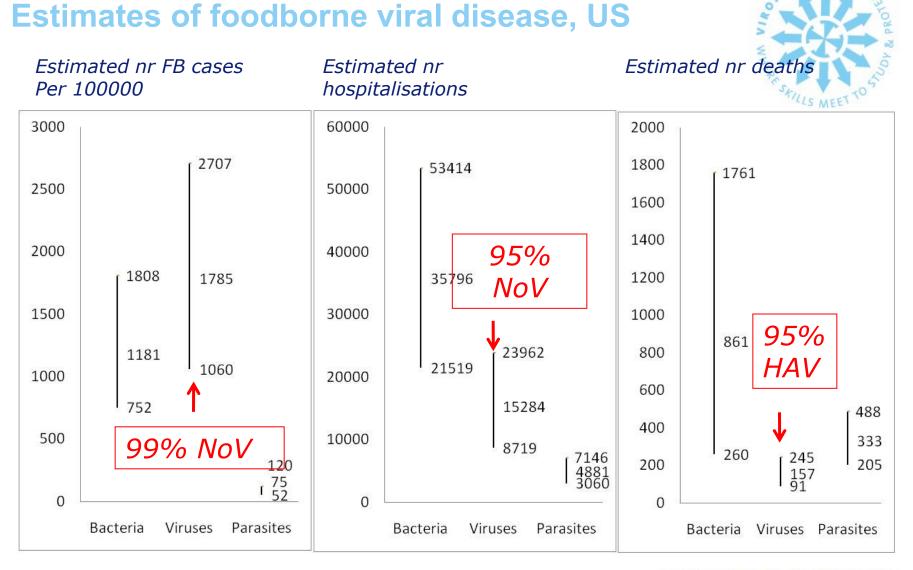


Challenges and opportunities in risk assessment for viruses Marion Koopmans @MArionKoopmans



Scallan et al., 2011

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Viruses and food, sources



Category	Source	Mode of Tx	Example	Clinical syndrome
Fecal viruses human	Fecal contamination at source irrigation Food handler	Fecal oral	Norovirus Sapovirus Astrovirus Rotavirus Enterovirus Hepatitis A virus Hepatitis E virus?	Diarrheal disease Hepatitis Neurological disease
Fecal viruses animal	Fecal contamination at source Irrigation (shellfish, berries)	Fecal oral, zoonotic	Hepatitis E virus Rotavirus 	Hepatitis Diarrheal disease
Systemic infection animal	Meat, animal products	Zoonotic, including fecal oral	Hepatitis E virus TBE Nipah Avian influenza Rift valley fever MERS 	Hepatitis Neurological disease Respiratory disease Hemorrhagic fever

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Issues to address in microbial risk assessment

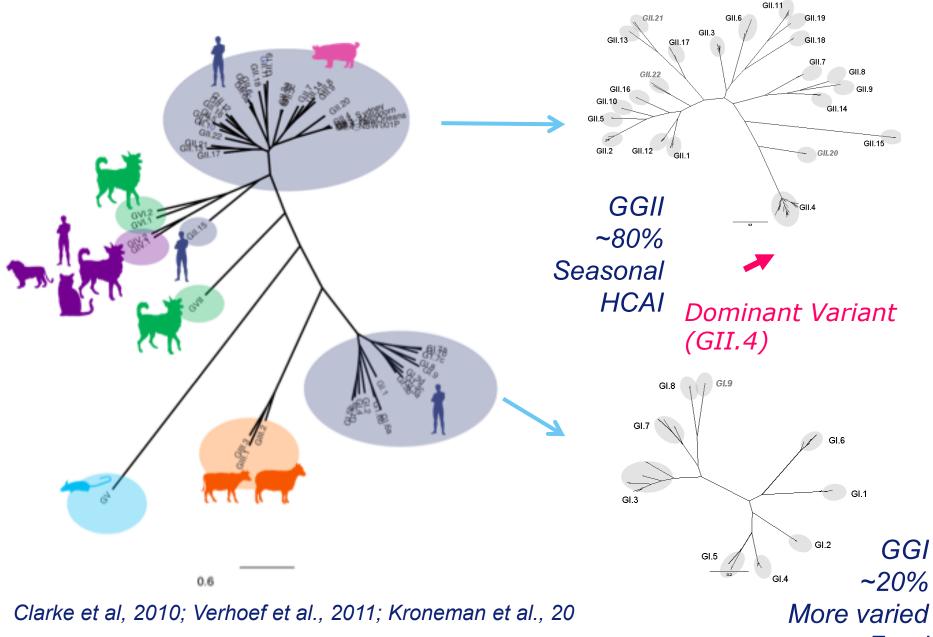
- Microbial growth and death
- Genetic diversity and evolution
- Potential for secondary transmission
- Immunity
- Diversity of health endpoints
- Heterogenous distribution in environment
- Detection method sensitivity
- Population, community, and ecosystem level dynamics
- Routes of exposure
- • • •
- Infectome, microbiome and exposome

Microbial Risk Assessment Guideline: Pathogenic Organisms with Focus on Food and Water. FSIS Publication No. USDA/FSIS/2012-001; EPA Publication No. EPA/100/J12/001.

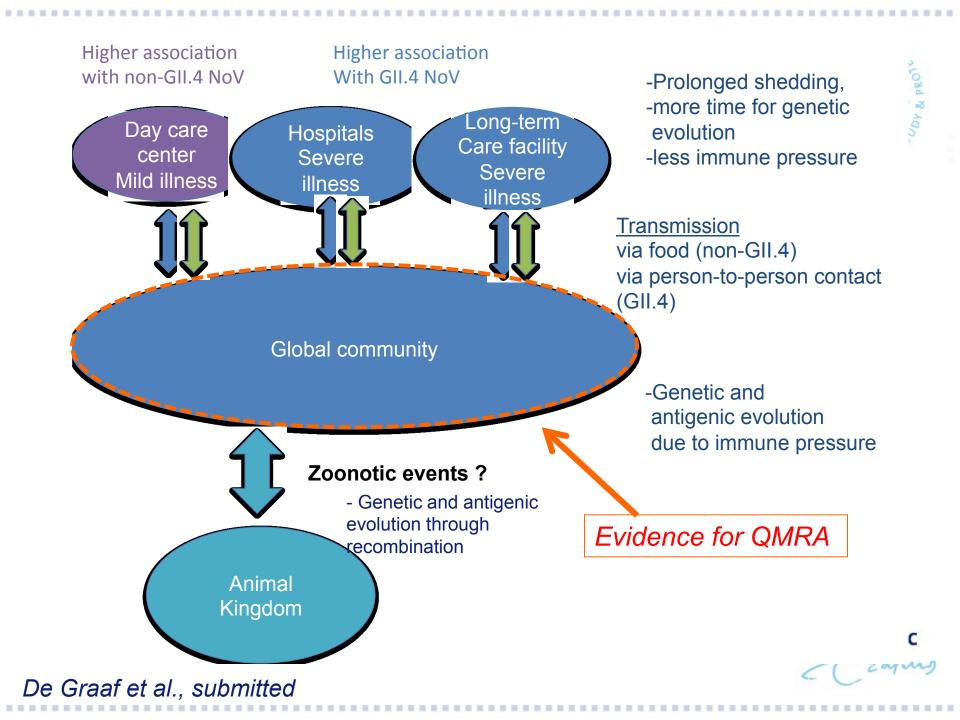


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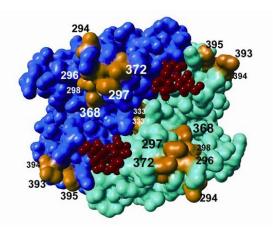
Family, genus, genogroups, genotypes, variants



Food



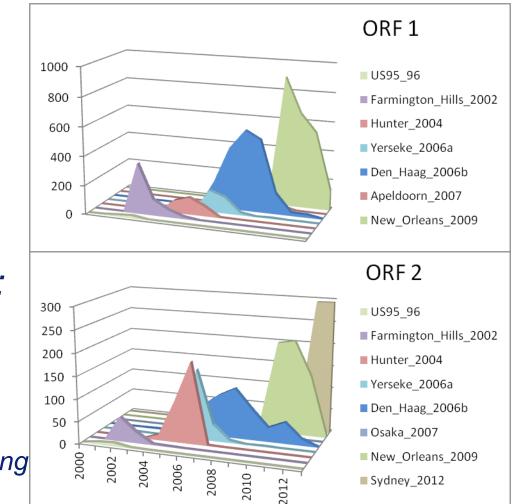
Noroviruses persist in the population through evolution



Effects of mutations:

Escape mutants (drift)No protective immunity

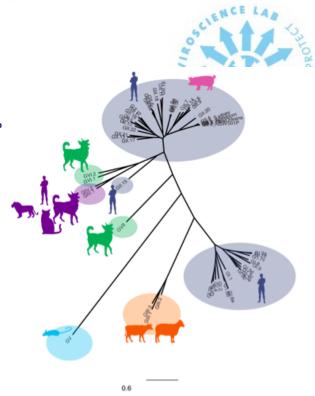
Differences in host cell binding > New host range

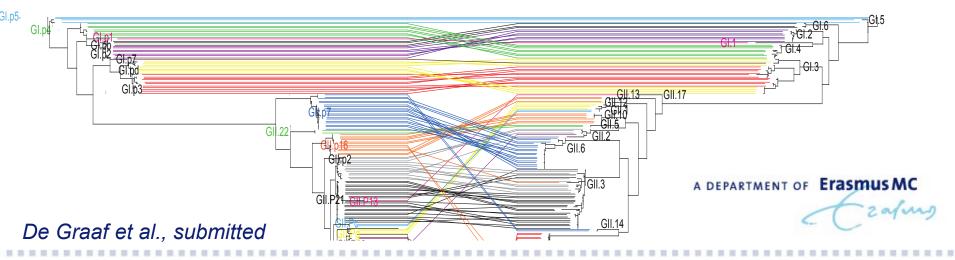


Siebenga et al., 2008; Allen et al., 2008; Parra et al., 2012; Tan et al., 2003 Lindesmith et al., 2008; Bok et al., 2009; Siebenga et al. 2010

Reassortment

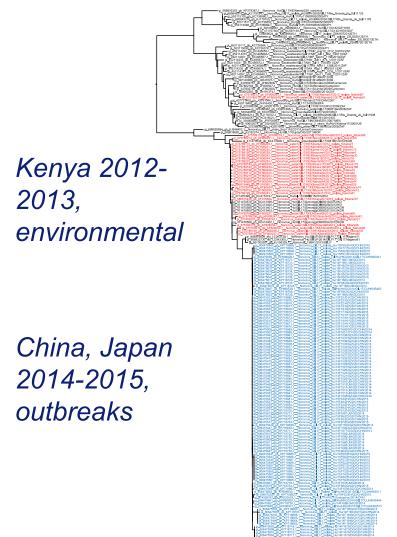
- Infection of a single cell with more than one strain > progeny virus with mixed genome
- More likely when humans are exposed to multiple viruses
 - Eg foodborne outbreaks
 - Daycare centres
 - Immunocompromised persons
- Potential for introduction of novel viruses into the population (mixed genomes)



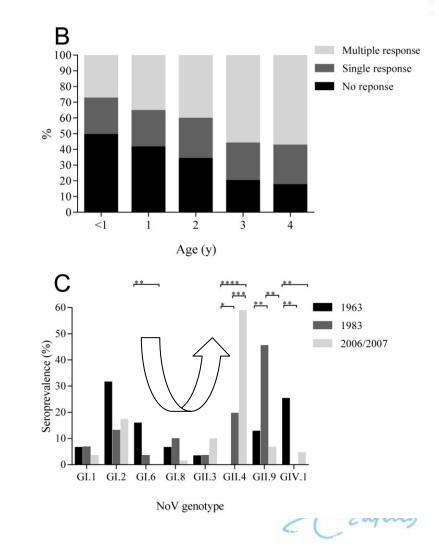


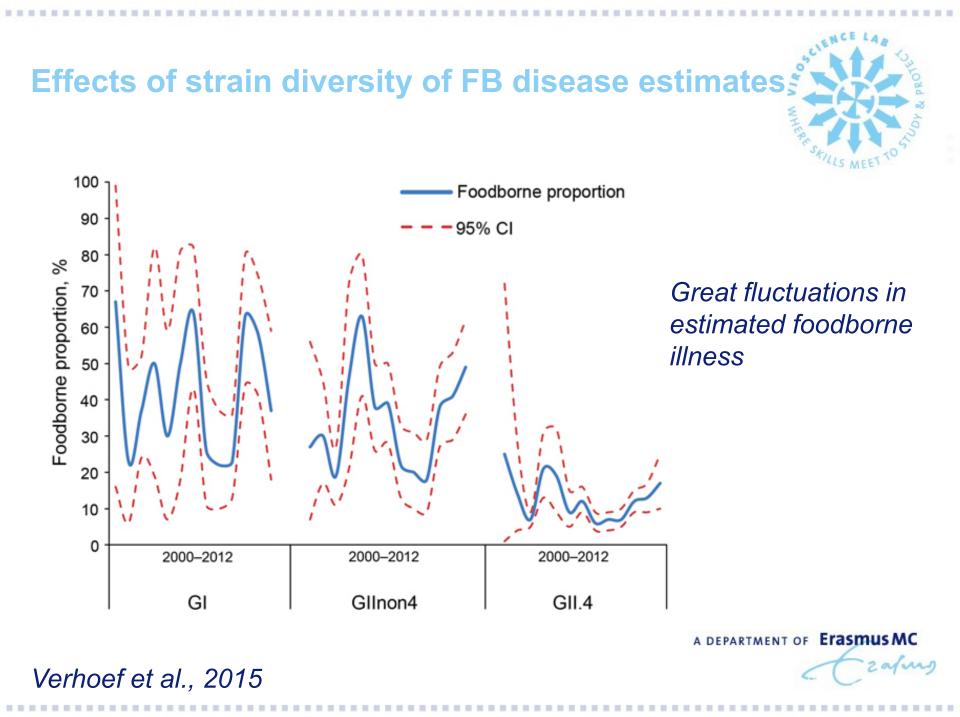
Emergence of recombinant strains may change impact

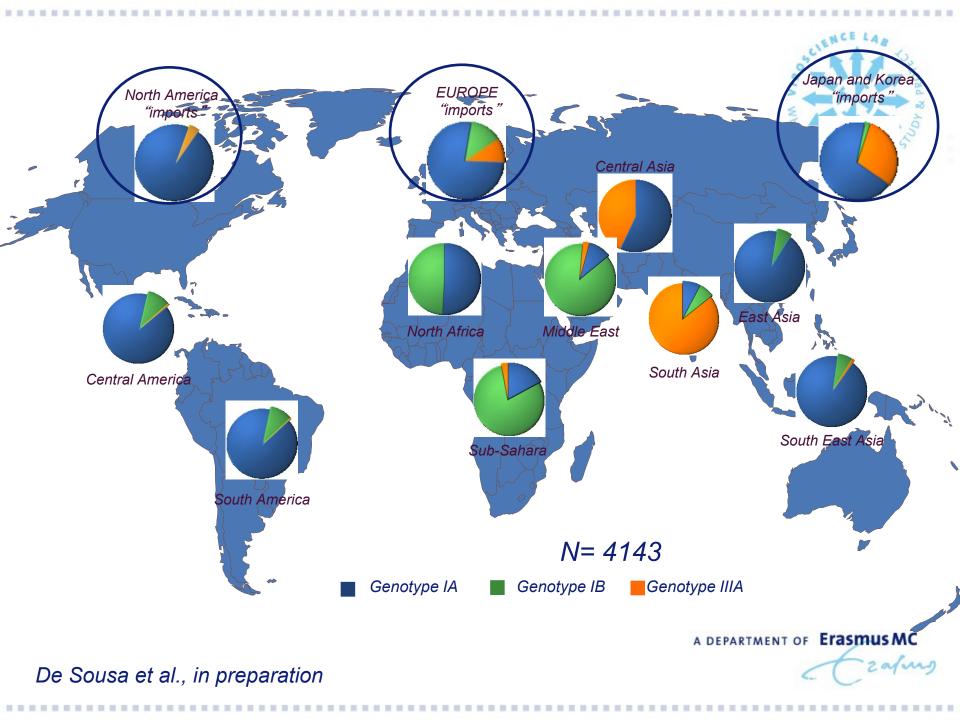




De Graaf et al., 2015; van Beek et al., submitted

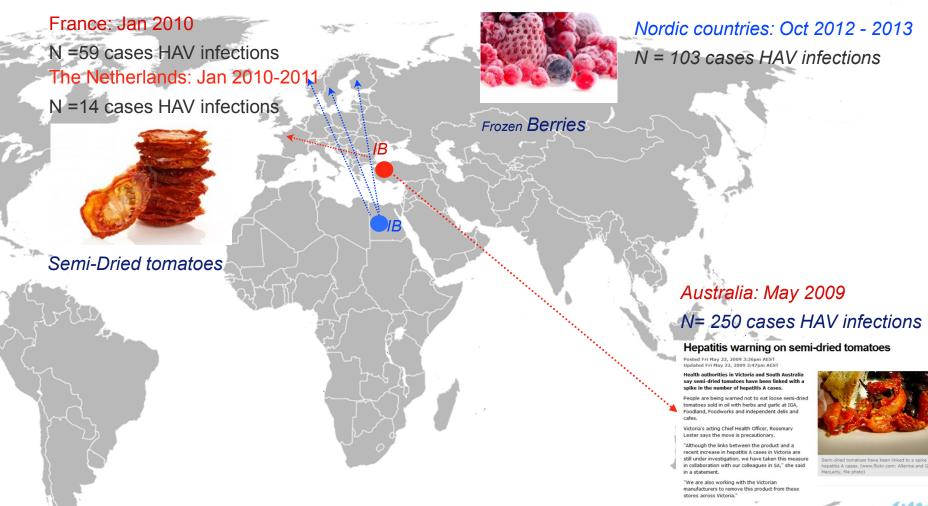






Sequence based cluster detection, HAV example





Petrignani et al., 2010, 2014

Hurdle: getting robust evidence for foodborne outbreaks

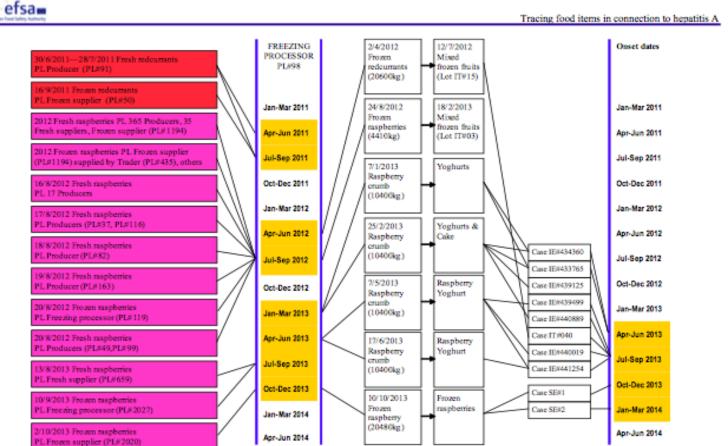


Figure 24: Time line of deliveries from "hotspot" PL#98 to lots and cases in Italy, Ireland and Sweden (only confirmed cases are represented)

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EFSA Journal 2014;12(9):3821



Intensive farming under scrutiny as French oysters face viral wipeout

From Hugh Schofield in La-Trinité-sur-Mer, Brittany

6 Jun 2010

French oyster farmers are facing ruin after a viral epidemic – which many believe is linked to intensive farming methods – struck for the third year in a row, wiping out millions of the baby shellfish.

From the Mediterranean coast, up to the Bay of Arcachon on the Atlantic and now Brittany, farmers have watched in dismay in recent weeks as the virus once again moved northwards, keeping pace with the rising sea temperature.

In 2008 and 2009 the industry was ravaged by the same epidemic, with many farms losing 80% to 100% of their stocks of naissains – first-year baby oysters.

Because it takes three years to grow a commercially viable oyster, so far the economic impact of the crisis has been limited.

But now all pre-2008 production has been depleted, so major shortages are predicted next winter when demand peaks around Christmas and New Year. In France that is when 90% of oysters are sold.

The Committee to Save Oyster-Farming – an ad-hoc group set up in answer to the crisis – has warned that 40% of the country's 4,800 mainly family-run businesses could be forced to close, with the loss of thousands of jobs.

66 Unofficially everyone thinks the hatcheries are to blame, even if no-one says so openly 39

Jacques Cadoret

Ifremer – the state marine research agency – described the situation as "one of the worst crises in the history of French oyster-farming".

Last month, hundreds of farmers staged a much-publicised protest in central Paris, dumping lorry-loads of oyster shells on one of the Seine River bridges. Their pressure paid off, because the French government has now promised a €150 million rescue package to tide the industry over the next three years.

Low tide on the sand flats at Le Po - a hamlet near Carnac in southern Brittany - exposes the poches or bags of overers being cultivated by Brung Lemoine. Last week he detected the first signs of the enidemic with many of the

Oysters have been hit by the OsHV-1 virus

Oyster herpes virus

Massive die-out

Market shift

Top producers: China, Japan, Korea

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Emerging viruses and the food chain

- Pig Hepatitis E immunocompromised
- SARS wild animal markets
- Nipah fruits>pigs>people
- Avian influenza- food handlers, blood
- Rift valley fever animal products, slaughter
- Ebola bush meat
- MERS coronavirus: dromedary milk, slaughterers immunocompromised

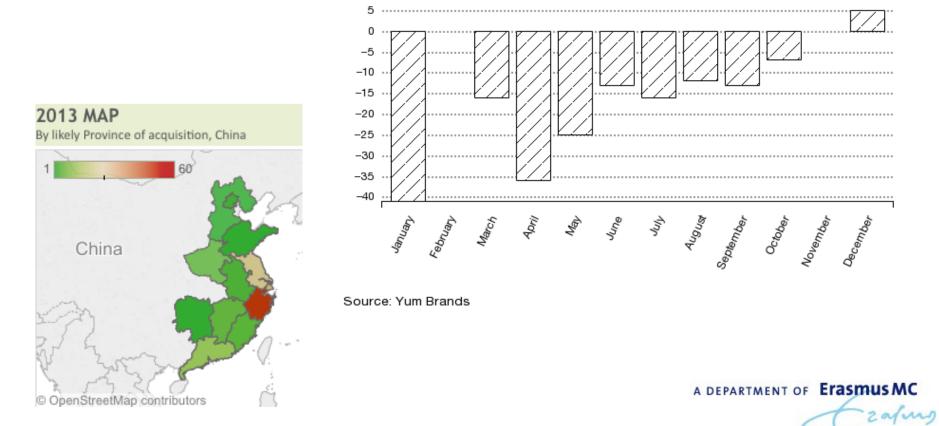


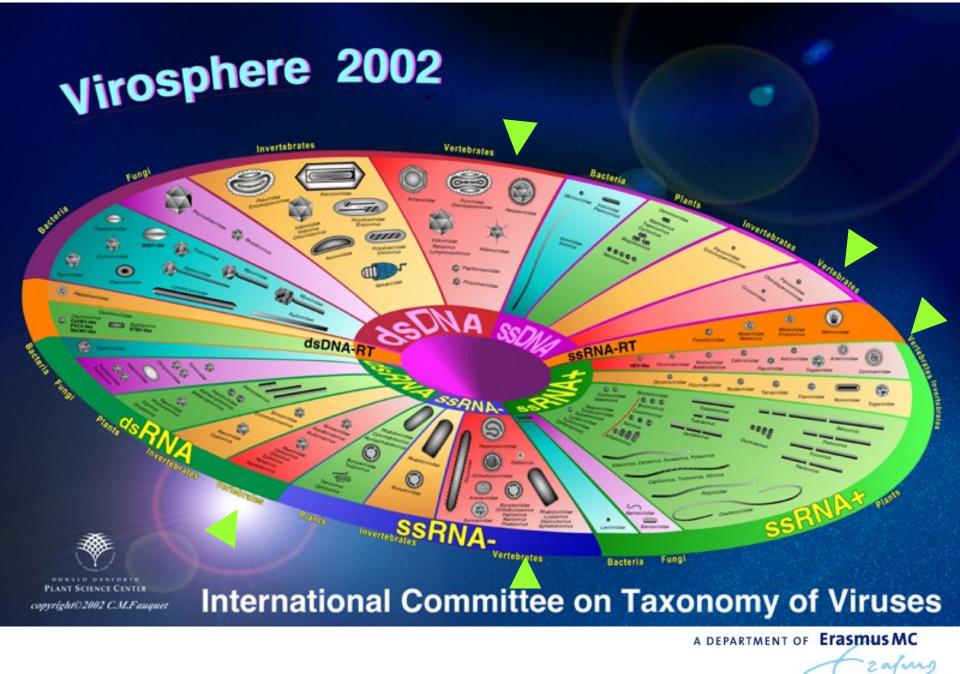


Impact of avian influenza H7N9 emergence



KFC China Same Store Sales (2013) % Change





http://www.ncbi.nlm.nih.gov/ICTVdb/Images/virosphere.htm



COllaborative Management Platform for detection and Analyses of (Re-) emerging and foodborne outbreaks in Europe

A global platform for the sequence-based rapid identification of pathogens - Start date December 1st 2014

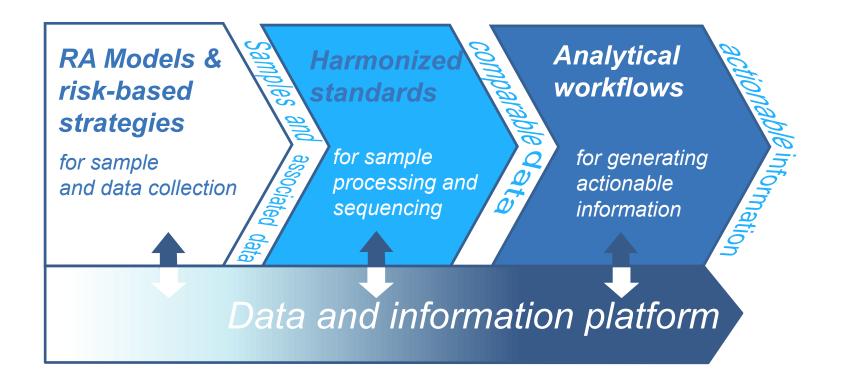
Coordinated by Frank M. Aarestrup (Technical University of Denmark) Marion Koopmans (Erasmus Medical Center, the Netherlands)

www.compare-europe.eu

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Analytical framework and globally linked data and information sharing platform.



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The future of risk assessment science for viruses?



Develop capacity to deal with emerging diseases > agenda setting for research Develop capacity to take stock of evolving data sources



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