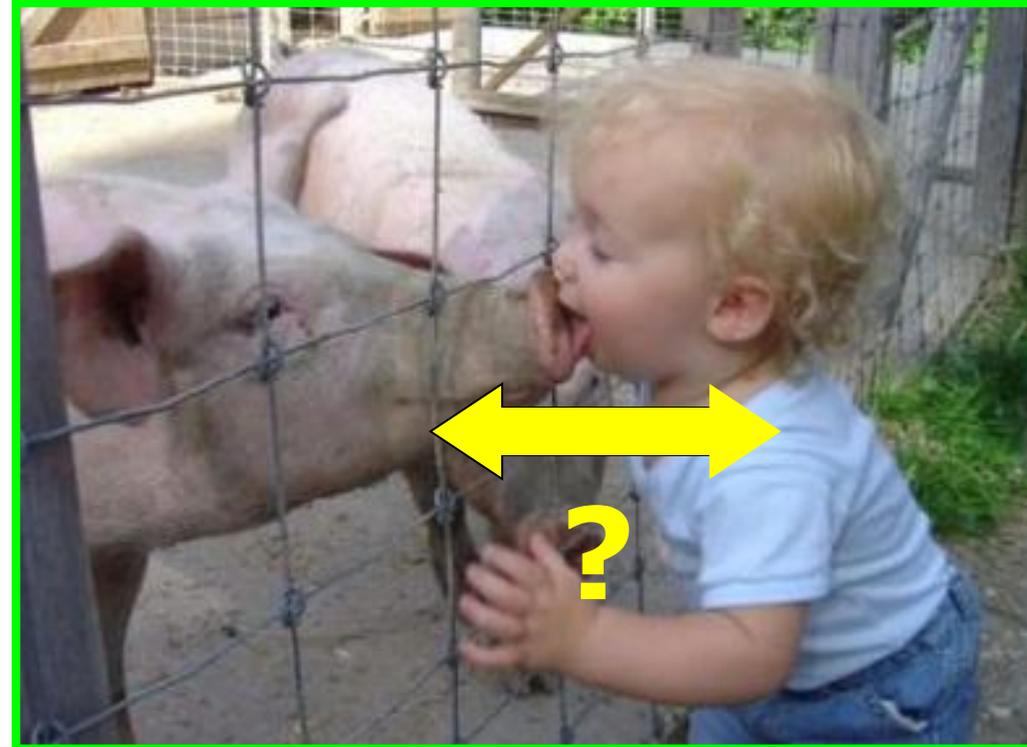
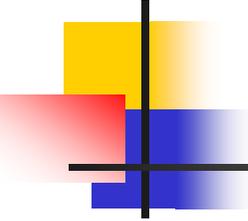


# Livestock Associated MRSA

What is the appropriate  
level of concern?

Peter Davies BVSc, PhD  
University of Minnesota



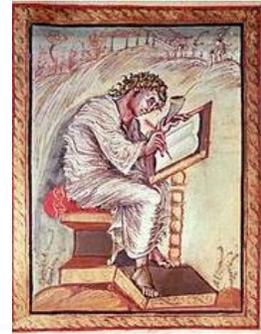


# Questions: LA-MRSA

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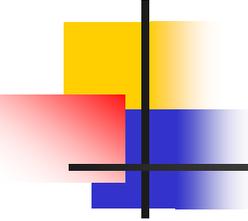
- What are the human health implications?
- Why/how did it emerge?
- What should be done about it?

*Matthew 7:7 - "Seek and ye shall find"*



## MRSA in market hogs (de Neeling et al, 2006)

- National survey of slaughter pigs in Holland
  - 39% of 540 pigs positive (nasal swabs)
  - 81% (44 of 54) of farms positive
- All isolates a 'single clonal group'
  - MLST: ST 398
  - 3 closely related spa types
- Uniformly resistant to tetracycline
  - Use of tetracyclines may be selecting for MRSA?

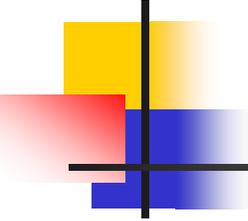


# MRSA in market hogs and swine veterinarians in the USA

(Funded by National Pork Board)

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- MRSA prevalence in swine veterinarians
  - 150 attendees at 2008 AASV meeting
  - 5 of 8 isolates spa type t034
- MRSA prevalence in market hogs
  - 539 hogs from 45 herds (9 plants)
    - All pigs negative at 3 plants
  - 25% prevalence (pigs)
    - Diverse spa types and variability among plants
    - t034 spa type most common
    - Cross-contamination in lairage?



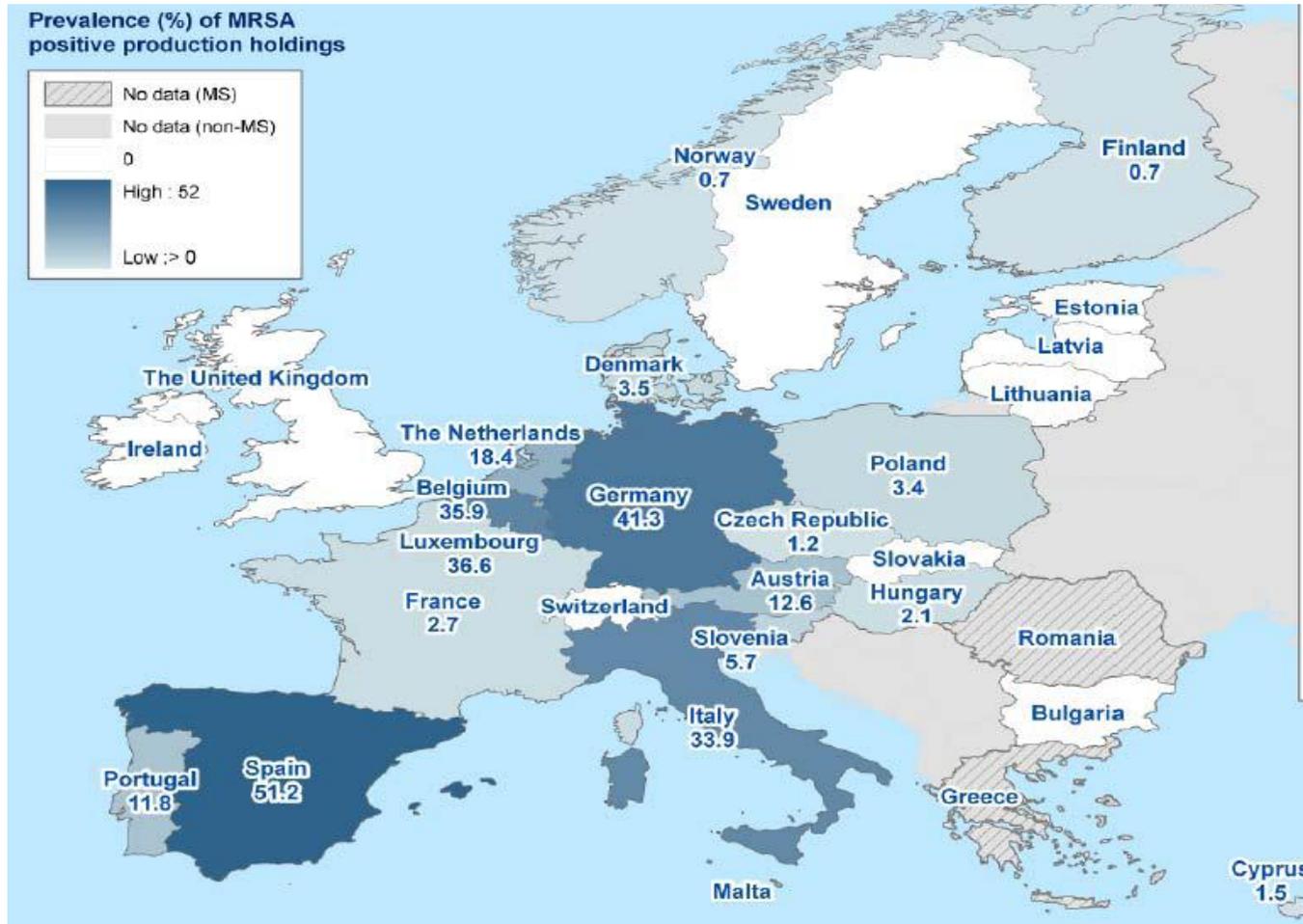
# Multi-state study (IA-MN-OH)

(Funded by National Pork Board)

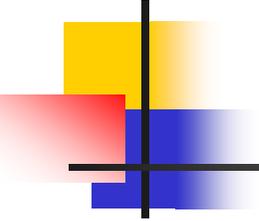
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- 45 farms
  - 18 IA-IL (9 ABF – 9 conventional)
  - 18 MN (9 ABF – 9 conventional)
  - 9 OH-NC (3 ABF – 6 conventional)
- All 21 ABF farms negative
- 4 of 24 (16%) conventional farms positive
- All in IA – why?

# EFSA farm prevalence study (dust samples)



Variability among countries – why?



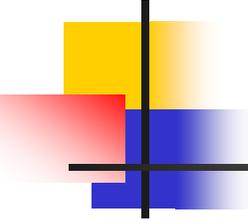
# MRSA from slaughter-age pigs in Canada

Weese et al. BMC Vet Res. 2011 7:41

Province	Herds	Herd prevalence
A	10	0
B	5	0
C	10	30% (3/10)
D	13	7.7% (1/13)
E	8	25% (2/8)

Variability among provinces?

Or uncertainty?



# MSSA isolates in MN

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- 92 isolates of MSSA characterized in MN pigs
  - PCR negative for *mecA* gene
  - Clustered on 4 farms (3 conv., 1 ABF)
- 65 (71%) of pig isolates were spa type t034
  - 1 isolate from human
- Need further characterization
  - Phenotypic and genotypic
  - *mecA* variants?

# What are the risks?

- To whom?
- How?
- How often?
- How severe?

## PROFILE



Maryn  
McKenna

Maryn  
McKenna is

MISINFORMATION

21 JULY 2010

News break: "Pig MRSA" ST398 involved in the death of a child?

OP-ED COLUMNIST

## Our Pigs, Our Food, Our Health



Nicholas D. Kristof/The New York Times

One of the many industrial hog farms outside Camden, Ind.

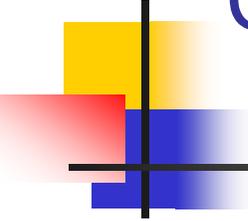
By NICHOLAS D. KRISTOF

Published: March 11, 2009

CAMDEN, Ind.

 TWITTER

 LINKEDIN

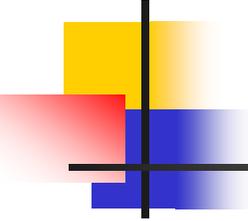


# Community dissemination of MRSA ST398

Cuny et al (2009)

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- Study of German farming community where MRSA ST398 prevalent on pig farms
- Nasal swabs from
  - Pig farmers and family members
  - Swine vets and family members
  - 462 pupils (10 to 16 yo) in villages in the high density pig farming area

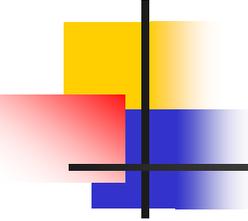


# Nasal colonization with ST 398

Cuny et al (2009)

<b>Group</b>	<b>N</b>	<b>Pct Pos</b>
Pig farmers	113	86%
Farmer family members	116	4.3%
Swine Veterinarians	18	45%
Vet family members	44	9%
Schoolchildren*	462	0.007% (3)

**\*All 3 positive children lived on pig farms**

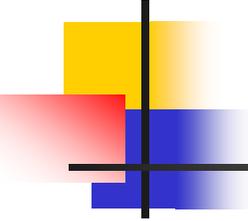


# Communities in pig dense areas in Holland

(van Cleef 2010)

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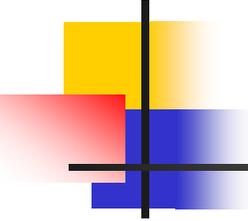
- 3 municipalities in Netherlands where livestock MRSA prevalent
- Adults completed questionnaire and nose swab (n = 583)
  - One of 534 persons without livestock-contact positive (0.2%)
  - 13 of 49 of farm residents/workers positive (26.5%)
- Conclusions:
  - High prevalence of livestock-associated MRSA in people with direct contact with farm animals.
  - Not spread into the wider community



# Occupational exposure!

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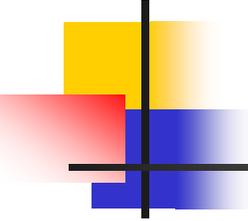
- Consistent observation: occupational exposure to animals increases risk of **MRSA positive culture**
  - Farmers, veterinarians, abattoir workers
- Colonization vs. contamination
  - Duration of 'colonization'
  - Role of regular animal contact
- Consequences of colonization
  - Infection risk
  - Transmission risk



# Duration of colonization

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- Research workers (short term exposure) *van Cleef et al (2011)*
  - 33 of 199 exposures led to positive culture on farm
  - Only 1 of 33 retested positive after 24 hours
- Evidence suggests most short term exposure leads to short term 'colonization'
- Veal farmers in Holland *Graveland et al (2011)*
  - Rapid decline in prevalence during absence of animal contact
  - LA-MRSA poor persistent colonizers in most humans.

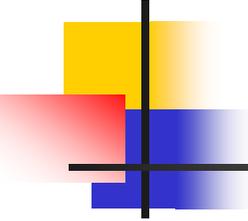


## Transmission of ST398 MRSA among people

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- Studies of transmission in Dutch hospitals (*Bootsma, 2010*)
  - ST398 is 5.9 times less transmissible than non-ST398 MRSA in Dutch hospitals'
  - 'Spreading capacity per admission insufficient to lead to an epidemic'
- 'Nosocomial transmission of ST398 MRSA is 72% less likely than non-ST398 MRSA strains'

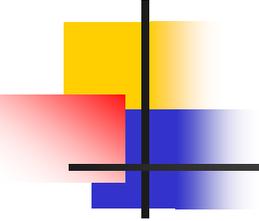
*Wassenberg (2011)*



## Burden of disease from ST398 MRSA

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- Many reports don't distinguish infection from colonization
- Small number of serious infections
  - Often no livestock association
  - One reported fatality with ST398 MSSA without livestock contact
- Retrospective study of human isolates in Canada
  - 5 ST398 out of 3,687 MRSA isolates
  - 4 skin/soft tissue infections (Golding et al 2010)
- CDC has examined >12,000 isolates in USA
  - ST398 not identified in a human clinical case (June 2011)



# Self reported disease in US swine farmers

Leedom Larson et al (2010)

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- Mail survey
  - 135 (17.2%) surveys received from 783 pork producers actively farming hogs
- Five (3.7%) respondents reported a history of physician-diagnosed MRSA SSTI
  - Time period unclear (ever?)
- No bacteriological information or control group

# Distribution of LA-MRSA and other MRSA clinical isolates in Europe

(van Cleef et al., 2011)

Sample source	No. (%) typed clinical isolates		p value†
	MRSA ST398, n = 113	Other MRSA, n = 3,435	
Blood	2 (1.8)	343 (10.0)	<b>0.004</b>
Respiratory tract	20 (17.7)	451 (13.1)	0.16
Skin and wound	76 (67.3)	2,312 (67.3)	0.99
Urinary tract	6 (5.3)	173 (5.0)	0.90
Other	9 (8.0)	156 (4.5)	0.09

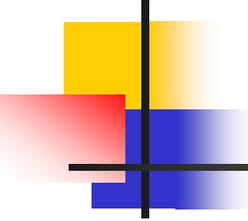
\*Only data from 9 national or regional laboratories in the 7 countries that reported clinical isolates and typed all these isolates were included.

# Geographic distribution of *S. aureus* causing invasive infections in Europe

Grundmann et al 2010



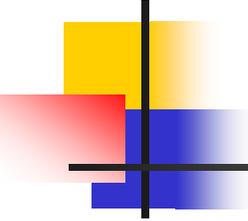
- 357 laboratories serving 450 hospitals in 26 countries (2006-2007)
  - 2,890 MSSA and MRSA isolates from invasive infections
- ST 398 spa types (t011, t034, t571, t1255, and t2383) identified on 12 occasions (1.3%)
  - None harbored the mecA gene.
- No cases of ST398 MRSA invasive disease



# Public health risk of ST398 MRSA

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- Current evidence suggests low transmissibility
- Current evidence suggests low pathogenicity?
  - Lack of reported fatalities (over 7 years)
  - Elevated **infection** risk in farm workers yet to be documented
  - Significantly less invasive disease in Europe
  - No reports of outbreaks



# Lethal pneumonia caused by an ST398 *S. aureus* strain

Rasigade et al (2010)

## ■ Observations

- Fatal necrotizing pneumonia in a previously healthy 14yo girl
- ST 398 - spa type **t571**
- PVL positive
- Tetracycline **susceptible**
- Methicillin **susceptible** (MSSA)
- No livestock contact

## ■ Inference

- “spread of *S. aureus* ST398 among livestock is a matter of increasing concern because strains of this sequence type were able to acquire PVL genes”

# "One Health" means more than one inference

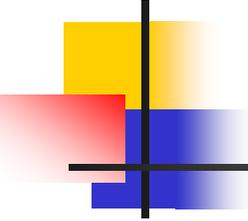


- ‘.. concern because strains ST398 strains were able to acquire PVL genes’ Rasigade et al (2010)
- But....
  - Livestock strains almost uniformly tetracycline resistant and PVL negative
  - Spa type t571 uncommon in animal isolates
- Could adaptation of ST398 to livestock hosts include loss of human virulence factors?
- Could some ST398 variants persist in people without any role of livestock?

# More of the story

Davies et al, EID June 2011

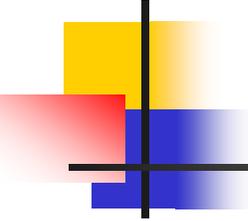
- **t571** ST398 MSSA detected in 9 families from the Dominican Republic living in Manhattan
  - with no apparent contact with livestock (Bhat et al., 2007)
- **t571** the sole MSSA spa type in Dutch study of ST398 clinical isolates, including 3 cases of nosocomial bacteremia
  - with no apparent livestock contact (van Belkum et al., 2008)
- **t571** the predominant (11%) MSSA type at a Beijing hospital
  - Livestock contact unlikely (Chen et al 2010)
- Recent case report of **t571** MSSA from Colombia
  - With no apparent livestock contact (Jimenez et al 2011)



Emergence of unusual bloodstream infections associated with pig-borne-like *Staphylococcus aureus* ST398 in France. (van de Marquet et al., 2011)

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- Study of **t571 MSSA** strains from cases of bloodstream infections in France
- The 30 isolates differed from pig-borne strains
- Isolates shared similarities with strains from humans in China and virulent USA300 strains
- Epidemiologic diversity in ST398 lineage



# Human origin for avian pathogenic

## *Staphylococcus aureus*

(Lowder and Fitzgerald 2010)

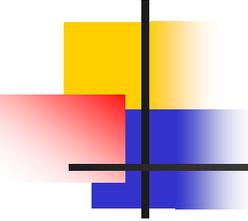
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- Traced origin of *S. aureus* CC5 causing morbidity in the broiler industry.
- Evidence that it evolved from a single human to poultry host jump followed by extensive genetic diversification
  - Acquisition of novel mobile genetic elements
  - Loss of virulence gene function
- “Loss of function of genes involved in human disease suggests that the clone may now be attenuated for virulence in humans”

# Evolutionary genomics of *S. aureus* reveals insights into the origin and molecular basis of ruminant host adaptation

(Guinane et al, 2010)

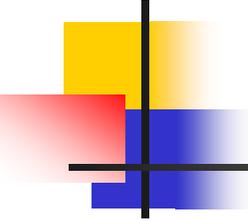
- Evidence that CC113 clone of *S. aureus* resulted from a human to ruminant host jump followed by adaptive genome diversification.
- Whole-genome sequencing
  - Molecular evidence for host adaptation
  - Gene decay and diversification of proteins involved in host-pathogen interactions
- Interspecies transmission leads to genetic adaptation and changes in virulence and transmissibility?



# Diversity of genome of human and LA ST398 MRSA strains

(Hallin et al., 2011)

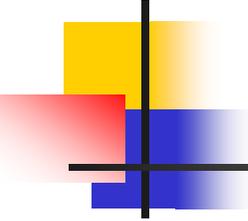
- 'LA-MRSA strains represent a homogenous lineage distinct from co-local HA- and CA-MRSA strains
  - characterized by a lack of human-associated virulence and adhesion determinants
- Absence of detectable enterotoxin gene among ST398 LA-MRSA strains from a wide host range is reassuring regarding their foodborne pathogenic potential.'



# LA-MRSA: Implications

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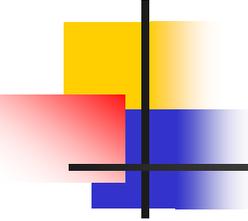
- Not good news for swine/livestock industries
- Priority to understand occupational risks
- Preventive measures in industry
  - Personal hygiene
  - Showers (soaps, towels)
  - Clothing
- Wound treatment, covering
- Awareness and medical treatment



# The role of antimicrobial use?

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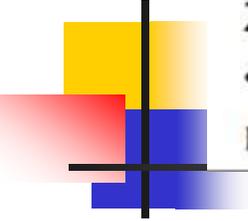
- Tetracycline resistance
  - MRSA emergence in horses
- Long term use of beta lactams in food animals
- 'Therapeutic' vs. 'non-therapeutic' uses
- Role of antimicrobial growth promotants
  - Most are not beta lactams
- Newer injectable products
  - Long acting cephalosporins



## Selective pressures (Aarestrup 2010)

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- Tetracyclines
  - MSSA in pigs also tetracycline resistant
- Zinc
  - 74% of ST398 MRSA in DK had high MIC to zinc
  - All 60 MSSA strains had low MIC to zinc
- Widespread use of zinc since AGP ban
  - Prevention of enteric disease in weaned pigs
  - Used on most DK swine herds
- Law of unintended consequences?



Zinc resistance of *Staphylococcus aureus* of animal origin is strongly associated with methicillin resistance

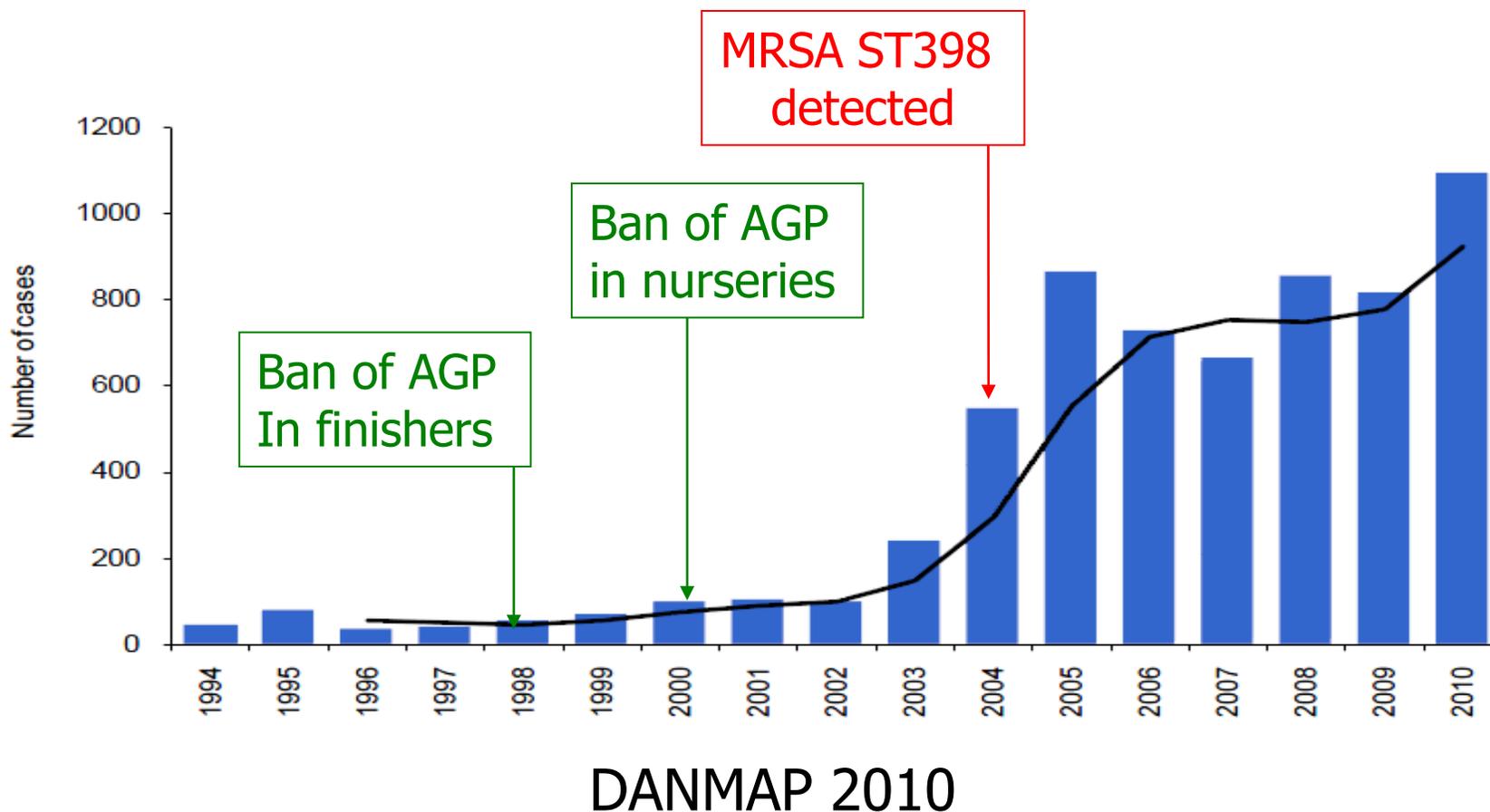
Lina M. Cavaco <sup>1,\*</sup>, Henrik Hasman <sup>1</sup>, Frank M. Aarestrup <sup>1</sup>

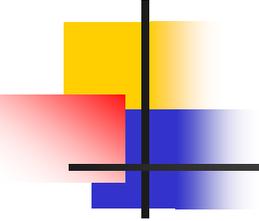
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- Global strain collection
  - MRSA and MSSA
  - Swine and veal isolates
- Study showed that zinc resistance and the *czrC* gene widespread among CC398 MRSA
- Suggests use of zinc in feed might have contributed to the emergence of MRSA.

# Growth promotants as the evil

## MRSA cases in Denmark (all types)





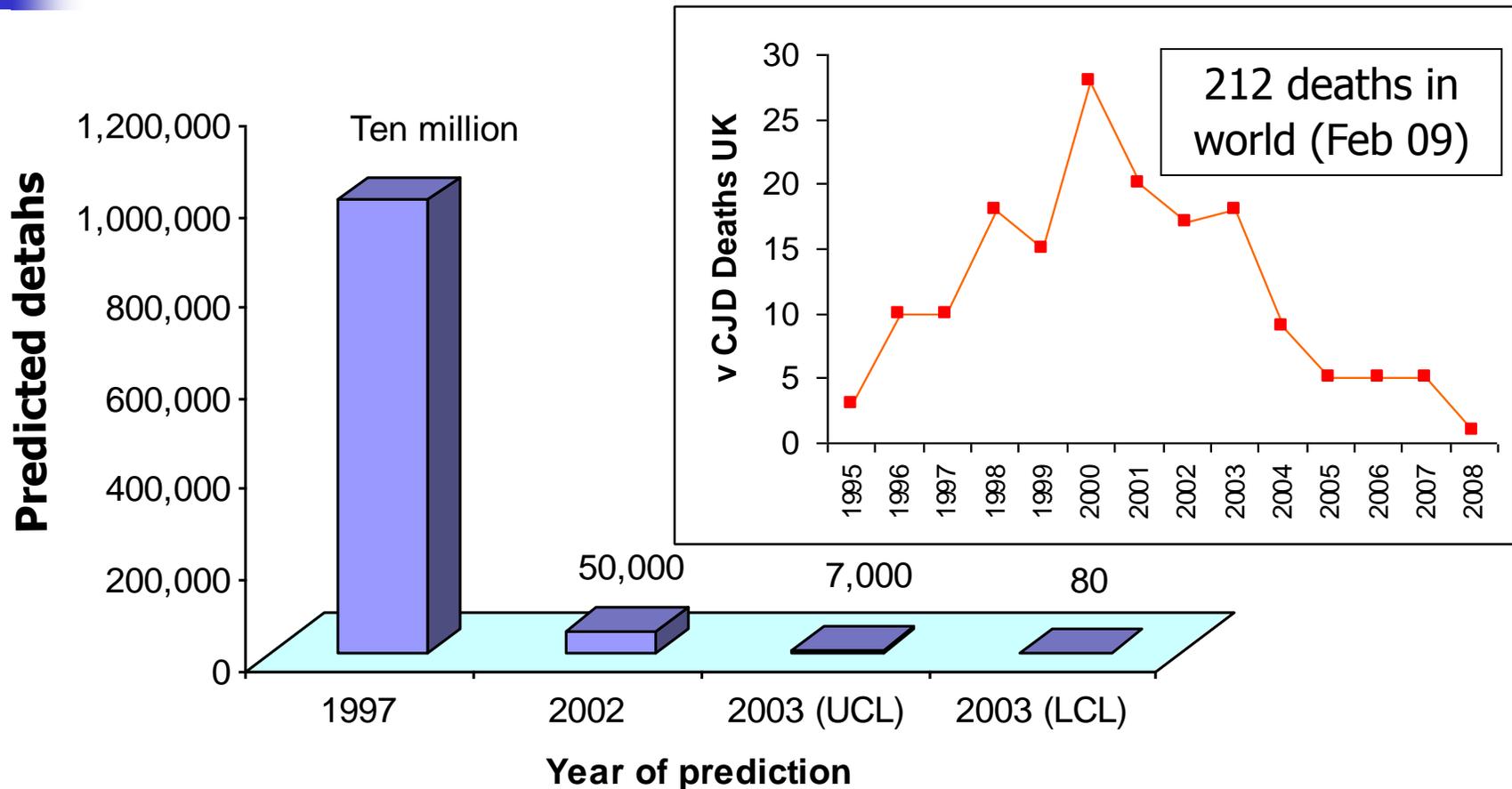
# The rest of the story.....

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- Understanding LA-MRSA is embryonic
- Naïve perceptions of complex epidemiology
  - All MRSA found in livestock are ST398
  - Livestock are the only reservoirs of ST398
- ST398 isolates of diverse genotype and geographic origin may also be epidemiologically distinct
- Requires systematic investigation of *S. aureus* epidemiology in animals and humans.
- Occupational health and public relations concerns
  - Not yet a public health concern!

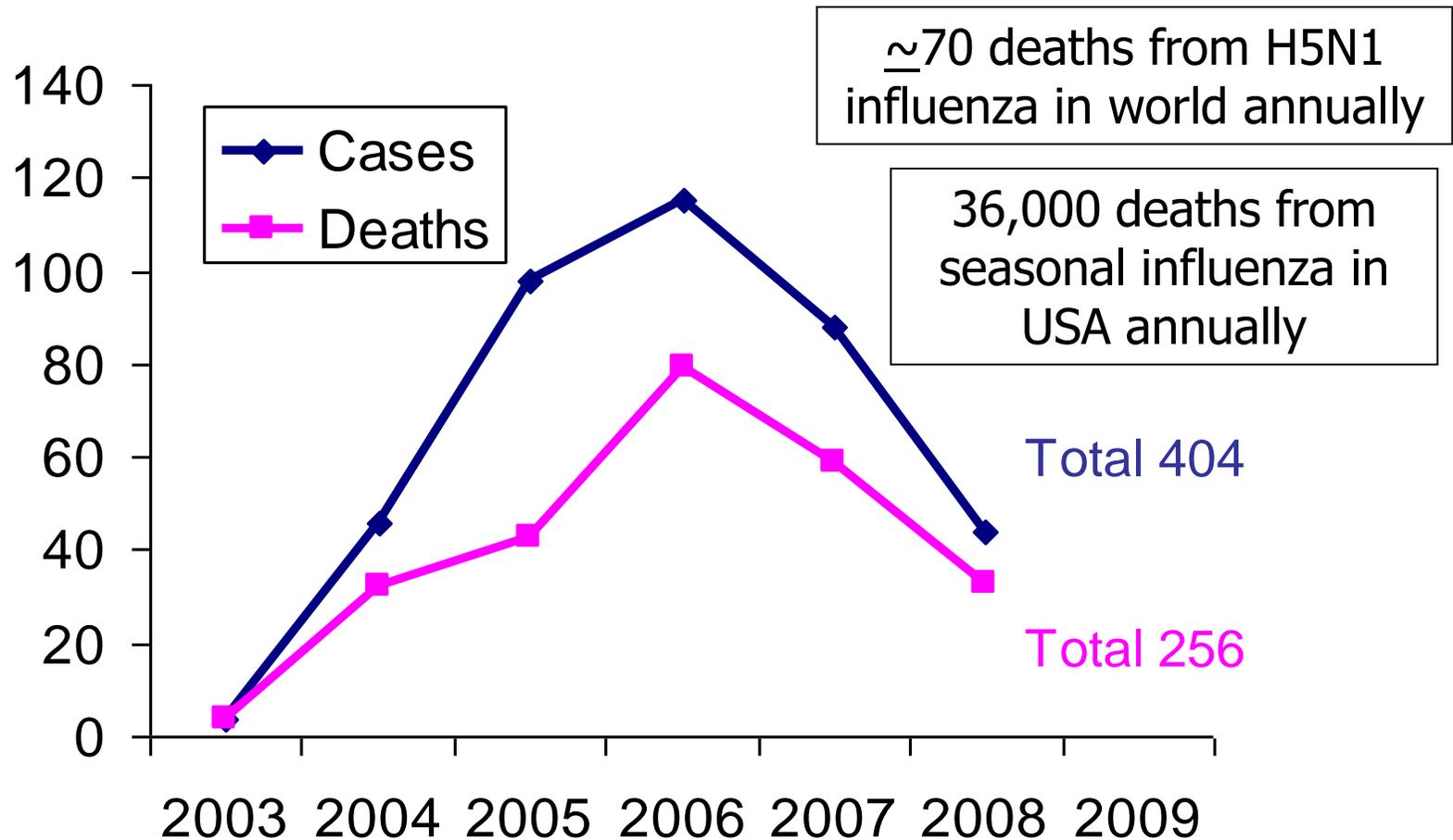
# Panic and Prediction

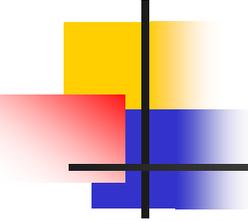
Predicted of size of vCJD epidemic



# Reported deaths from H5N1 influenza

WHO





# Staphylococcal foodborne disease

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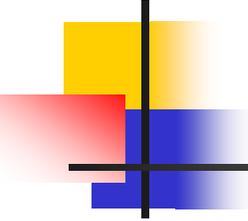
- Common foodborne disease
  - Ingestion of preformed toxins in food
  - Only rare reports of MRSA involvement
  - All livestock associated ST398 investigated have been negative for enterotoxins
- Antimicrobials not indicated
  - Resistance of isolates irrelevant
- MRSA in meat relevant only due to exposure risk from food handling

# MRSA prevalence in Dutch meat

de Boer et al (2009)

Meat type	<i>n</i>	Number (%) positive
Beef	395	42 (10.6)
Veal	257	39 (15.2)
Pork	309	33 (10.7)
Lamb/mutton	324	20 (6.2)
Chicken		
– Total	520	83 (16.0)
– NL+ other EU countries	302	
– Import (non-EU countries)	162	
– Biological	56	
Turkey	116	41 (35.3)
Fowl	118	4 (3.4)
Game	178	4 (2.2)
Total	2217	264 (11.9)

Most isolates  
ST398

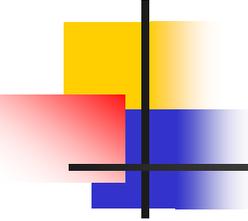


# MRSA in meat: what are the implications?

De Boer et al (2009)

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- High prevalence of MRSA in raw meats not shown to contribute to the dissemination MRSA to humans.
- MRSA numbers so low that risk of colonization is slight.
- Conditions for growth of *S. aureus* in raw meats are poor.



# MRSA in meat – what are the implications?

---

- VWA Risk Assessment Bureau (Holland)
  - Foodstuffs play a negligible role, in the spread of MRSA.
- Weese and van Duijkeren (2009)
  - Evidence implicating food as the source of infection appears to be tenuous.
  - Clinical relevance of MRSA contamination of food, ... is currently unclear
- DANMAP 2010
  - Frequent occurrence of MRSA in meat combined with no/very few cases in urban areas makes it safe to conclude that there is **very little if any risk for meat being a risk for contracting MRSA CC398.**