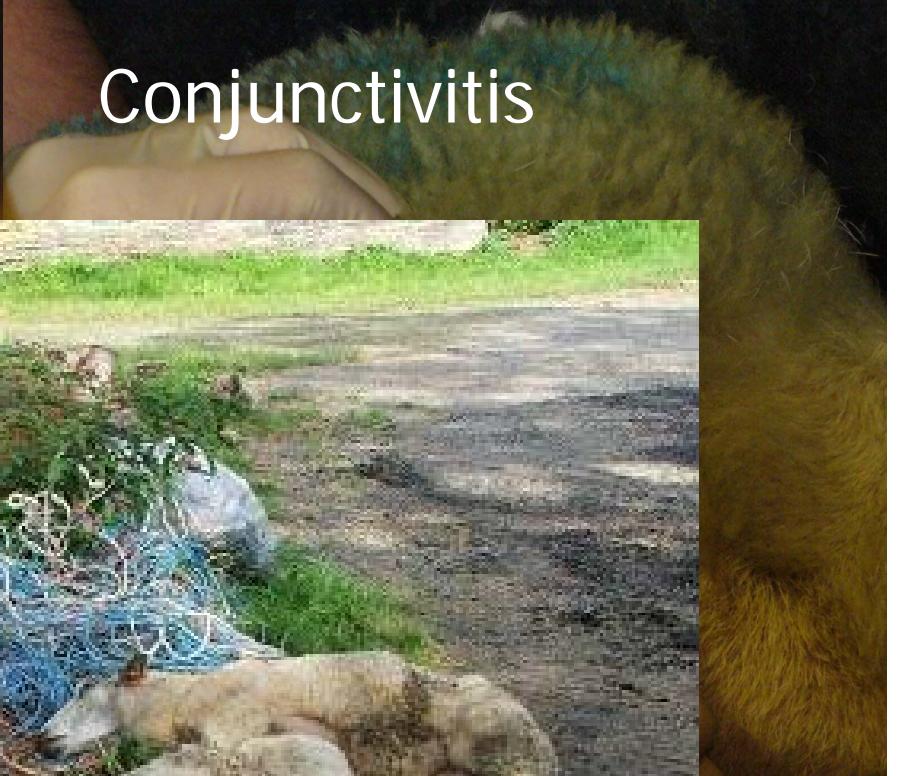
**“On a global basis, the second most economically important disease of livestock after Foot and Mouth”**







Coronitis



Conjunctivitis

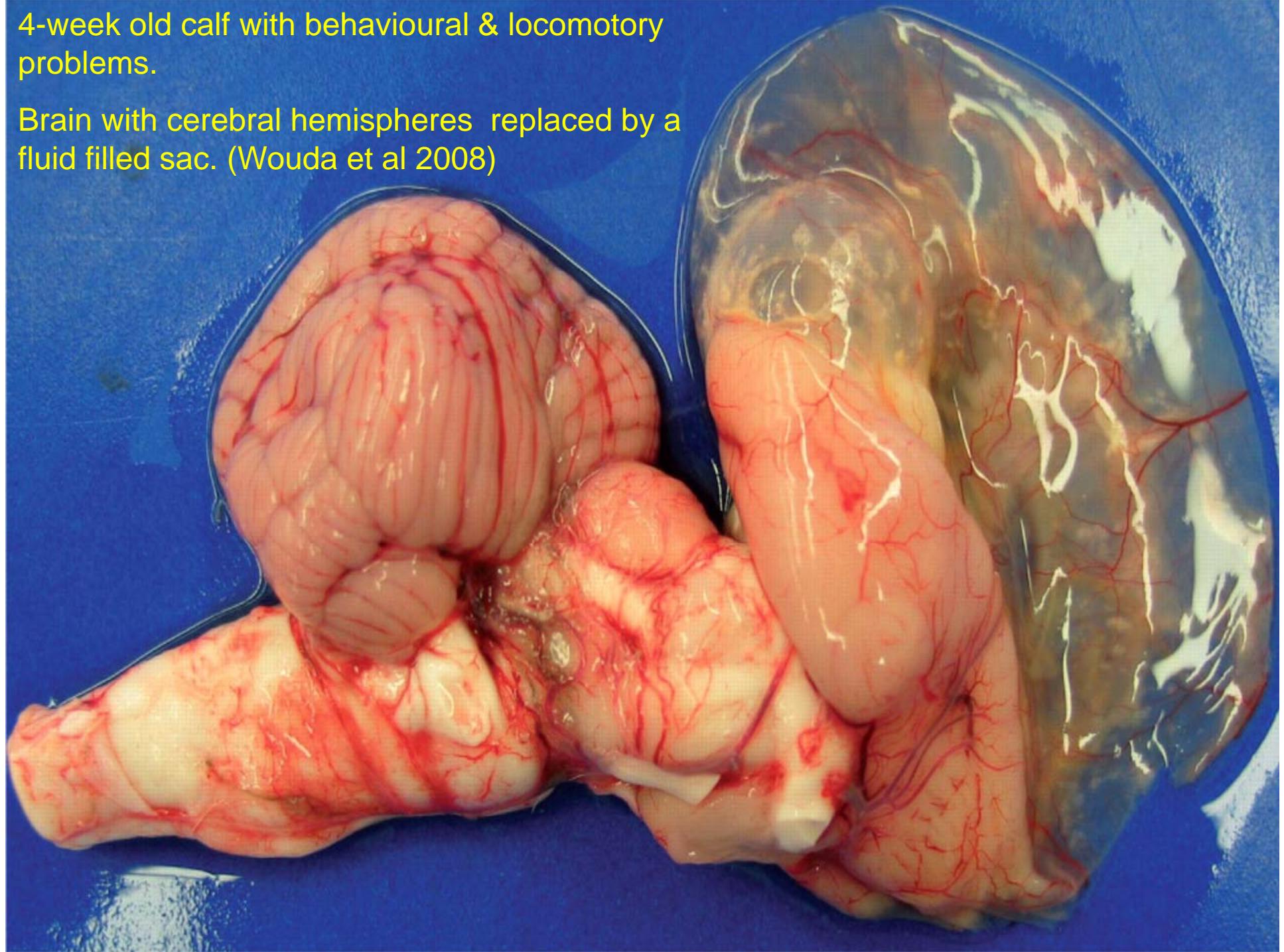


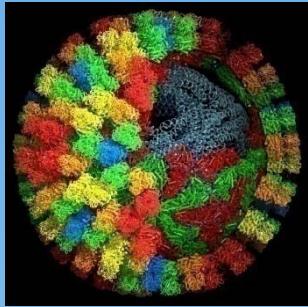


: Dummy calf at pasture 1b - Williamson, et al Vet Record 20

4-week old calf with behavioural & locomotory problems.

Brain with cerebral hemispheres replaced by a fluid filled sac. (Wouda et al 2008)





# **Bluetongue virus:** Classification and distribution



## Classification of *Orbivirus* species (serogroups)

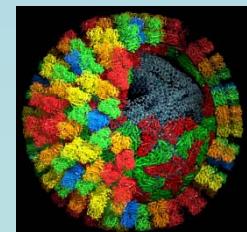
- 1-African horse sickness virus (AHSV)
- 2-Bluetongue virus (BTv)
- 3-Changuinola virus (CGLV)
- 4-Chenuda virus (CNUV)
- 5-Chobar Gorge virus (CGV)
- 6-Corriparta virus (CORV)
- 7-Epizootic hemorrhagic disease virus (EHDV)
- 8-Equine encephalosis virus (EEV)
- 9- Eubenangee virus (EUBV)
- 10-*Ieri* virus (IERIV)
- 11- Great Island virus (GIV)
- 12- Lebombo virus (LEBV)
- 13-Orungo virus (ORUV)
- 14-Palyam virus (PALV)
- 15- St Croix River Virus (SCRV)
- 16-Umatilla virus (UMAV)
- 17-Wad Medani virus (WMV)
- 18-Wallal virus (WALV)
- 19-Warrego virus (WARV)
- 20-Wongorr virus (WGRV)
- 21- Peruvian horse sickness virus (PHSV)
- 22- Yunnan Orbivirus (YUOV)

**Total: 22 virus species, 161 serotypes**

### Tentative species / unassigned viruses

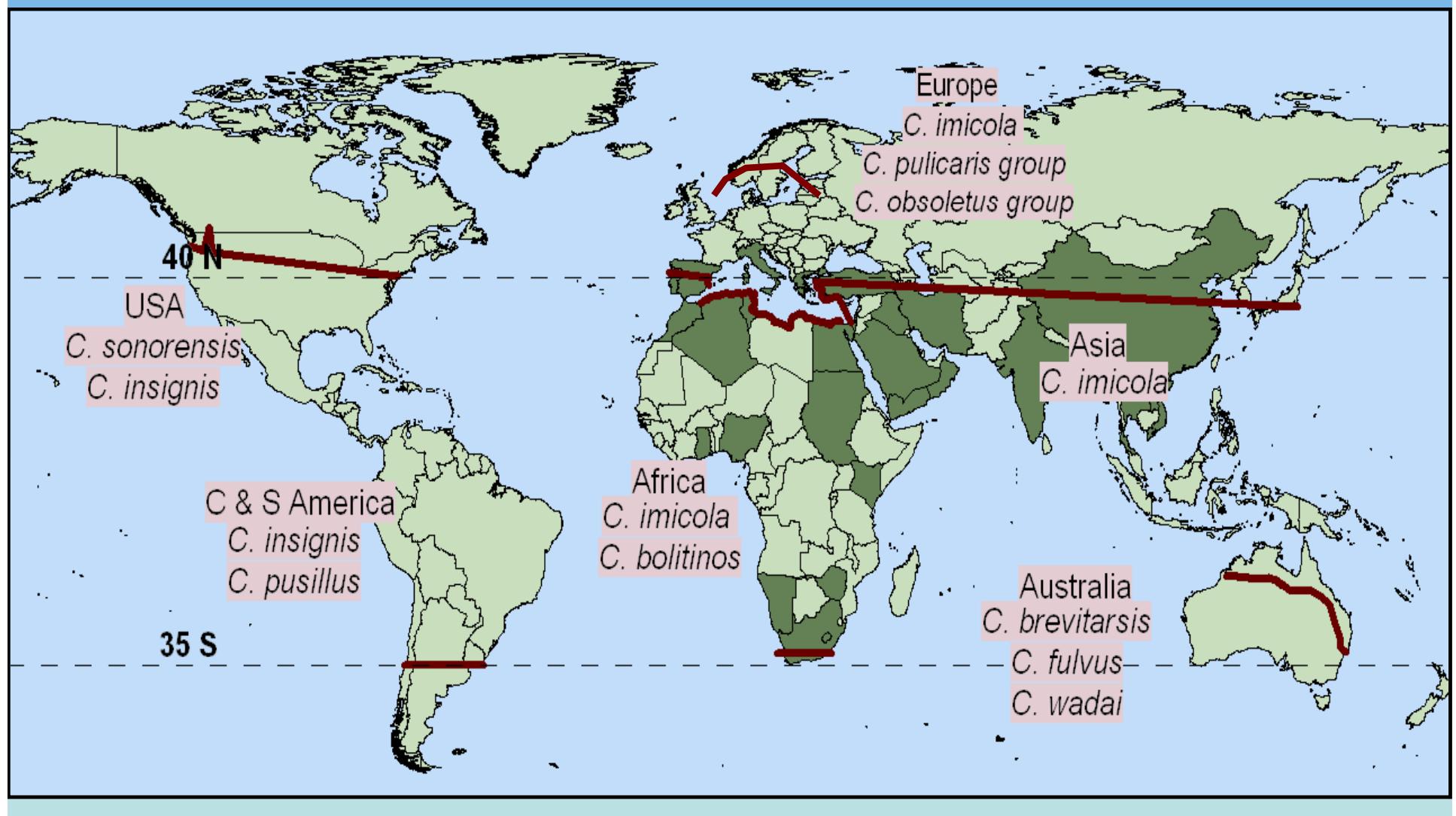
- Andasibe virus (ANDV)
- Ife virus (IFEV)
- Itupiranga virus (ITUV )
- Japanaut virus (JAPV)
- Kammavanpettai virus (KMPV)
- Lake Clarendon virus (LCV)
- Matucare virus (MATV)
- Tembe virus (TMEV)
- Codajas virus (COV)
- Tracambe virus (TRV)

Total : 12 unassigned viruses



## Global distribution of the 25 Bluetongue virus serotypes

# The global distribution of Bluetongue virus (pre 2006)



## ***Culicoides* vector species from Europe, Africa and the USA**

*C. obsoletus*



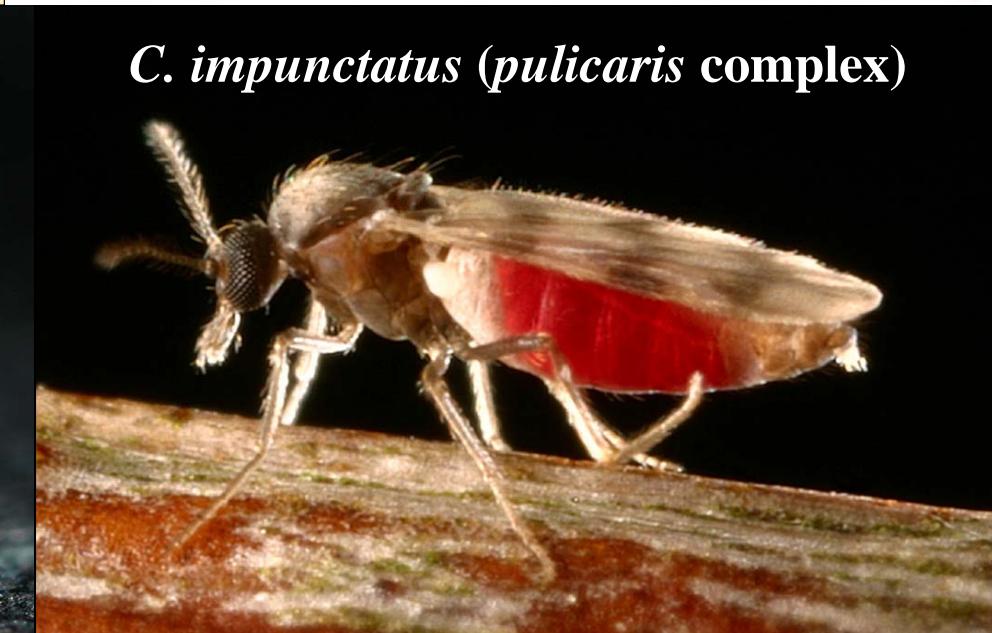
*C. sonorensis*



*C. imicola*



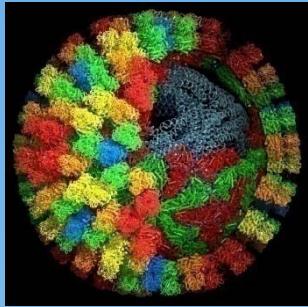
*C. impunctatus (pulicaris complex)*



Adult female *Culicoides nubeculosus* (from the insect colony at IAH Pirbright)



~1mm

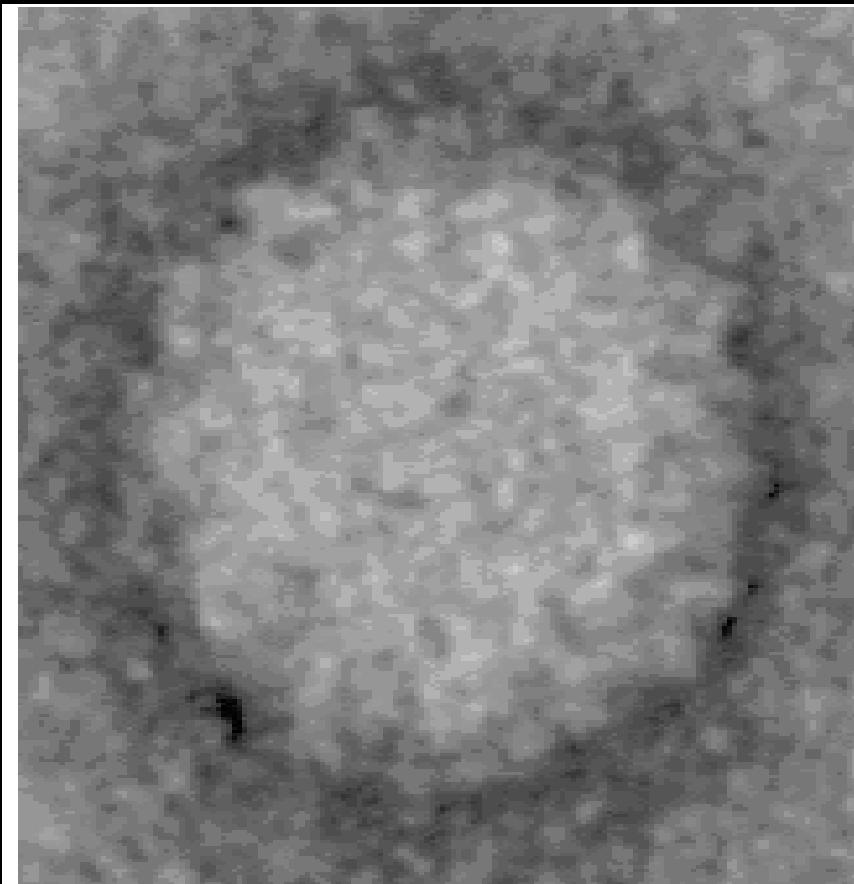


# **Bluetongue virus:**

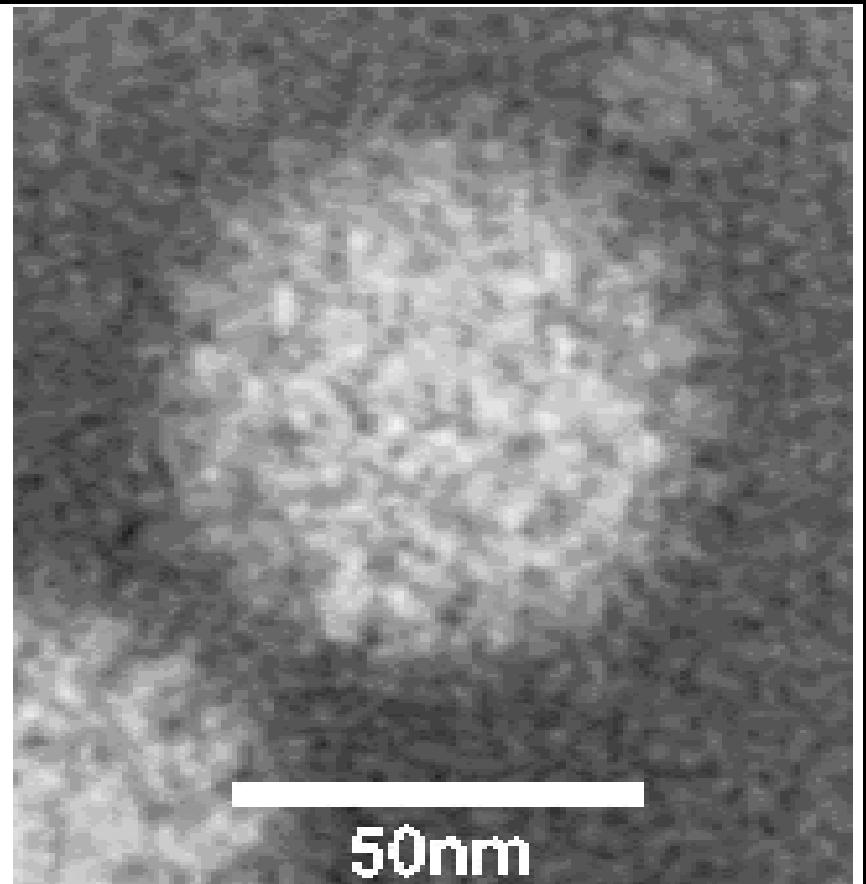
## Structure and serological properties



**BTV-1**  
**Virus particle**



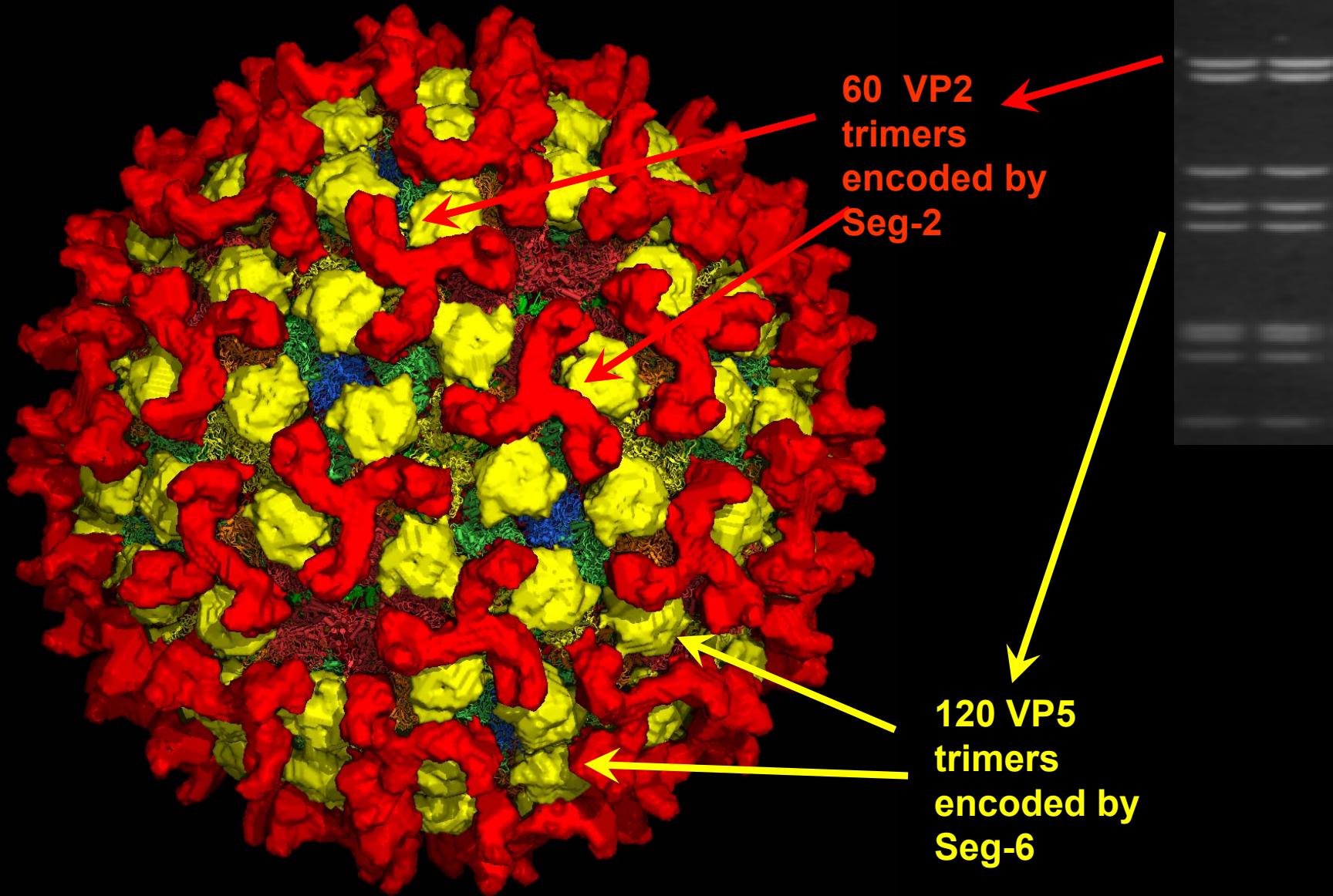
**BTV-1**  
**Core particle**



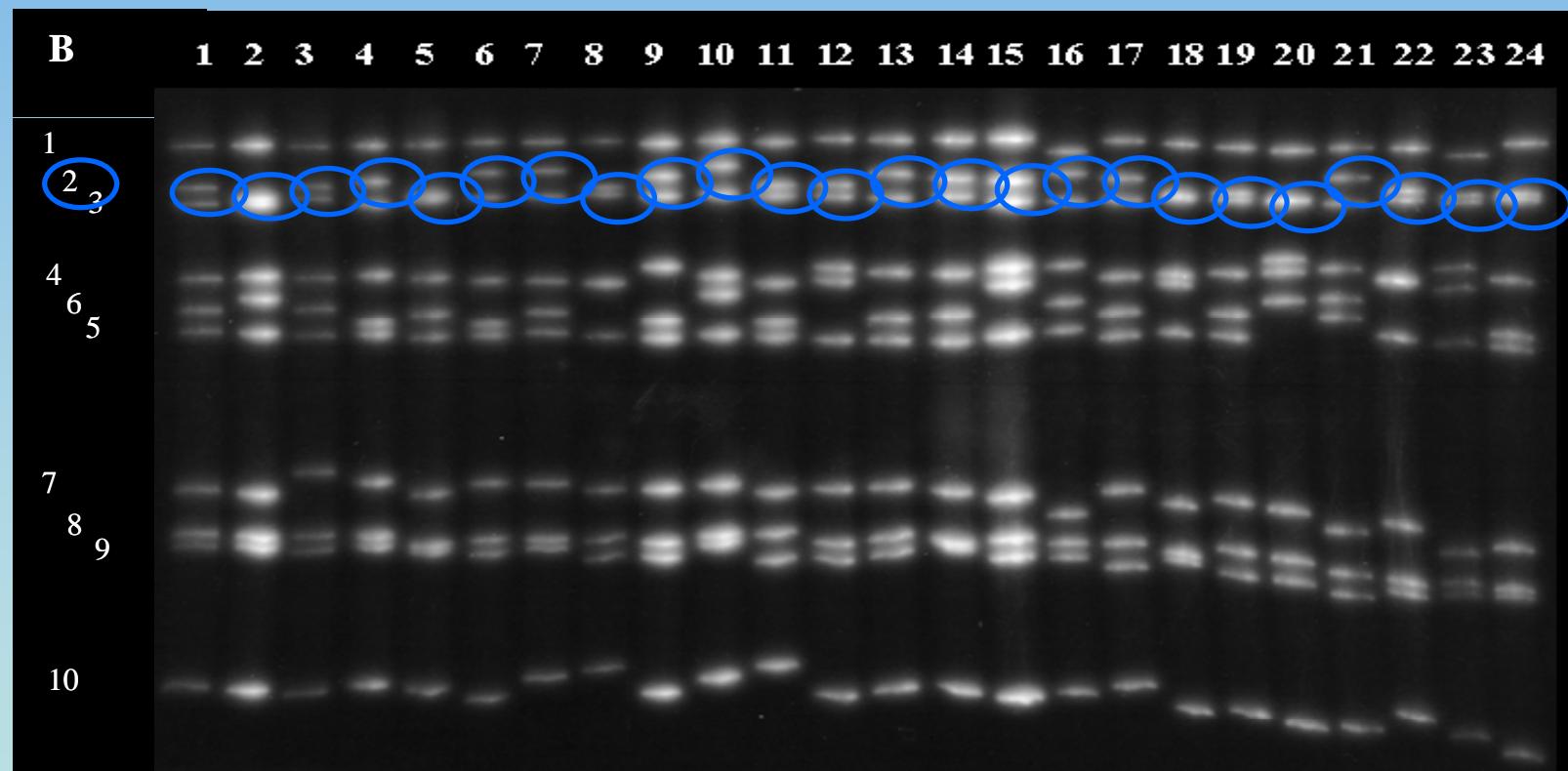
# Assembly of the major bluetongue virus capsid proteins



The outer surface of the BTV virus particle  
VP and VP5 - Encoded by segments 2 and 6  
(Cryo-EM and x-ray crystallography reconstruction)

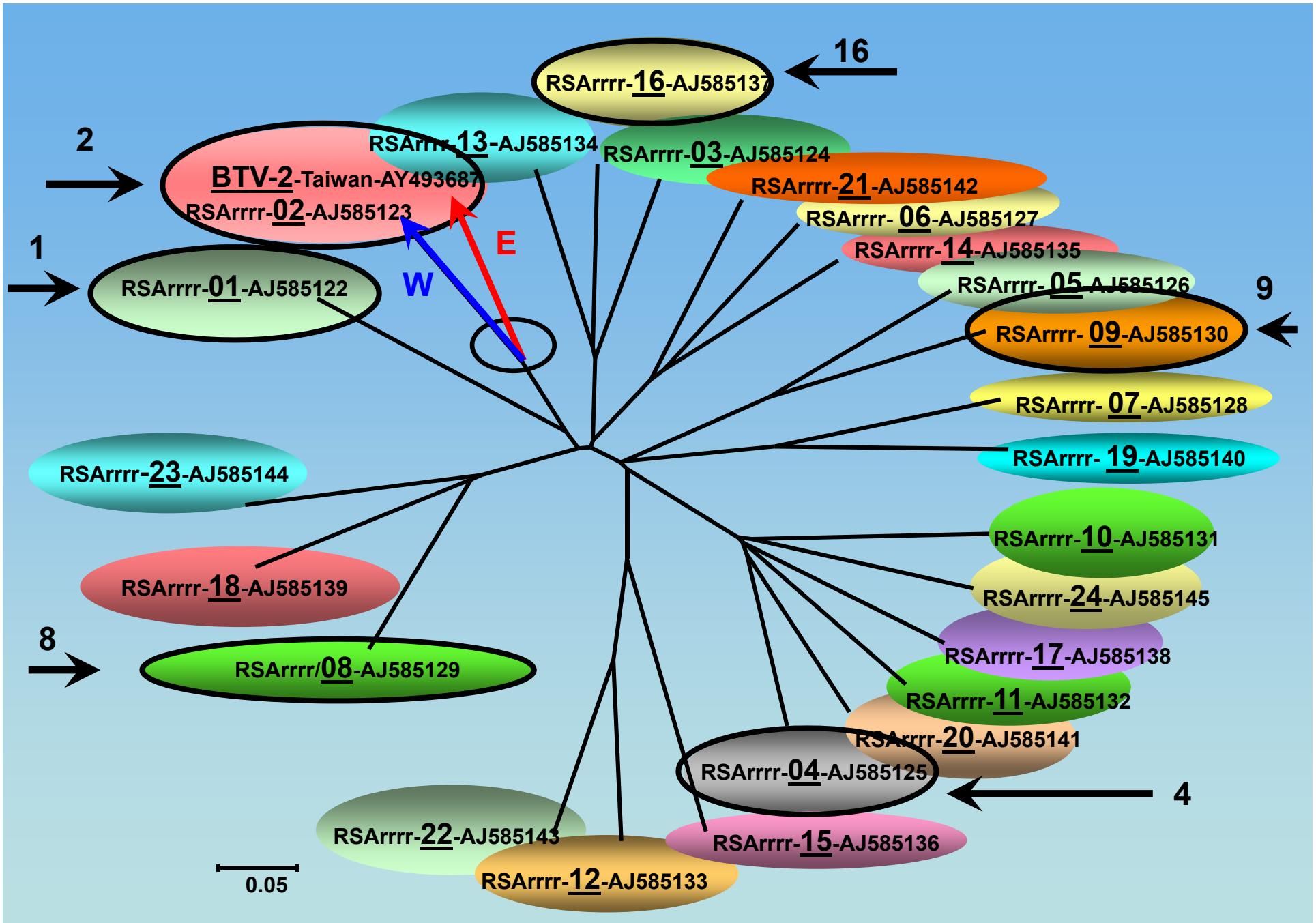


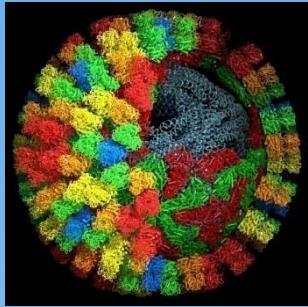
# The Genome segments of the 24 BTV serotypes analysed by SDS-PAGE



# Primers for identification of European BTV serotypes by Rt-PCR since 1998

Type	FWD Primer	RVS Primer	Product BP)
1	BTV-1/2/p406F GAGAACARGAARAATATAT	BTV-1/2/p968R TYATACGTTGAGAAGTTTGT	1686 (1A)
2	BTV-2/2/p198F TATCAGTTAATAGTCATT	BTV-2/2/p416R TACTAAAKATATACTTCTCCGT	653 (2A)
3C	BTV-3/2/p340F GAACTAATGAGGACAGAT	BTV-3/2/p840R TGAGATCGTCGTCTGAAGAT	1500
4	BT4/67/F GCTTAACATAAACCAACGAGG	BT4/775/R GTATCACCTGACCGCGTCG	2124
5	5NIG/123/F TACGGATCATAAATGGATGG	5NIG/791/R TATATGCTCACAGATATCT	2004
6	BT6/301/F GGTGGTATGTATAGAGGAAG	BT6/853/R CAAAGGGAACCTCGCGCGTAATC	1656
7	BT7/83/F GATGTATCCATAGCGGCAT	BT7/853/R GATAGATAAGCGAATCGAG	2310
8	BT8/101/F GAATGATGGTCATAGCGAG	BT8/500/R CGATGTGCGCATTCTCTC	1197
9	BTV-9/2/p491F GGSAATATATTCTRATGG	BTV-9/2/p867R GGAAAACGTATTGGCAAT	1128 (9C)
10	BT10/183/F GATGTTCCACATCTTACAG	BT10/803/R GTGAAACCTAATGAAATTG	1860
11	BT11/147/F TGTATTGTTAAGGCTAGG	BT11/770/R GTCATCGTATAGTATCAT	1569
12	BT12/111/F GGATCACAAATATAGATGTG	BT12/776/R CTACGATCATATGATAACTC	1995
13A	BT13/157/F CGAGGAAAGCGGATACCAC	BTV-13/2/p631R TGCTTAATGCTCAAACCTCG	1422
14	BT14/121/F GAAGGTTAGCTTAGGTTG	BT14/805/R CTCCGCTTCATCCAGCTC	2052
15	BT15/343/F GTGGCAGAACGCAGAGGCAG	BT15/806/R G TGAGACATATAATGTTCAAG	1389
16B	BTV-16/2/p266F GCGATAAGAGATAMTTGGAT	BTV-16/2/p924R GCCGAAGGTGCGATCTGGCCG	1974 (-Nig)
17	BTV-17/2/p146F GCGAATGCTGCCAACGCTG	BTV-17/2/p653R GCGCGACTTCCTAAACG	1521
18	BT18/160/F GTCTTATCATATAGAGCCAG	BT18/828/R GTACTCAGATAATAGTCGAG	2004
19	BT19/190/F TGTGCTCAAGCAAGCGCGTAT	BT19/841/R GTCATCGTCGAGTGCCTG	1953
20	BT20/135/F CATTACTCGATAGATTACC	BT20/812/R GTACGTCGTCAGCAATCTG	2031
21	BT21/116/F GAGGAATGGCTGAAGTGG	BT21/754/R GCCTCCACACAGCGAAGACAG	1914
22	BTV-22/311F GACCGCGTTGGATAGGATAT	BTV-22/682R GCTTATACCTCGCATCATCG	1113
23	BT23/459/F GCTTAGACCTGGCGATAAG	BT23/653/R GTTTAACATGCATACTCAG	582
24	BT24/313/F TGGATTATCTACACGATT	BT24/783/R CATAAGCTCCAACTTCAAC	1410





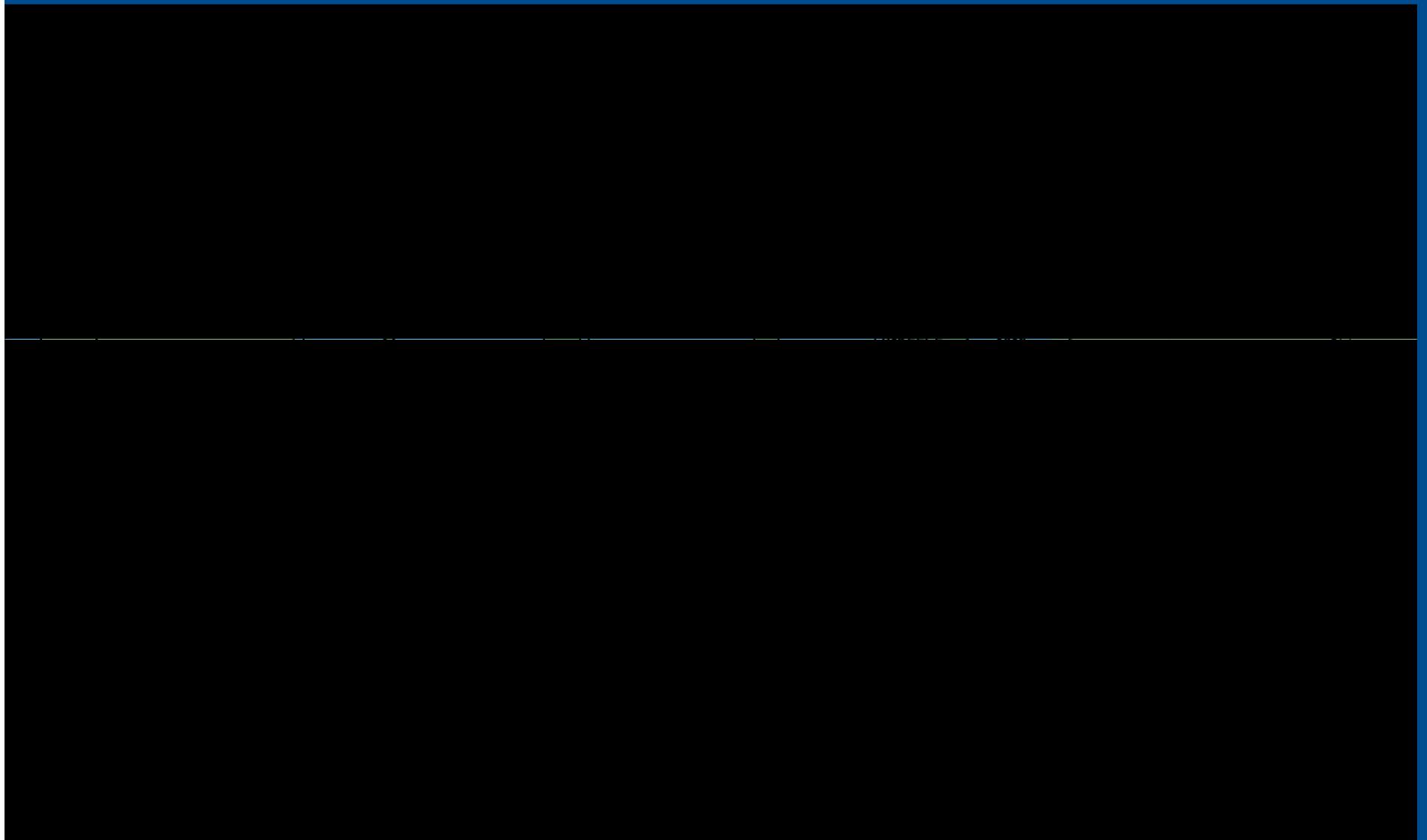
# BT in Europe 1998 to 2006: Transmission and spread

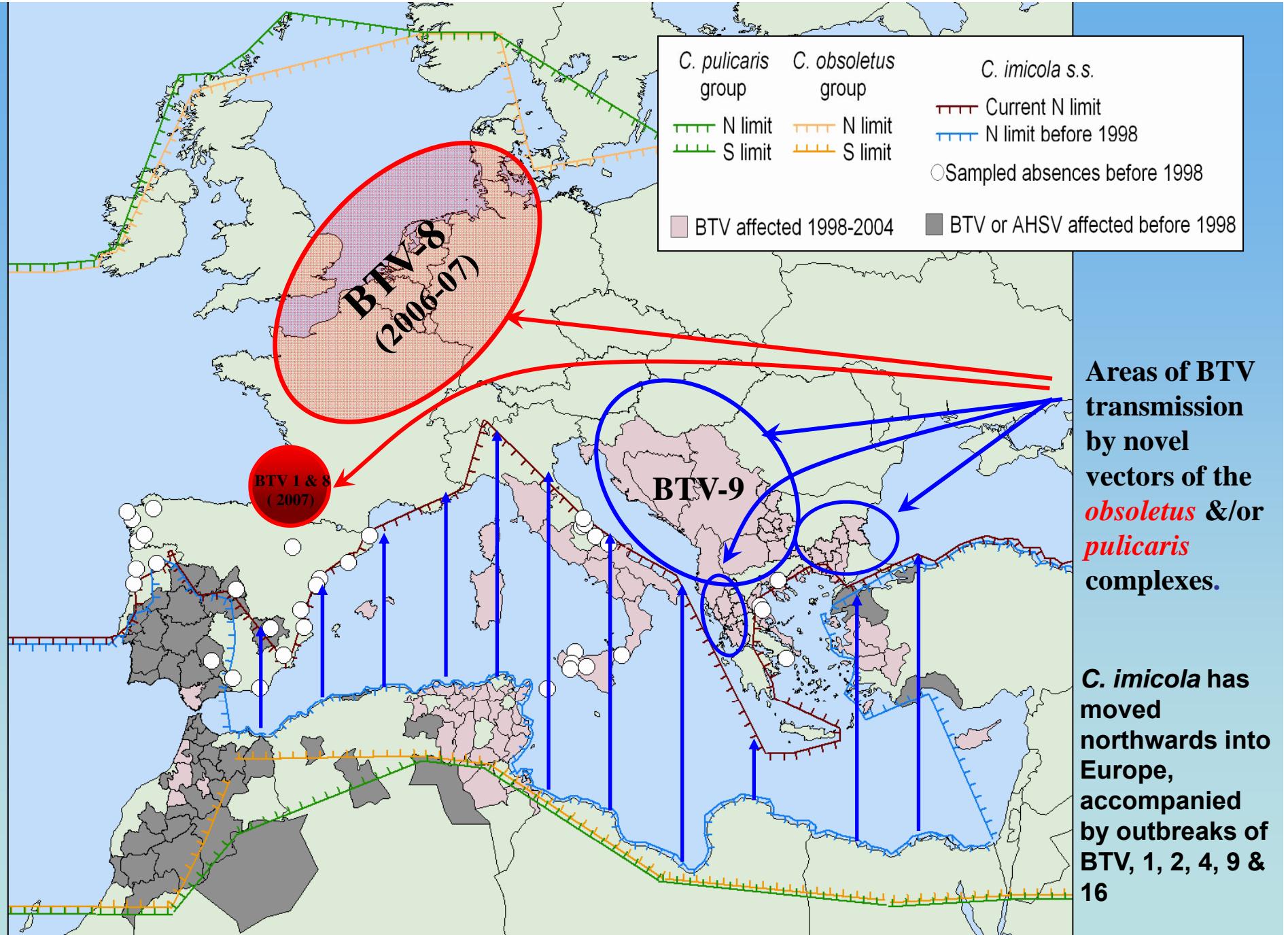


# **Northernmost Distribution of BTV to 1997**

BTV-10 in Spain & Portugal 1956 - 60, BTV-4 in Turkey and Cyprus during 1980's .  
BTV-3 in Lesbos and Rhodes 1979.

# Northern most distribution of *Culicoides imicola* (to 1997)

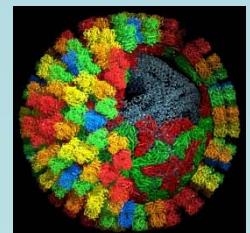




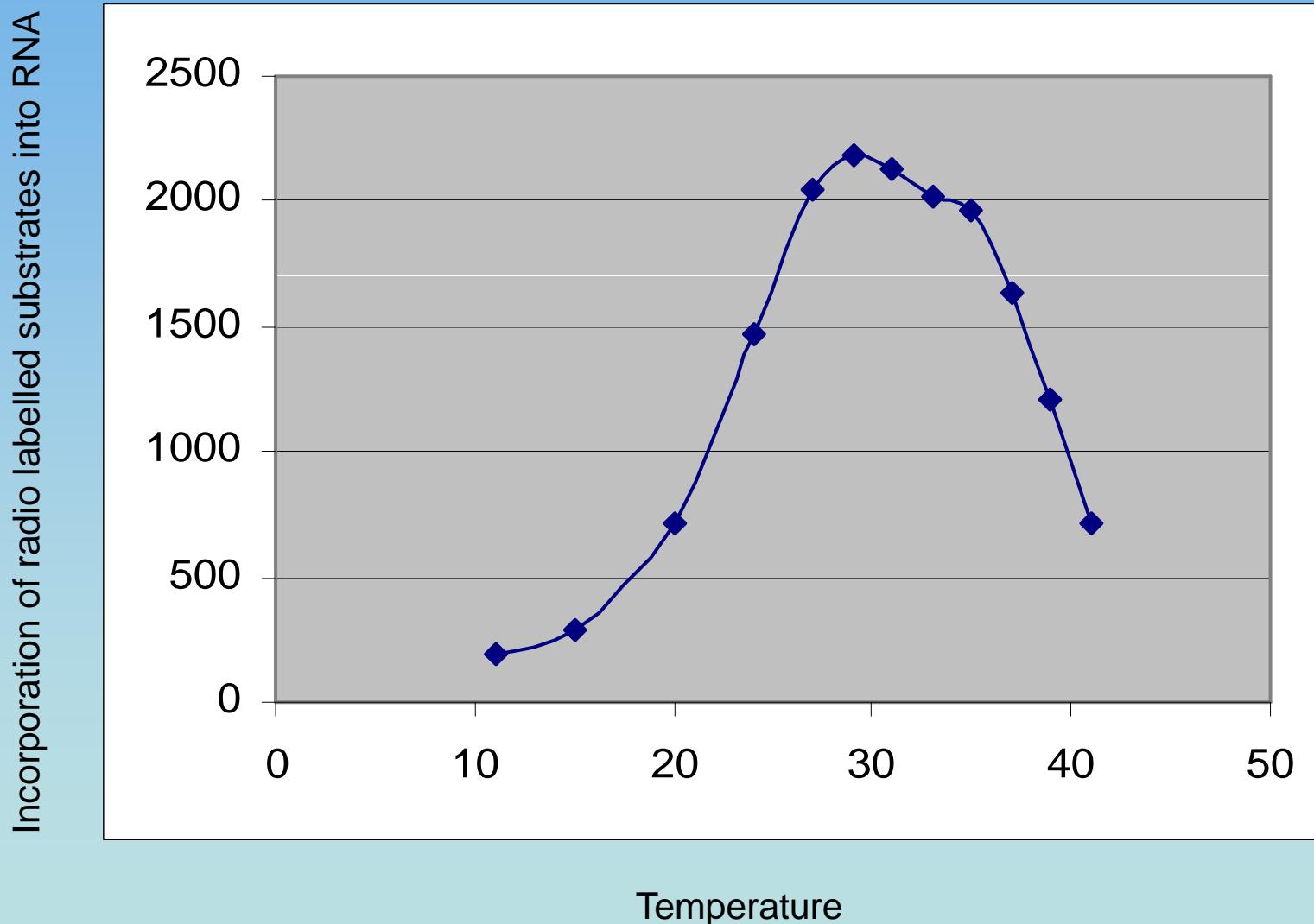


**15 distinct strains of nine different BTV serotypes have invaded Europe, with new introductions every year Since 1998**

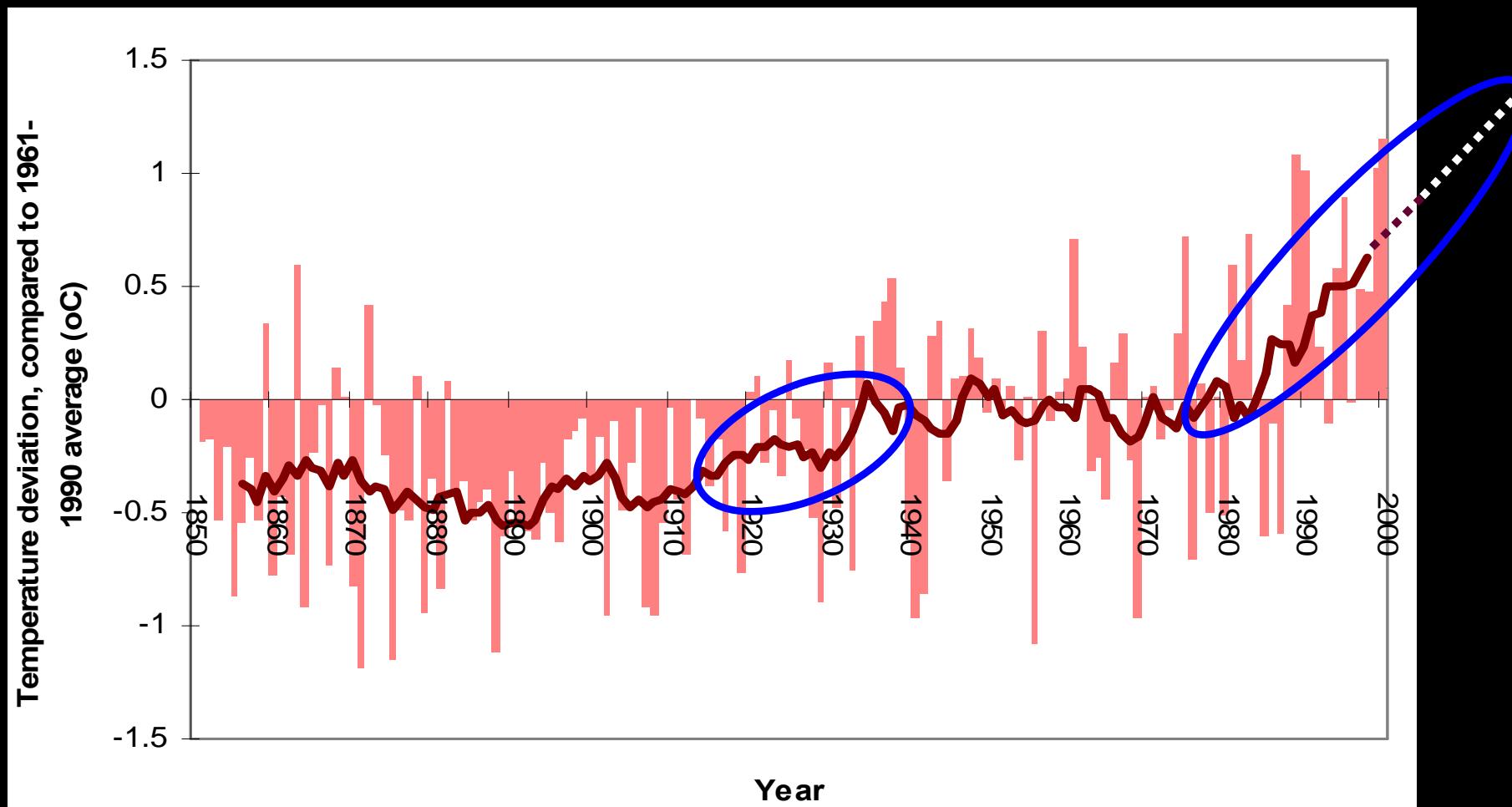
**This evidence supports that a fundamental ‘step-change’ in the epidemiology of BTV that has been related to climate change**



# BTV -1 polymerase activity



## Recent climate warming in Europe.

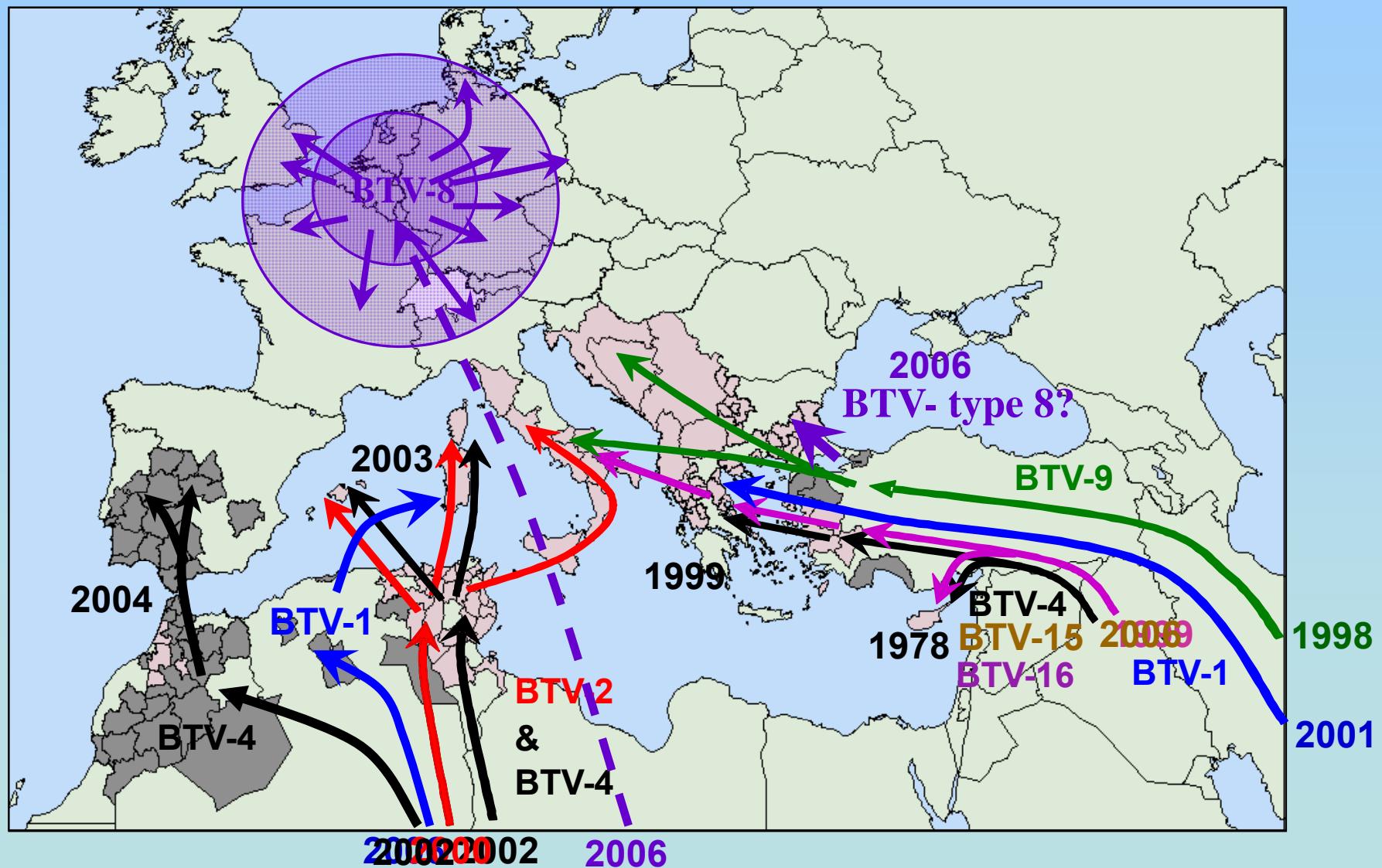


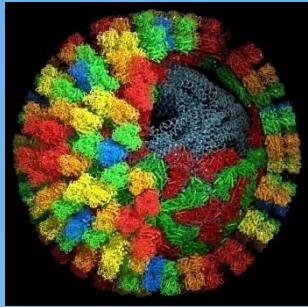
**Recent warming trend in Europe.** observed annual temperature deviations in Europe over the last 140 years (1850 to 2000) compared to the 1961 to 1990 average ( $^{\circ}\text{C}$ ) as light pink bars. Superimposed as a continuous dark pink line is the 10-year smoothed trend of these annual temperature deviations (reproduced with permission from EEA, 2004). (Purse et al 2005 Nature Reviews)

**Wind blowing dust from the Sahara across Italy and the Mediterranean Islands**



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



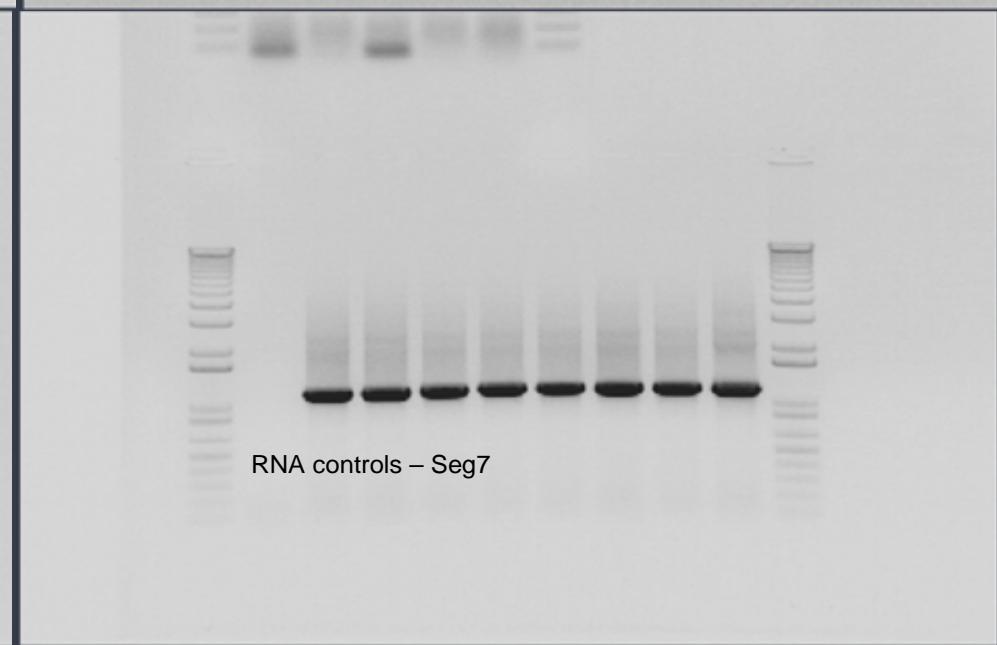
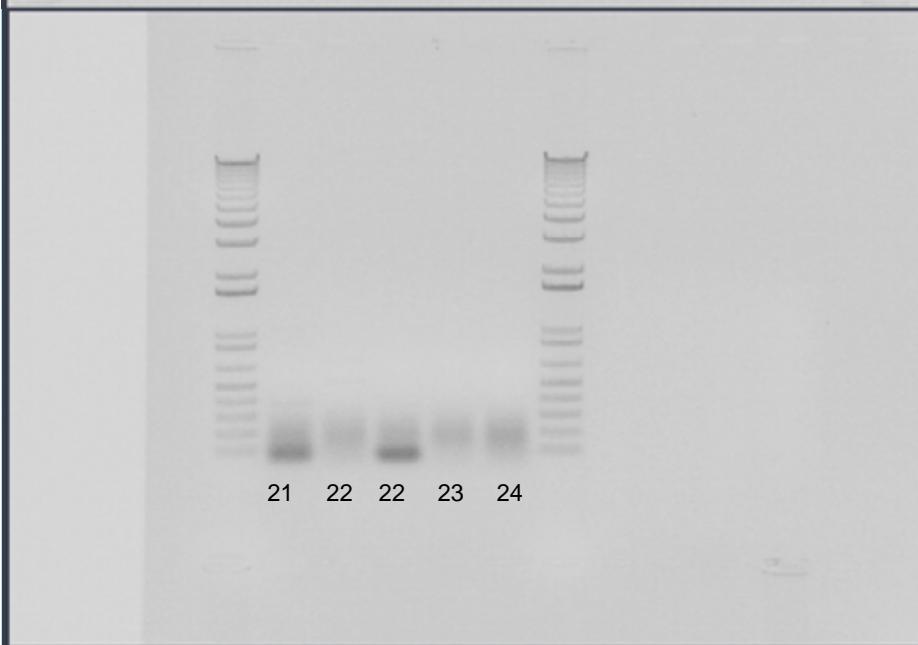
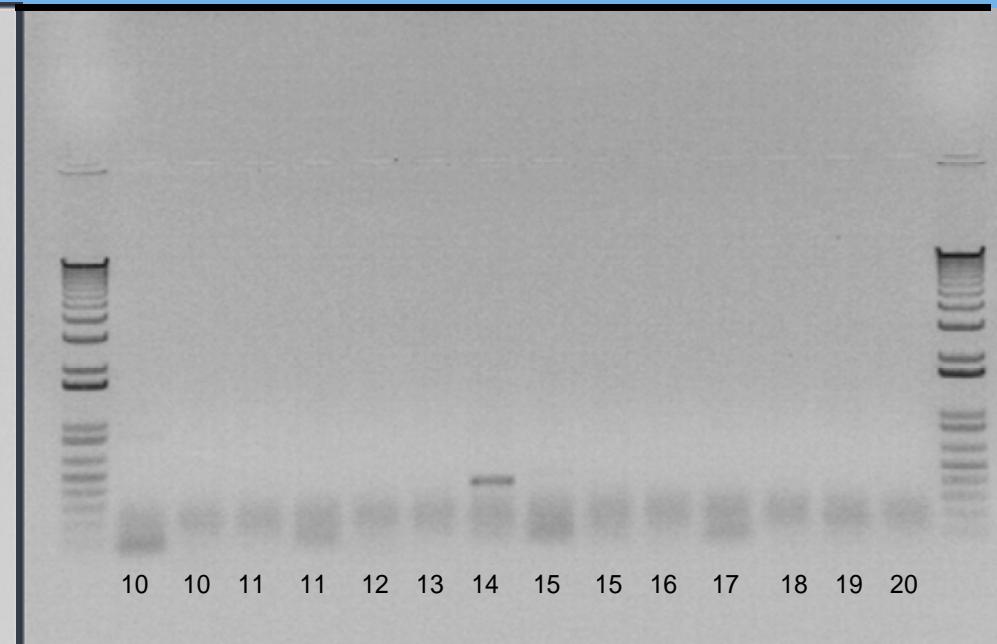
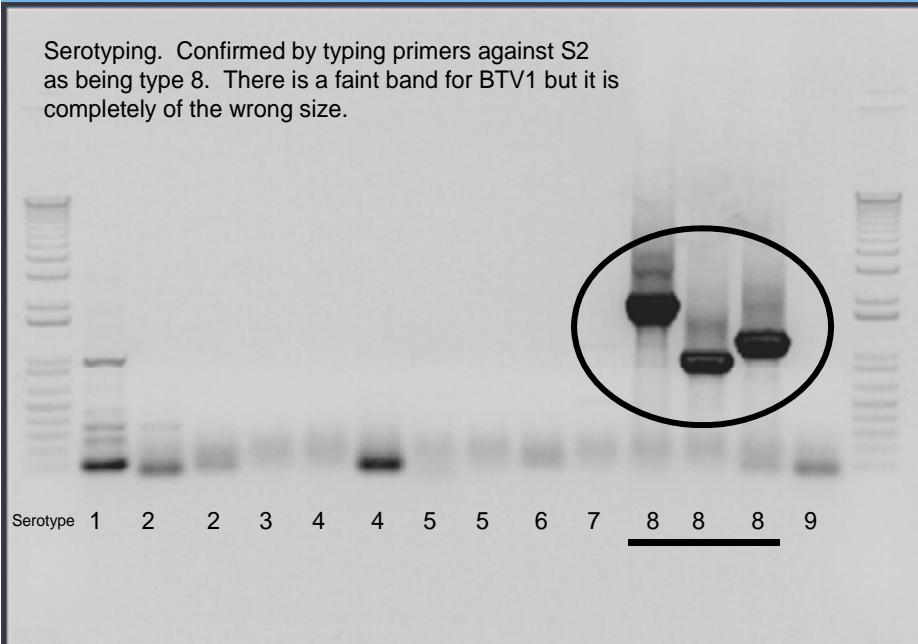


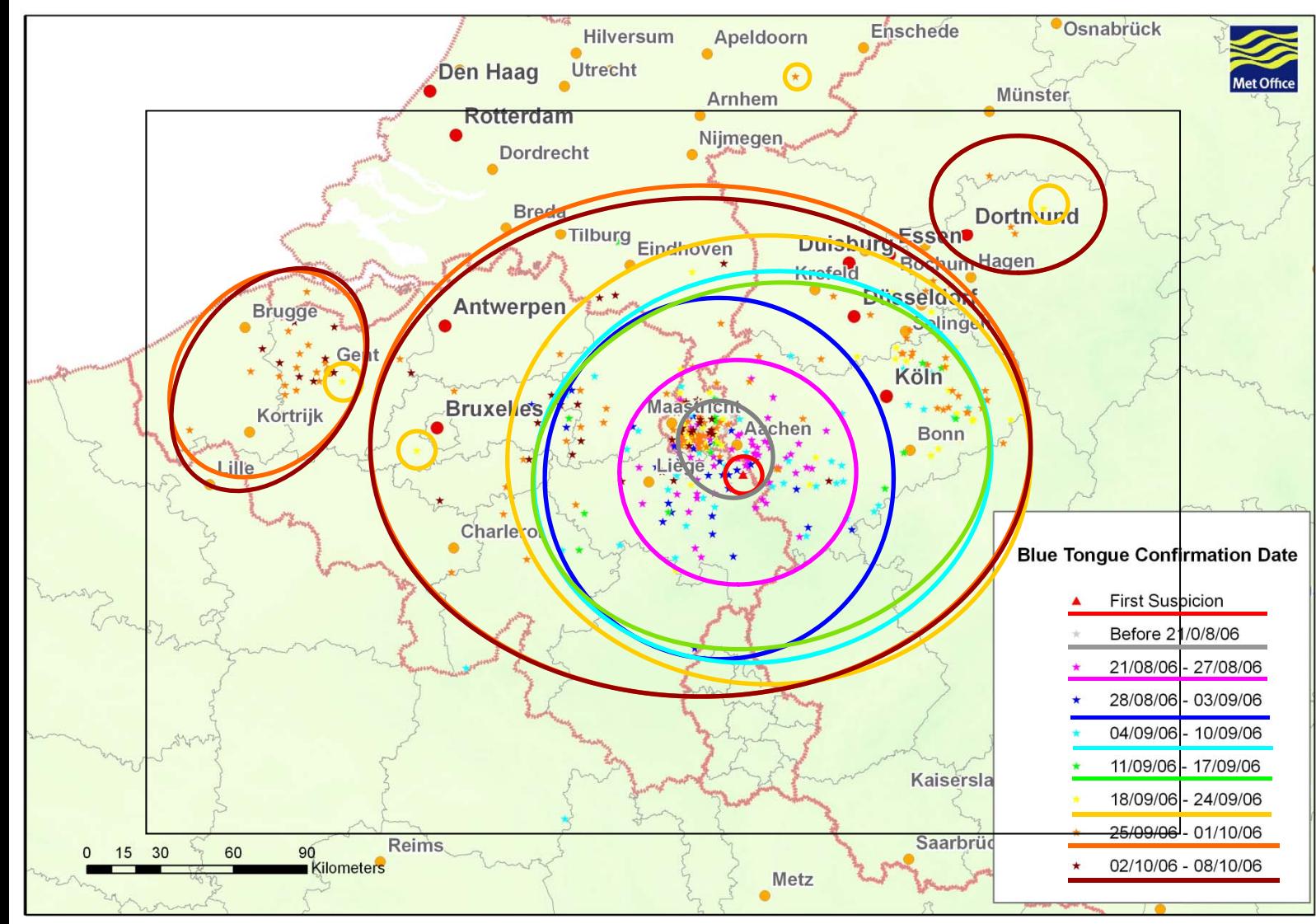
# BTV-8 in Europe: 2006 onwards



# Initial identification of NET2006/01 as BTV-8

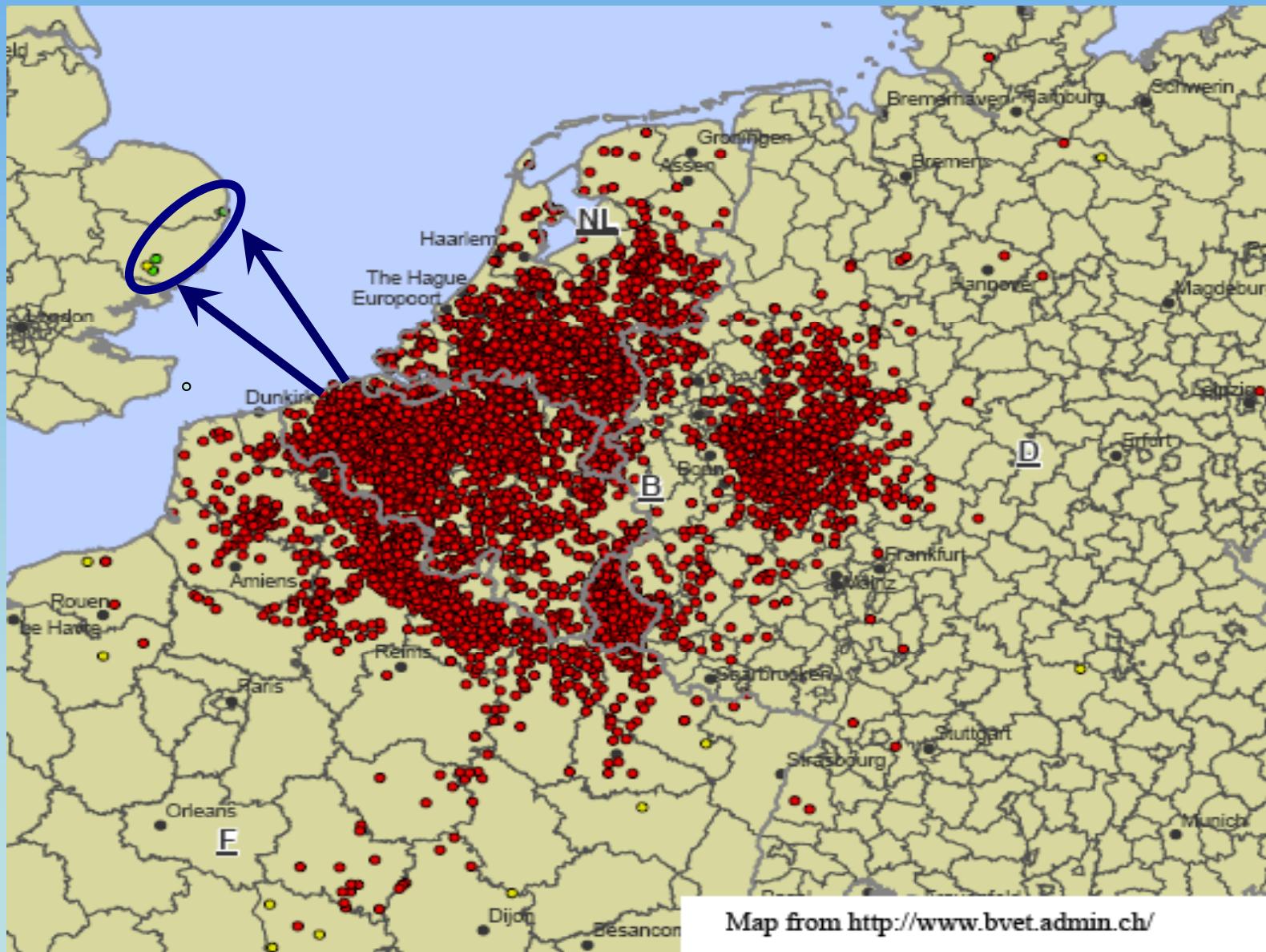
Serotyping. Confirmed by typing primers against S2 as being type 8. There is a faint band for BTV1 but it is completely of the wrong size.

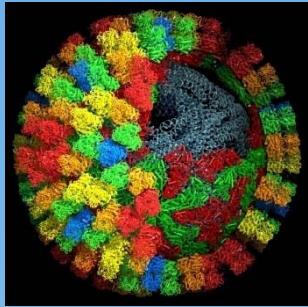




**The Spread of BTV outbreaks from 21<sup>st</sup> of August to 8<sup>th</sup> October 2006**

# European BTV situation 2007

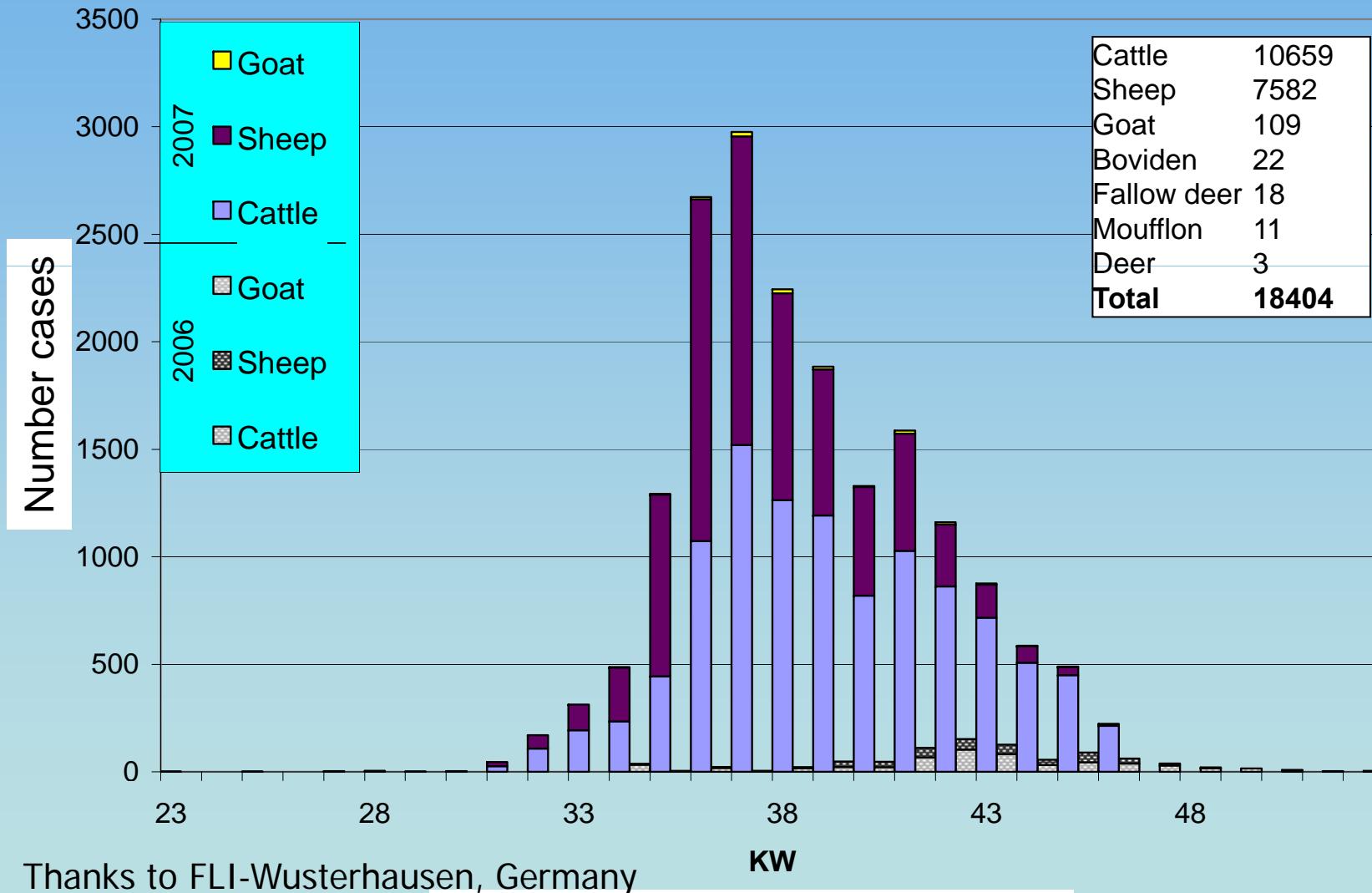




# BTV-8 in Germany



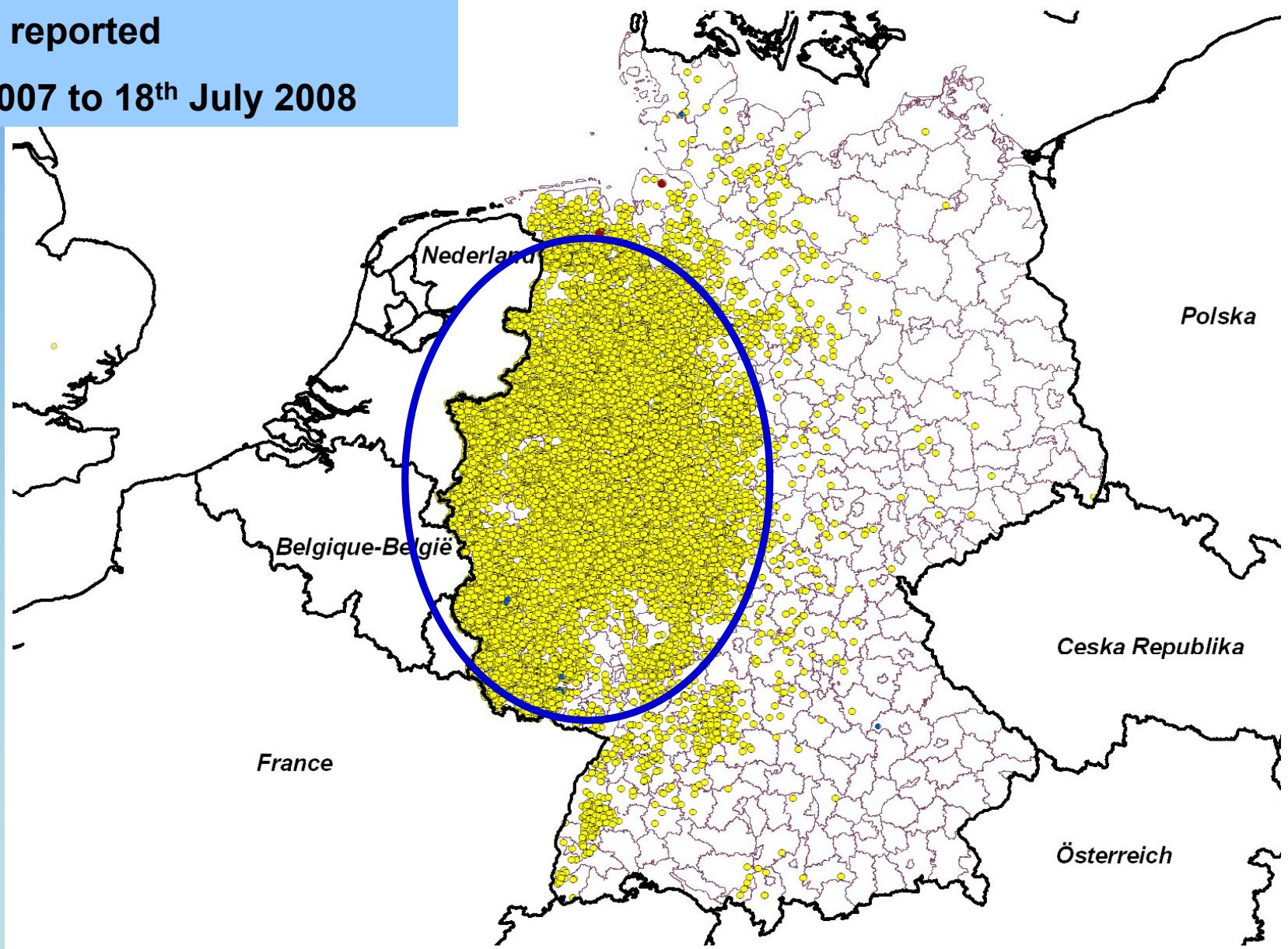
# Outbreaks 2006/2007



## Bluetongue in Germany

2033 cases reported

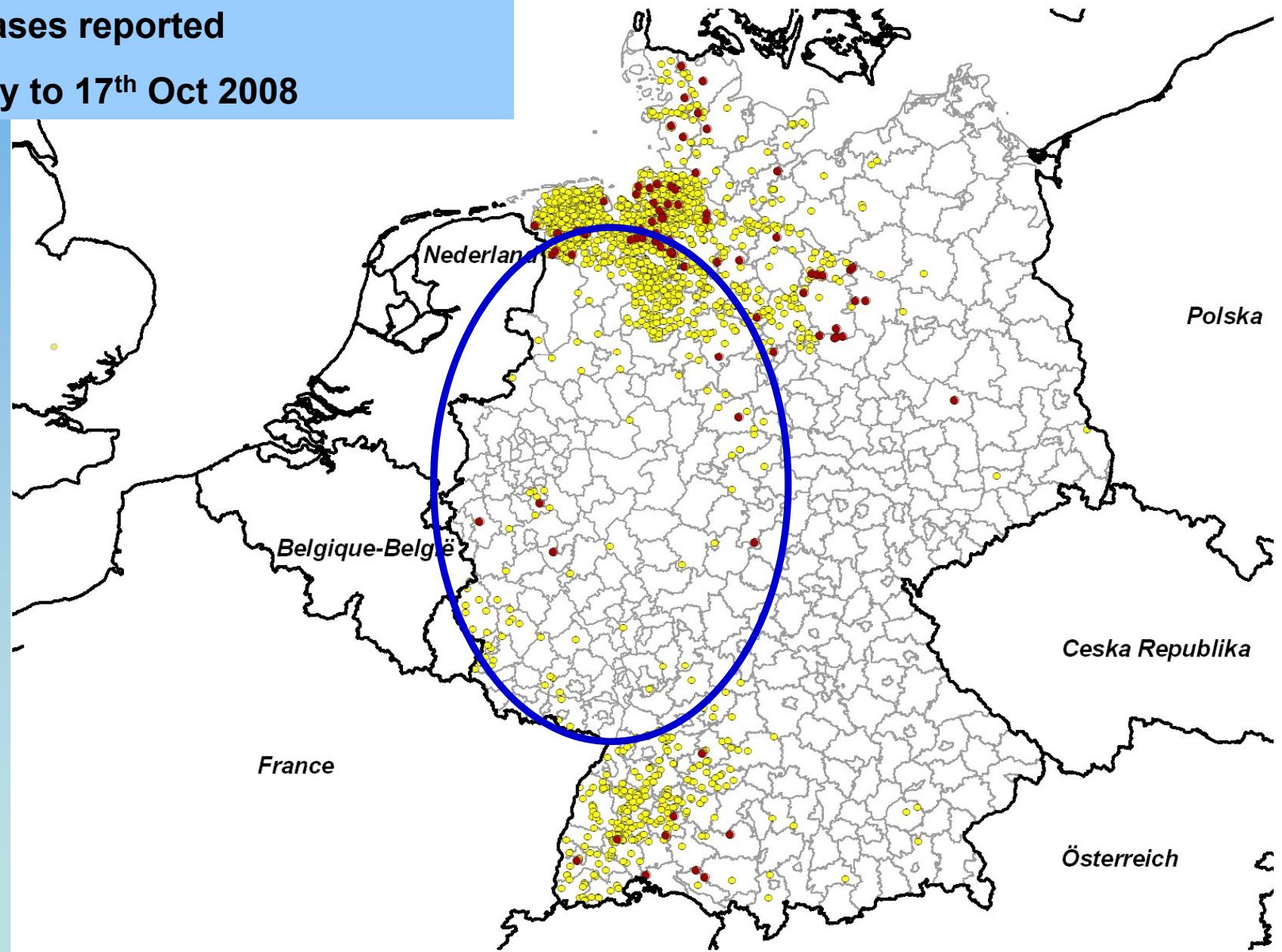
1st June 2007 to 18<sup>th</sup> July 2008

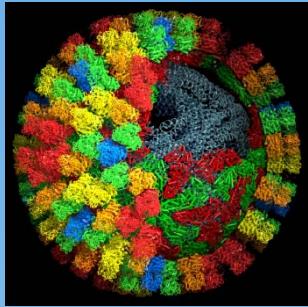


## Bluetongue in Germany

2033 cases reported

1st May to 17<sup>th</sup> Oct 2008

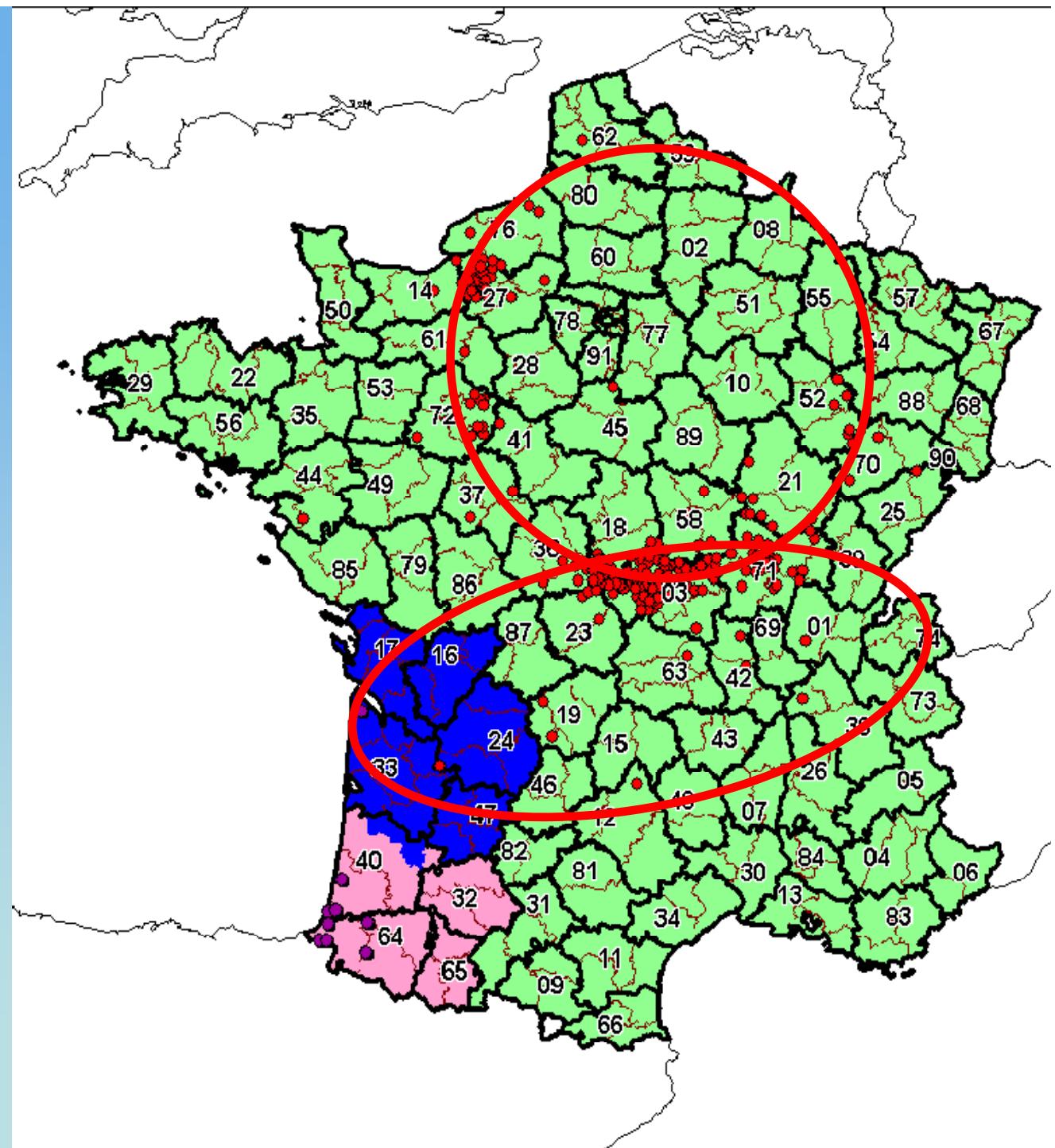




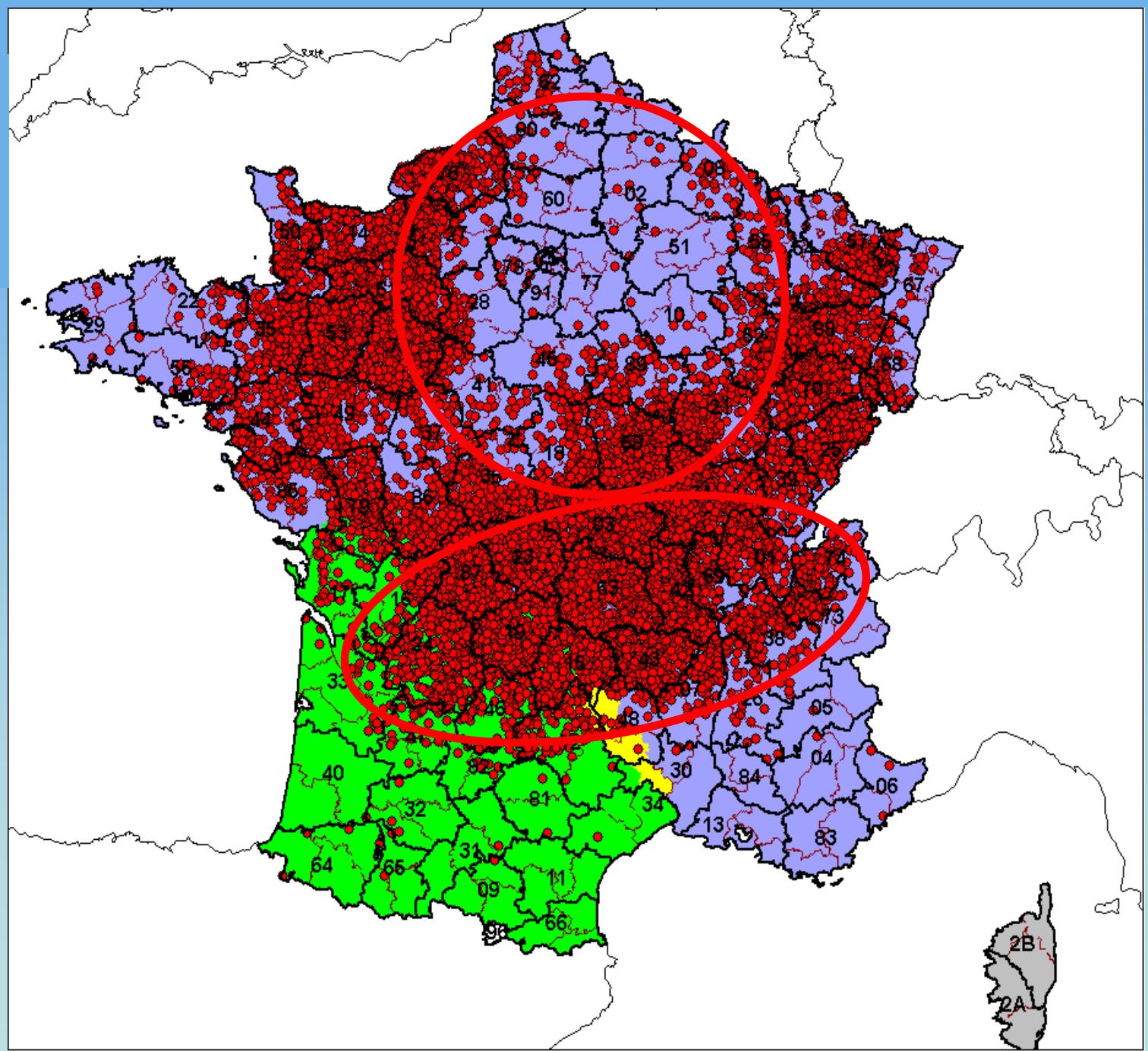
# BTV-8 in France



Outbreaks caused up to  
start of August 2008 by:  
**BTV-8** •  
and  
**BTV-1** •



**Outbreaks  
caused up to  
Late October  
2008 by :  
BTV-8 •**



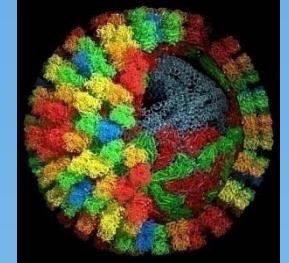
# BTV-8 outbreak Northern Europe

- **year 1:** low level initial spread (mainly in cattle), low mortality.
- **2nd year:** high level spread in cattle and sheep very severe.
- **3rd year:** Recovered animals but few susceptible animals remain, so fewer cases.
- **4<sup>th</sup> year (projected)** Few sources of infection remain and most animals immune. Potential for eradication

Vaccination



# BTV vaccines



- 3 Options
  - Live attenuated vaccine strains
    - (unsafe in Northern European sheep – transmitted in the field)
  - Inactivated virus vaccines
    - (used in the UK/France/Spain 2008/2009)
  - Next generation vaccines
    - (not yet commercially available)



# The effect of Vaccination with live attenuated BTV strains in UK sheep

## Pyrexia and viraemia duration recorded in Dorset Poll sheep after vaccination

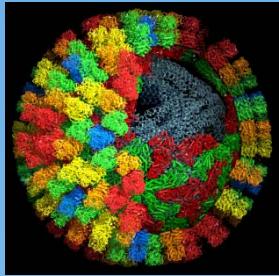
Serotype	No. of sheep showing pyrexia	Pyrexia duration range (days)	Max viraemia titre recorded*	Max viraemia duration (days)	Min viraemia duration (days)
BTV2	2/4	1	4.75	17	17
BTV4	2/4	2	4.25	16	11
BTV9	4/4	1-5	5.25	15	7
BTV16	4/4	6	6.83	19	17
BTV16	3/3	1	6.0	23	10

\*(log<sub>10</sub>TCID/50)

# Sheep inoculated with BTV-4 vaccine virus

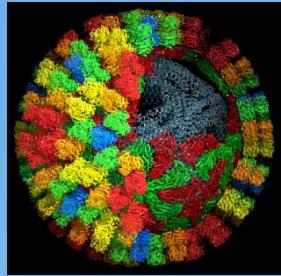


- Acute respiratory dysfunction
- Swelling and discolouring of the lips and tongue and copious salivation
- Cyanosis of the whole oral cavity including the tongue (acute oxygen deficiency)



**After compulsory vaccination, in France with inactivated vaccines, only three isolates of BTV-8 and no isolates of BTV-1 were made during 2009**

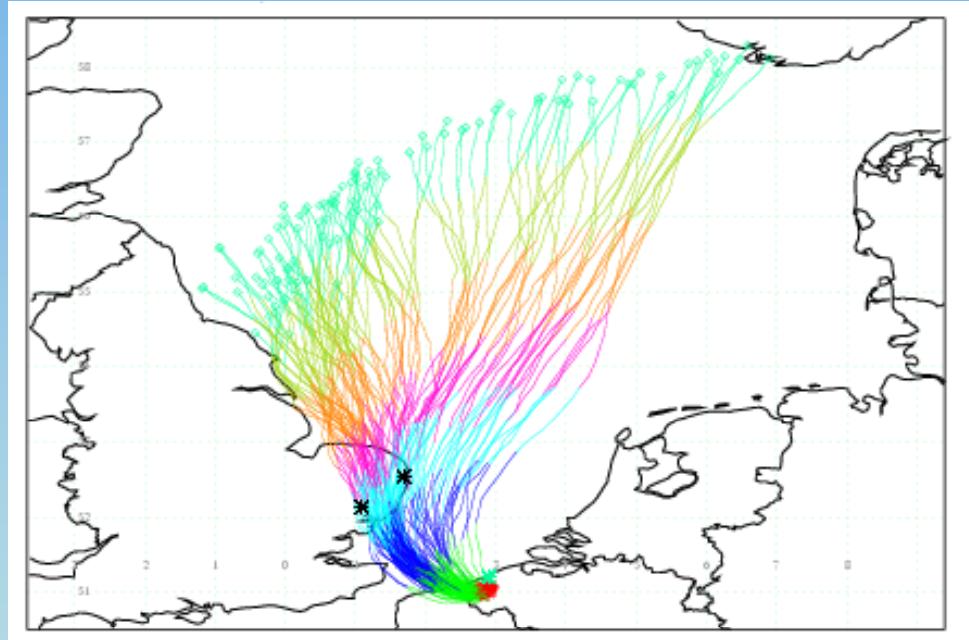




# BTV in the UK during 2007-9



# First Incursion of BTV-8 to the UK (2007)

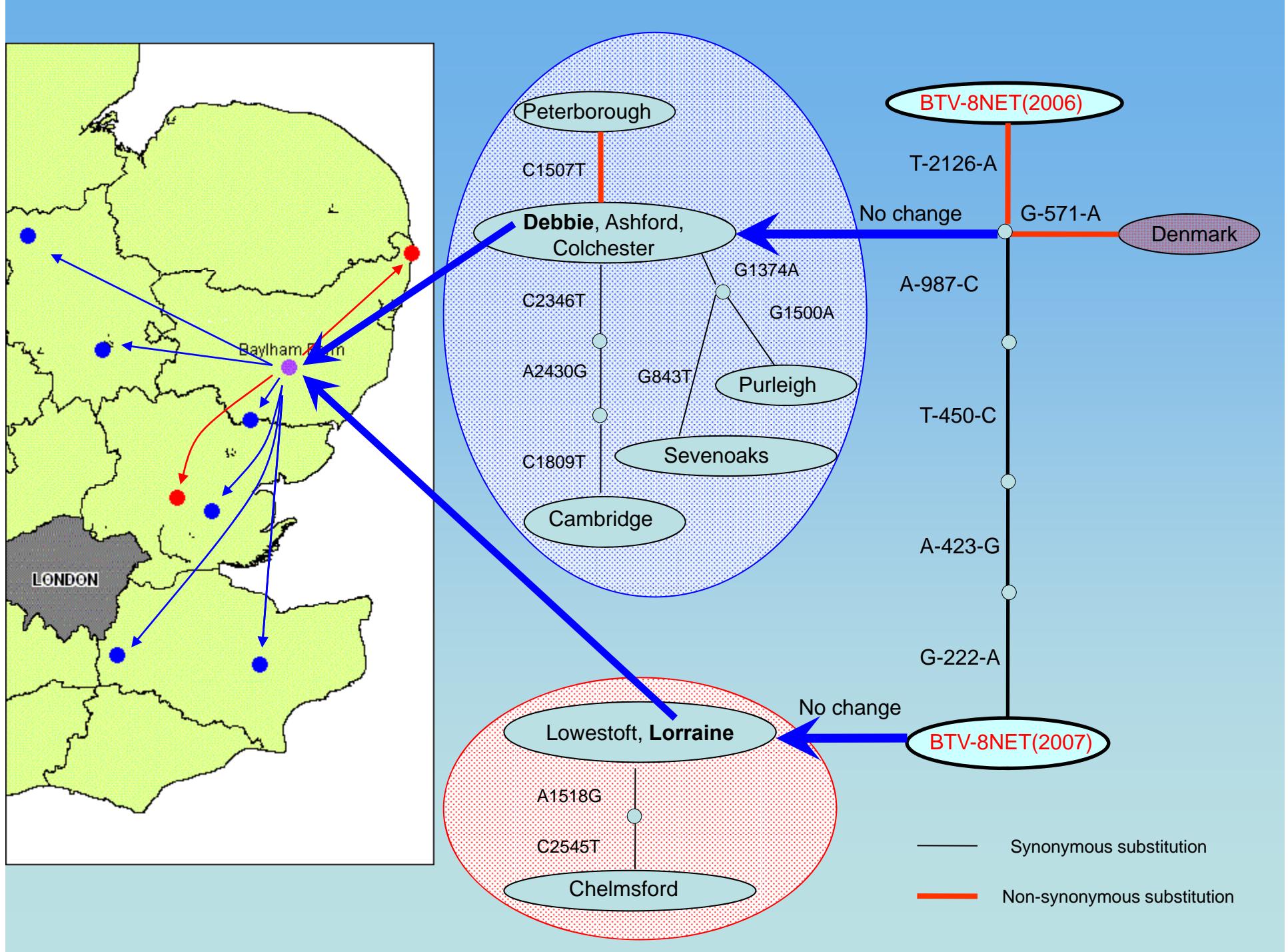


Air concentration plume and particle trajectories overnight on 4/5 Aug 2007 are believed to have caused the first outbreaks of BTV-8 in Suffolk

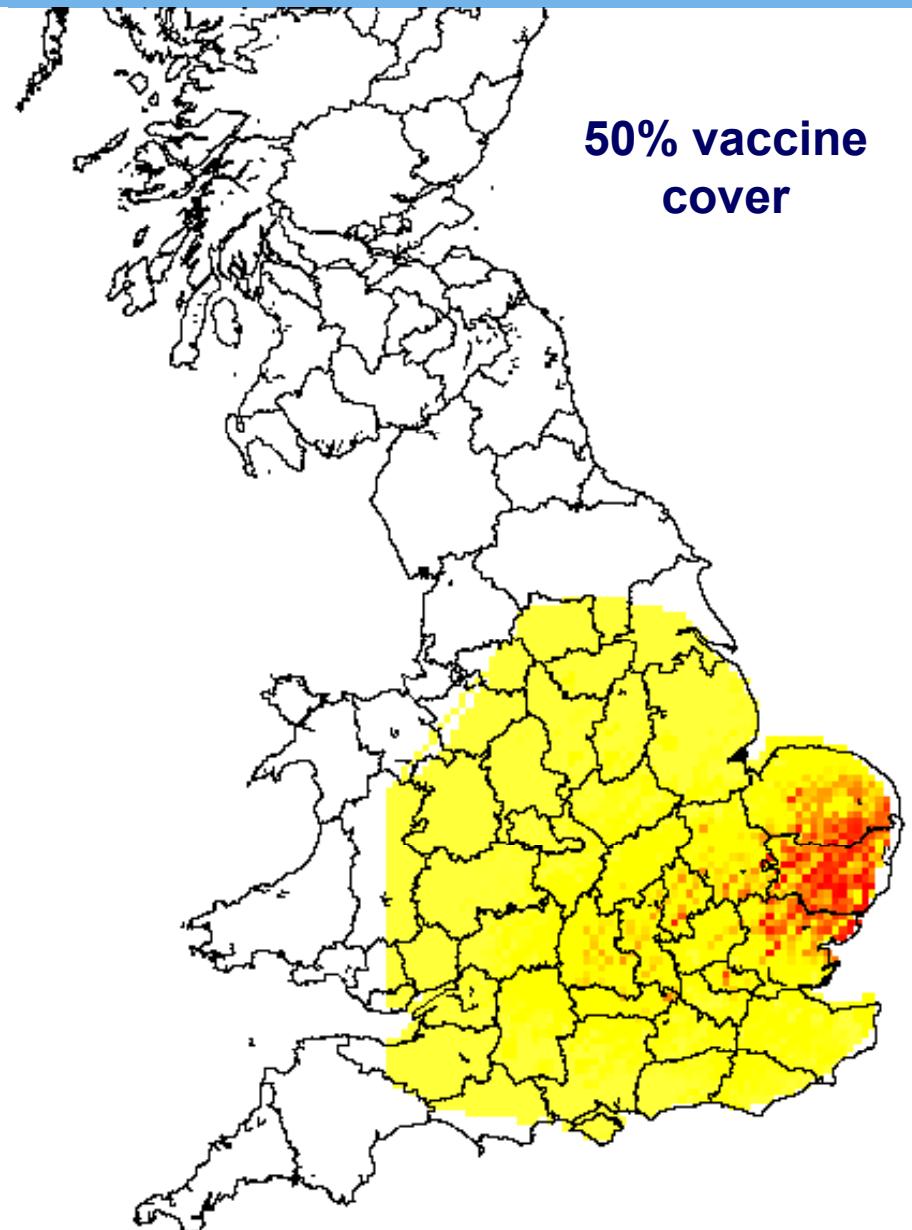
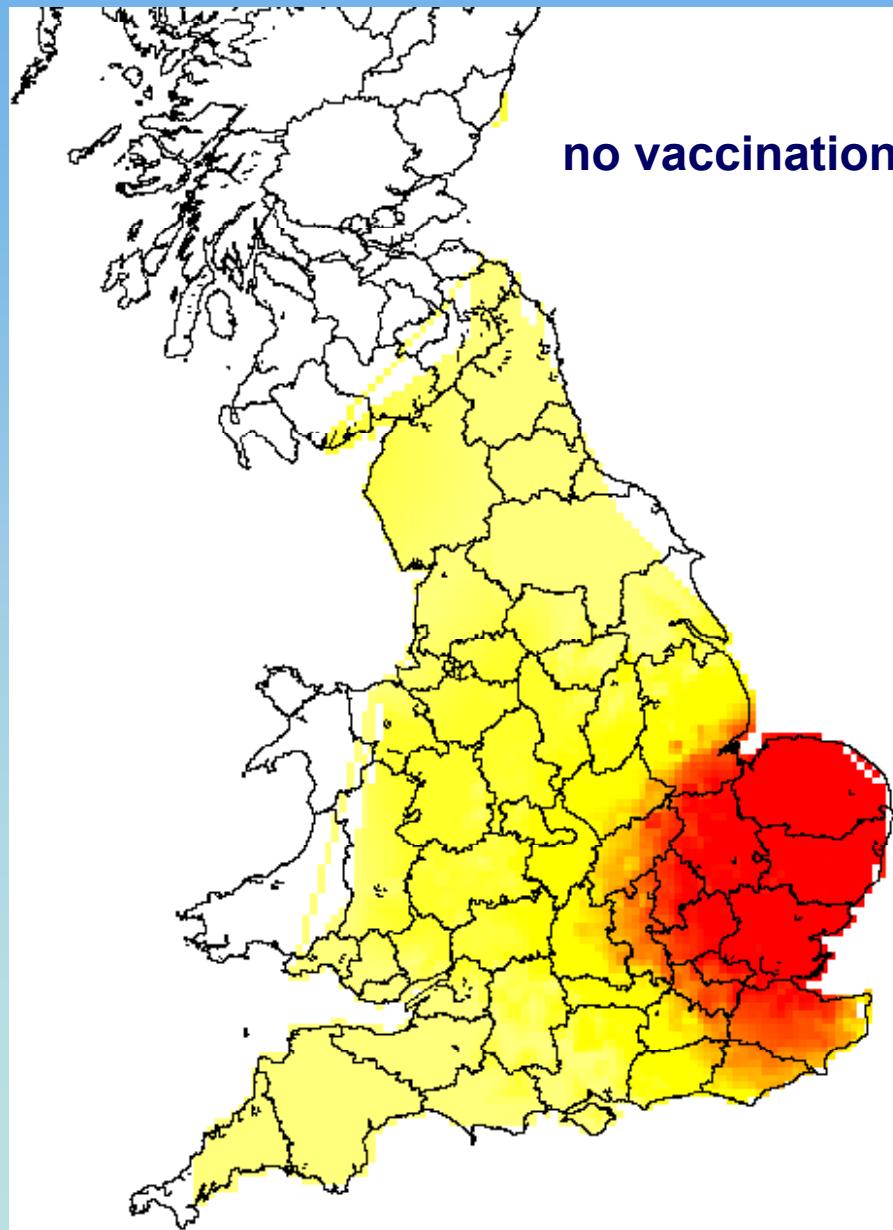
# BTV-8 Protection and Surveillance zones

(1st Jan 2008)





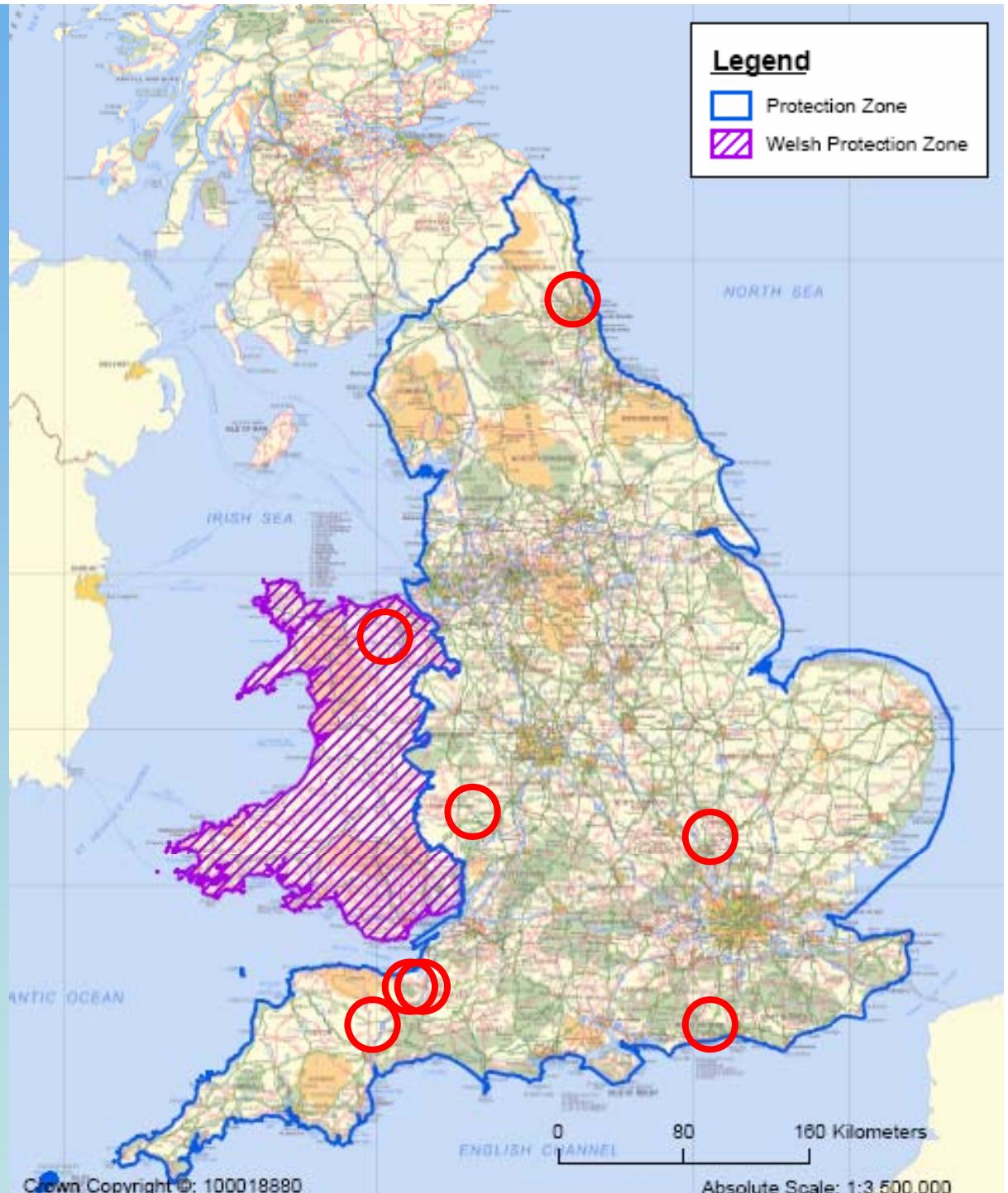
# October 2008: Modelling post vaccination

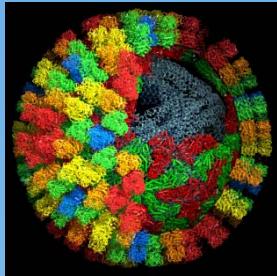


# Sales of BTV-8 vaccines in the UK (to August 2008)

- Protection zone:
  - Total required = 11,914,834
  - Total sold = 9,853,140 (= 82.7% cover)
- Outside the PZ
  - Total Required = >>10,000.000
  - Total sold (so far) = 1,230,000

**Zones as of  
Monday  
20<sup>th</sup> October  
2008:  
No insect  
transmission  
but  
importation of  
196 viraemic  
animals**

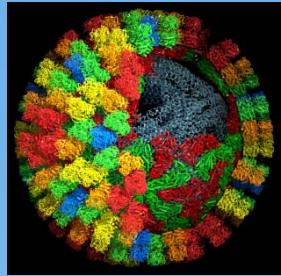




# No further cases of Insect transmitted BTV were observed after the intensive vaccination campaign against BTV-8 in the Summer of 2008

- The UK has now been reclassified a low risk zone.
- However, Vaccination in the UK has dropped to below 10%.
- It is also dropping very rapidly in the rest of northern Europe.
- **As the seropositive animals are replaced, the risk of further outbreaks will inevitably return**





# New BTV serotypes

## 2008-2010

**BTV-25 Switzerland**  
**BTV-26 Kuwait**



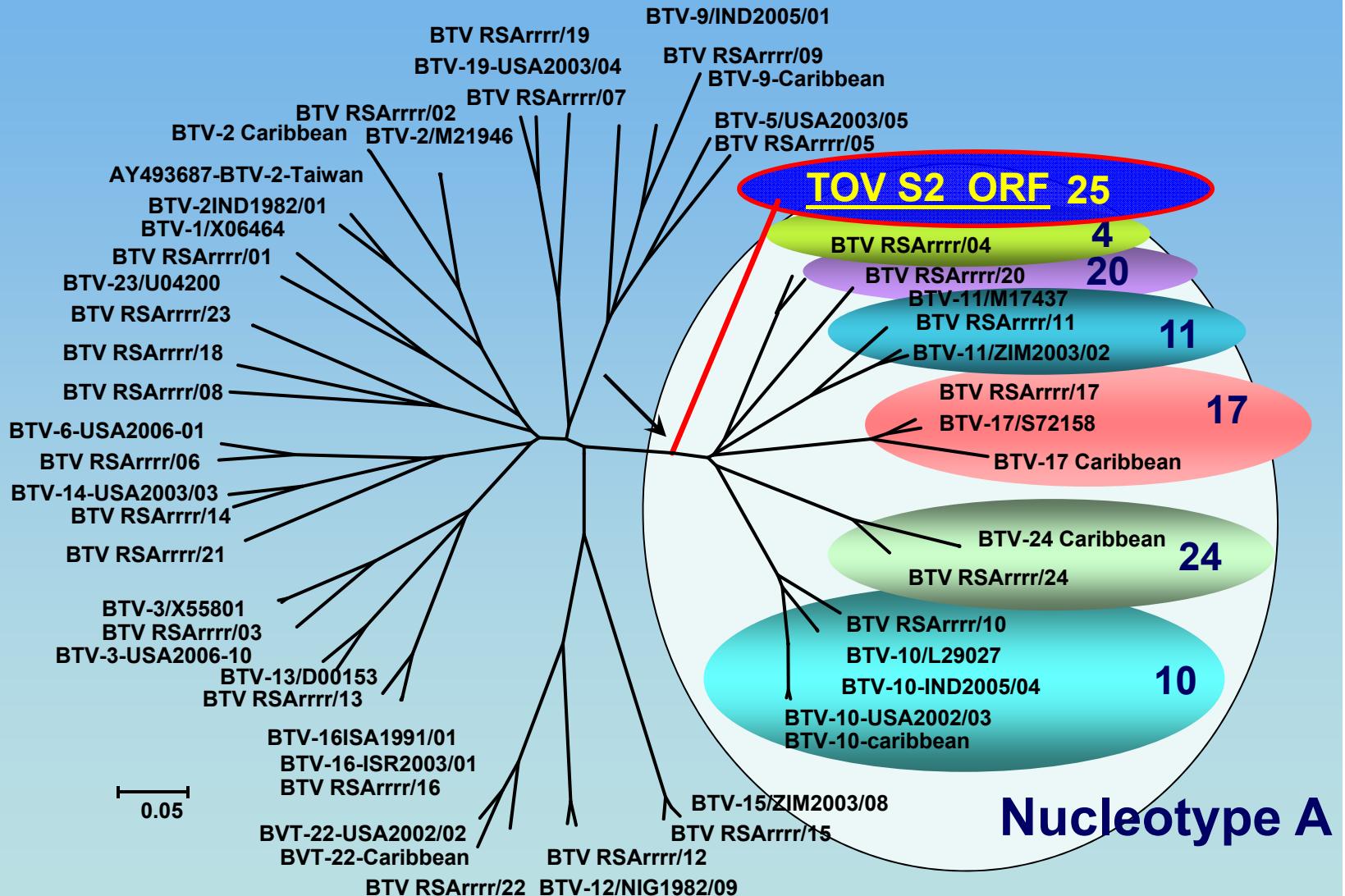
# BTV -25 2008

Toggenburg orbivirus, Cases in goats in Switzerland

No  
clinical  
signs of  
infection

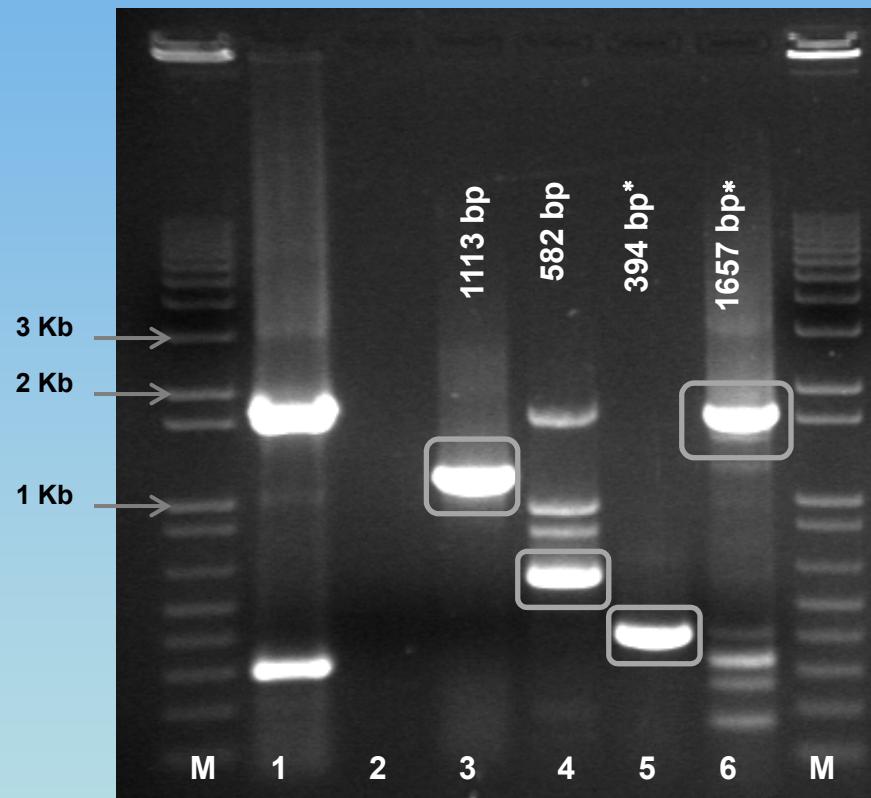


# *Phylogenetic analysis of full length Seg-2 of BTV serotypes*



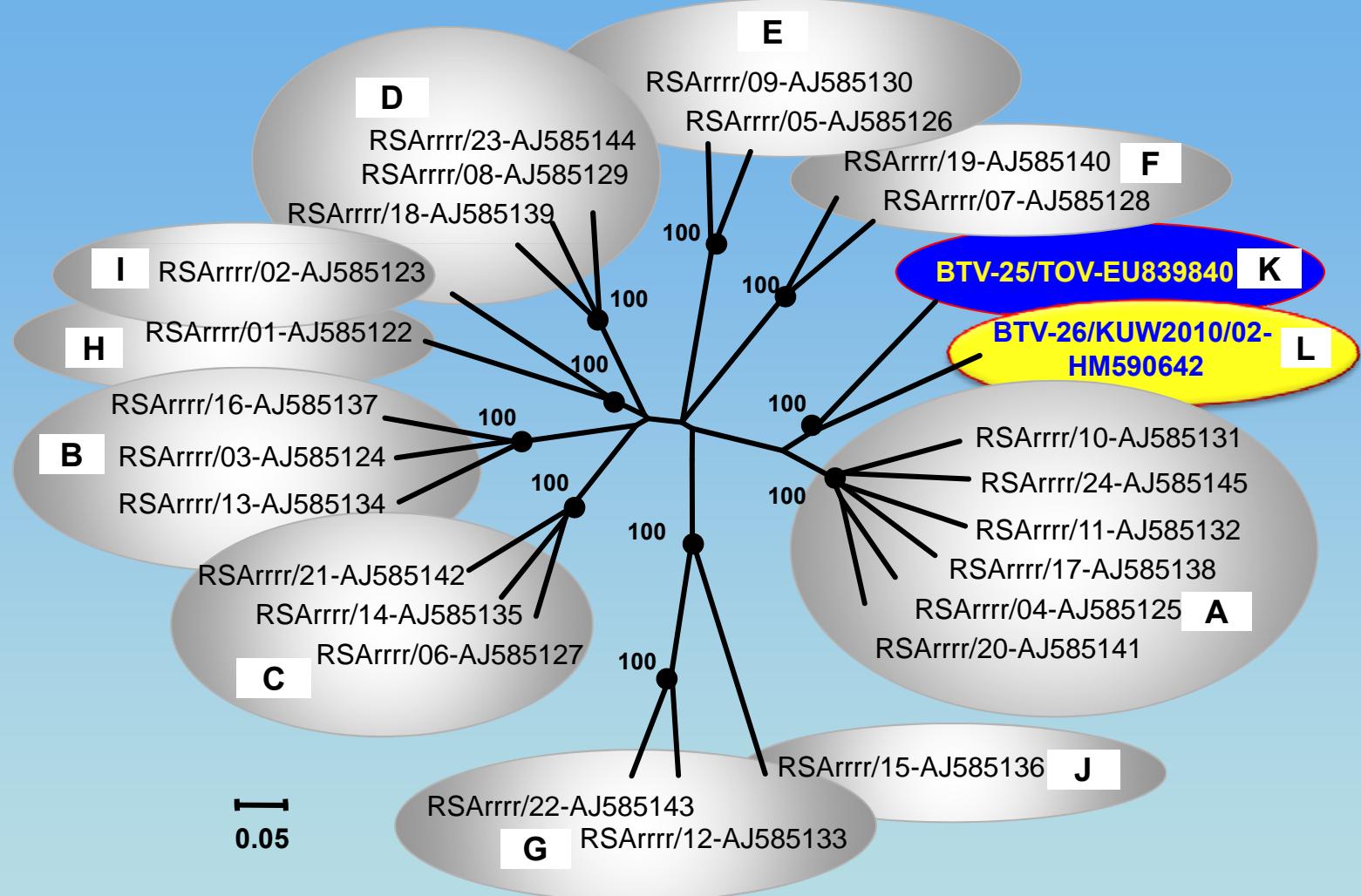
# BTV -26 2010

From sheep in Kuwait 2010

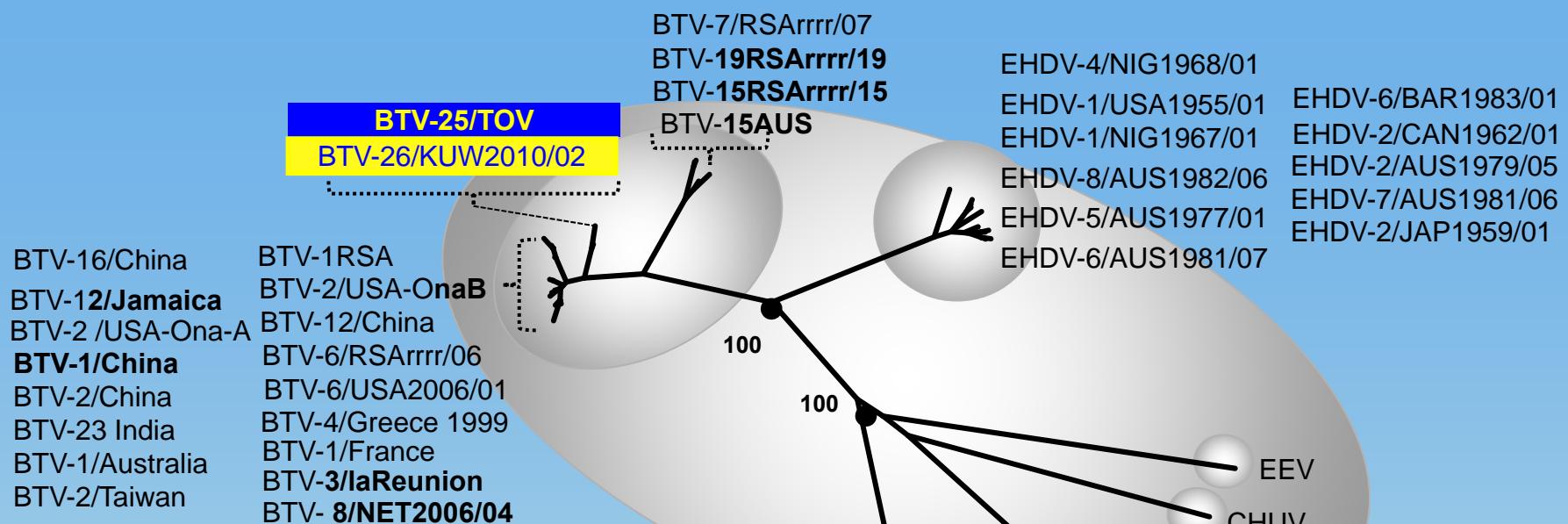


**Electrophoretic analysis of cDNA products from Seg-2 of KUW2010/02 using BTV-26 specific Seg-primer-pairs.**

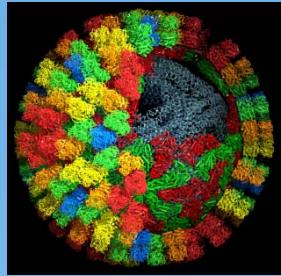
Maan et al., Identification of a novel bluetongue virus serotype (BTV-26) from Kuwait  
(Emerging Infectious Diseases – submitted 2010)



**Neighbour-joining tree showing relationships between VP2 of KUW2010/02 with BTV 1-25.**



**Neighbour-joining tree  
showing relationships  
between VP7[T13] of  
KUW2010/02 with other  
orbiviruses.**



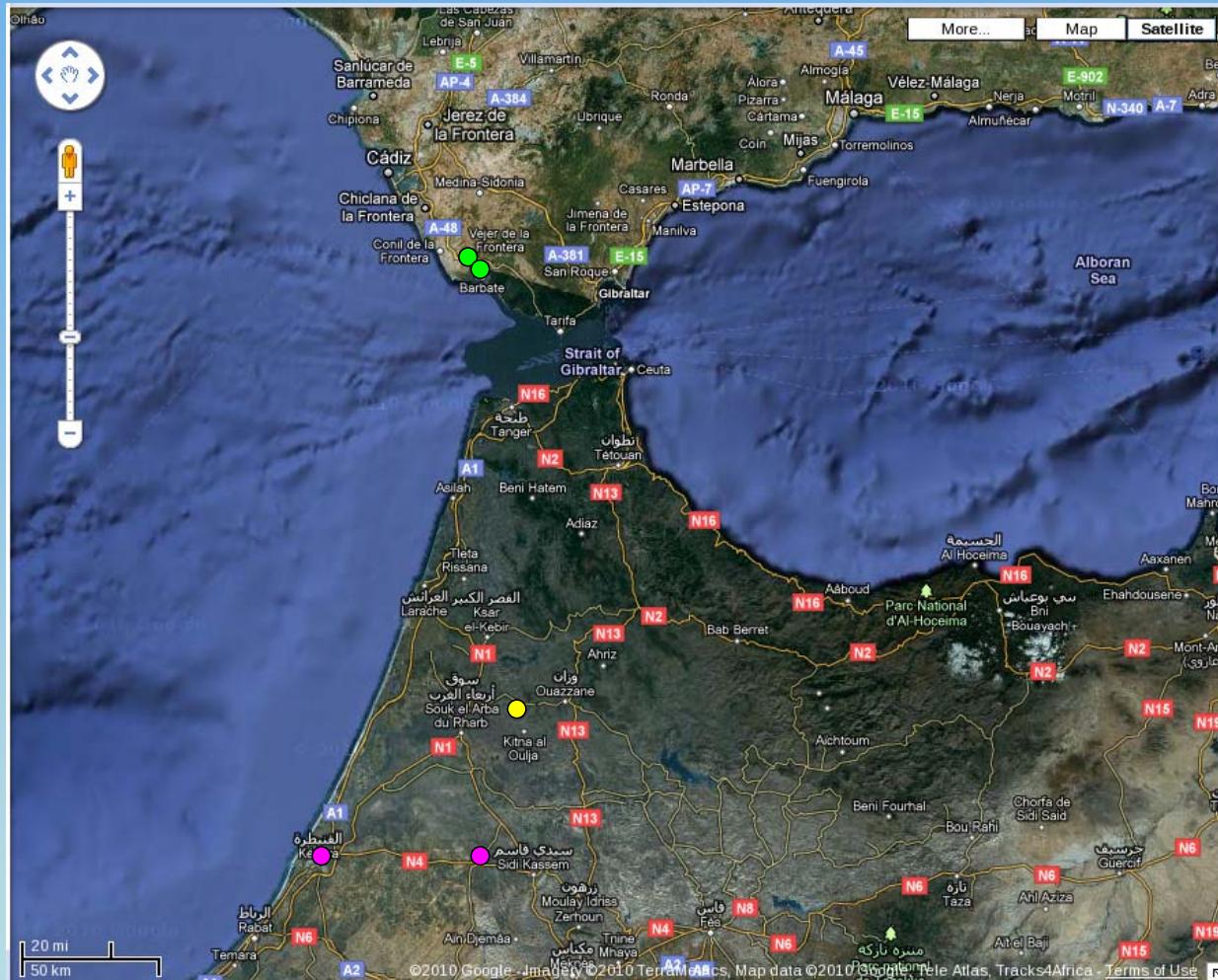
# New Virus strains and from “natural gene technology”

BTV-4

in Morocco and Spain (2010)

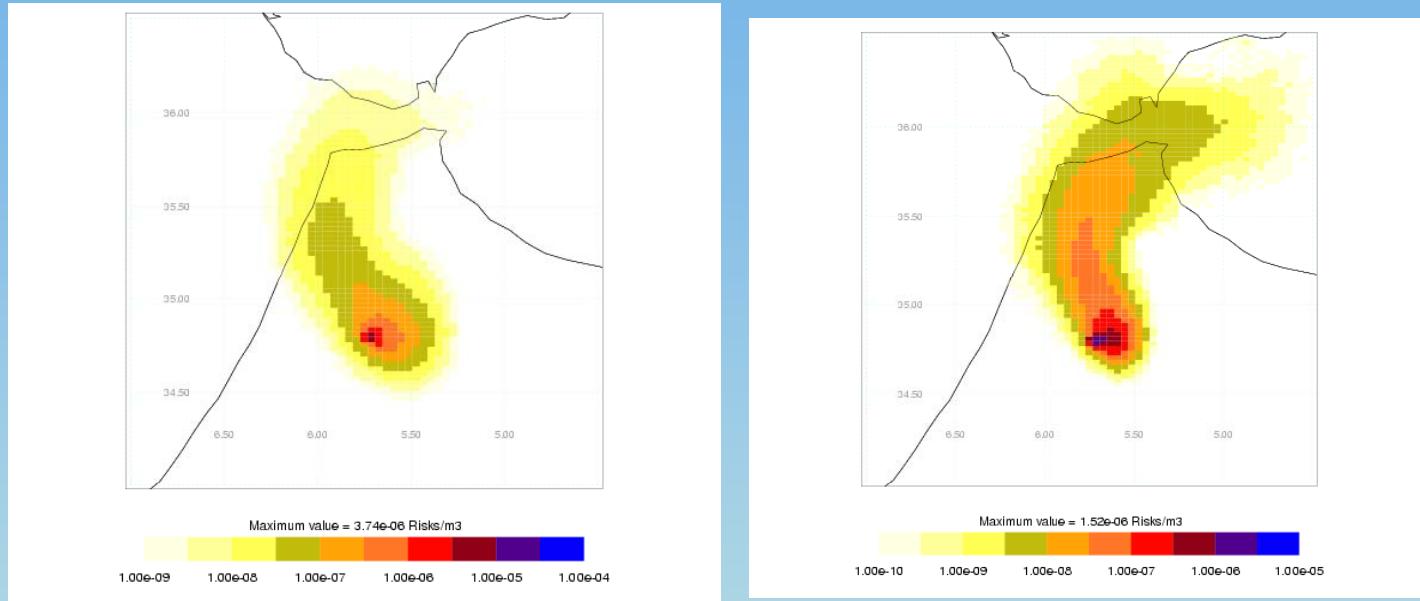


# BTV-4 in Spain and Morocco (2010)



**Fig 1: Locations of BTV-4 outbreaks in Spain (green dots) and Morocco (pink and yellow dots). The release location for the forward NAME runs is identified by the yellow dot.**

# Wind-borne spread of BTV-4 to Spain



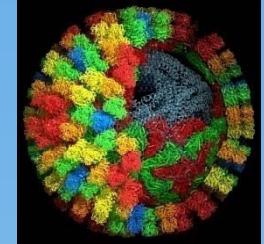
**From Climate modelling (NAME) BTV-4 could have spread from Morocco into Spain on the following dates: 20-21<sup>st</sup>, 29-30<sup>th</sup>, 15-16<sup>th</sup>, or 16-17th August 2010**

## Similarity between BTV-4 2009 (MOR 2009/09) and previous strains of BTV 1 and 4 from Morocco

BTV-4 MOR 2004/02 % similarity with MOR 2009-09	BTV-4 Morocco 2009 ( MOR 2009-09 ) Genome segment Number	BTV-1 MOR 2006/06 % similarity with MOR 2009-09
98.8%	1 ←	99.8%
99.6%	2 →	49.9%
99.4%	3 →	94.90%
95.9%	4 ←	99.8%
96.8%	5 ←	100%
99.8%	6 →	69.2%
94.1% (4MOR2009/10)	7 ←	100%
-	8	-
-	9	-
-	10	-



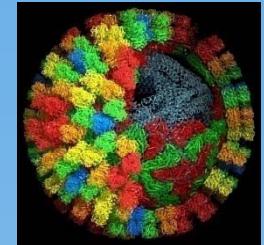
# Summary



- New European BTV strains every year since 1998 . **What will arrive next?**
- Multiple serotypes (e.g. BTV-1, 4, 6, 8 and 11) make detection and control (vaccination) more difficult .
- The whole of Europe is now ‘at risk’ from further incursions of BTV, (new serotypes)
- Genetic exchange (reassortment) can lead to novel virus strains (which may be more dangerous)
- New viruses will emerge, even within well characterised genera and species (BTV-25 and 26).
- New cross-reactive vaccines are needed
- Europe is under threat from other orbiviruses (e.g. AHSV and EHDV), and other arboviruses Recent outbreaks of West Nile, Chicungunya,



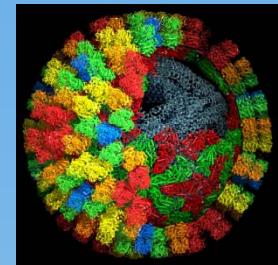
# Biosafety Considerations



- What containment level is appropriate for work on BTV
  - transfer of reagents from foreign animal disease labs to regional diagnostic labs
  - transfer of exotic virus strains to low containment labs
- Vector-borne transmission is most relevant in the field but not exclusive
  - oral transmission has been observed in the field



# Acknowledgements



I would like to thank

- Members of the Arbovirology Programme, at IAH Pirbright
- Many many International Colleagues who have provided virus samples and data.
- Funding support from DEFRA, BBSRC, OIE and the European Commission (OrbiVac - Grant no.: 245266; WildTech - Grant no.: 222633-2) EMIDA grant OrbiNet - K1303206.

