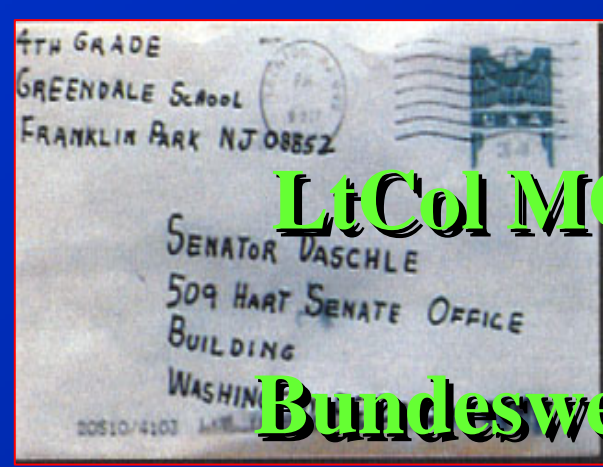


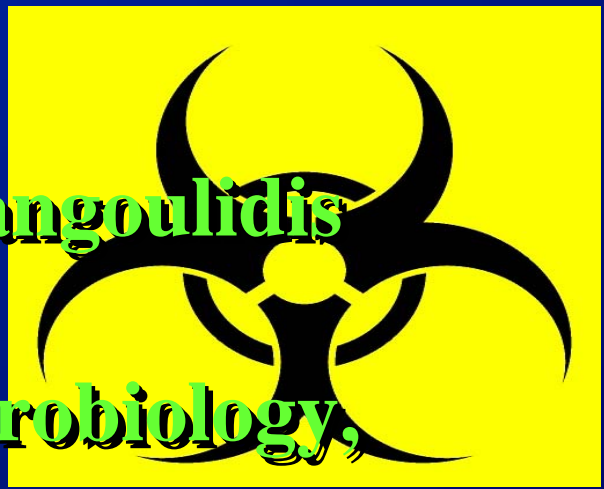


Realtime-PCR of Bioterrorism agents



LtCol MC Dr. Dimitrios Frangoulidis

**Bundeswehr Institute of Microbiology,
Munich**





- „Bioterrorism“ = 4.365
(1996-9/2001 = 205;
10-12/2001 = 240)
- „qPCR“ = 1.481
- „Bioterrorism“ & „qPCR“ = 132



The „dirty dozen“

Bacteria

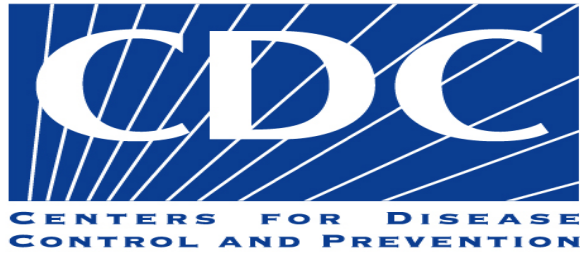
- *Yersinia pestis*
- *Bacillus anthracis*
- *Brucella* spp.
- *Francisella tularensis*
- *Burkholderia mallei*/
B. pseudomallei
- *Coxiella burnetii*

Virus

- Variolavirus
- Venezuelan Equine
Encephalitisvirus
- Marburgvirus

Toxins

- Botulinumtoxins
- Staphylokokken-
enterotoxins
- Ricin



Category Characteristics

A Easy dissemination or contact transmission
 Cause high mortality rates and have the potential for major public health impact
 May cause public panic and social disruption
 Require special action for public health preparedness

Anthrax (*Bacillus anthracis*)
 Botulism (*Clostridium botulinum* toxin)
 Plague (*Yersinia pestis*)
 Smallpox (*Variola major*)
 Tularemia (*Francisella tularensis*)
 Viral hemorrhagic fevers

B Moderately easy to disseminate
 Result in moderate morbidity rates and low mortality rates
 Require specific enhancements of CDC's diagnostic capacity and enhanced disease surveillance

Brucellosis (*Brucella* species)
 Epsilon toxin of *Clostridium perfringens*
 Food safety threats (*Salmonella* species, *Escherichia coli*, *Shigella*)
 Glanders (*Burkholderia mallei*)
 Melioidosis (*Burkholderia pseudomallei*)
 Psittacosis (*Chlamydia psittaci*)
 Q fever (*Coxiella burnetii*)
 Ricin toxin (from castor beans)
 Staphylococcal enterotoxin B
 Typhus fever (*Rickettsia prowazekii*)
 Viral encephalitis
 Water safety threats (*Vibrio cholerae*, *Cryptosporidium parvum*)

C Emerging pathogens engineered for mass dissemination
 Availability
 Ease of production and dissemination
 Potential for high morbidity and mortality rates and major health impact

Emerging infectious diseases such as Nipah virus and hantavirus



but....“most wanted“:



PubMed

**„PCR and bacteria“ =
95.976**





Identification of Bioterrorism



agents

- **UNAMBIGUOUS**
- Specific
- Sensitive
- Fast
- Reproducible
- Safe



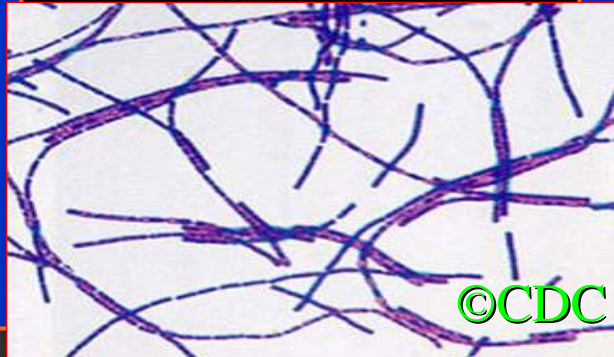
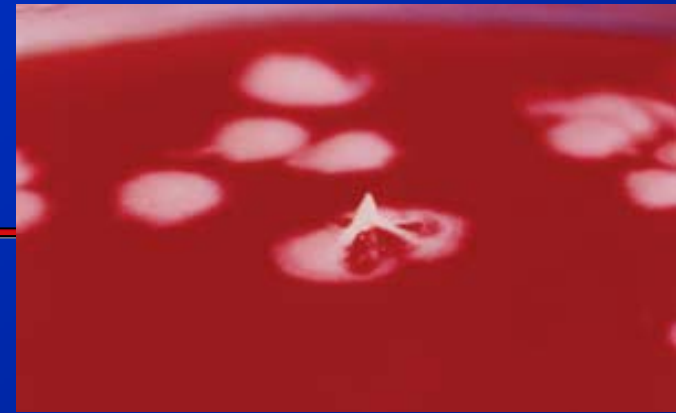
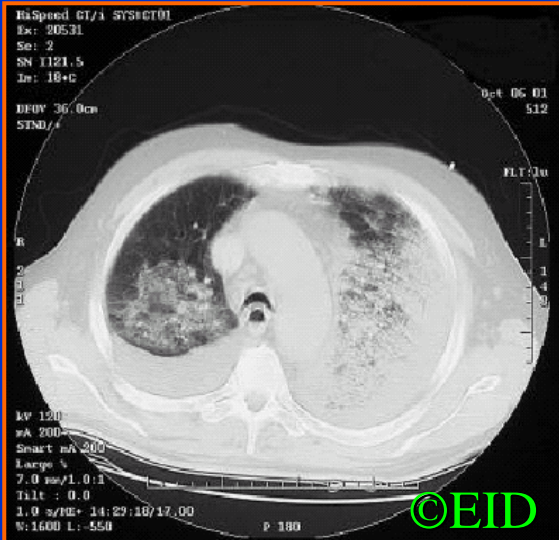


Kochs „Postulate“

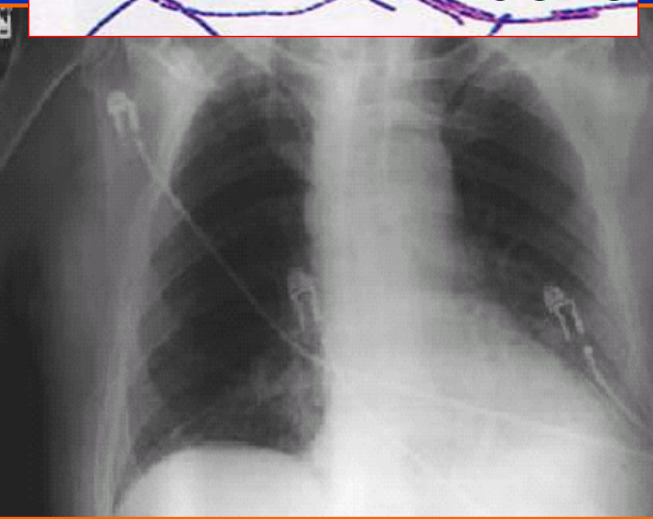


- The anthrax bacillus, *Bacillus anthracis*, was the first bacterium shown to be the cause of a disease. In 1877, Robert Koch grew the organism in pure culture, demonstrated its ability to form endospores, and produced experimental anthrax by injecting it into animals.





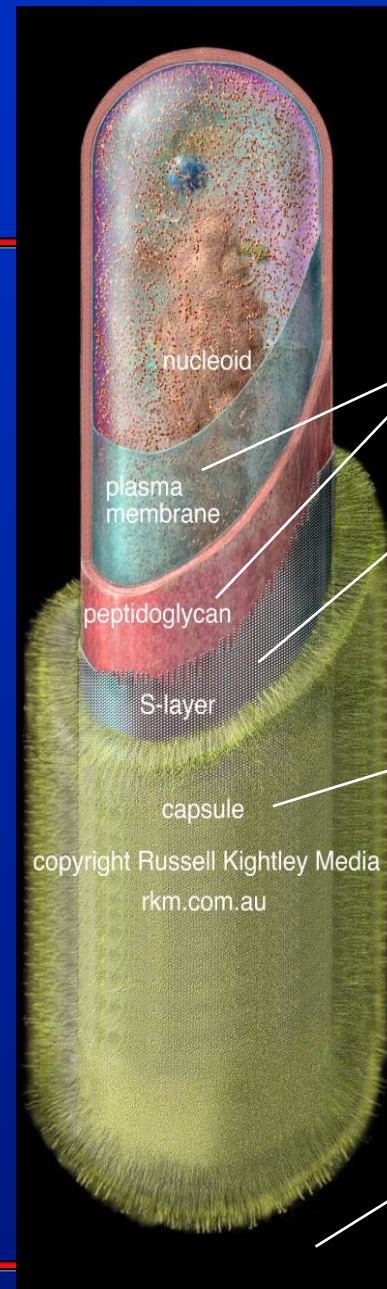
„Anthrax“





Bacillus anthracis:

- * Gram-positive
- * aerobe rod
- * 1 - 1.2 μ m in width x
3 - 5 μ m in length
- * sporeforming
- * Genotypically and phenotypically
very similar to *Bacillus cereus* and
thuringiensis
- * Toxin producer



Sacculus

** S-Layer
Sap + EA1

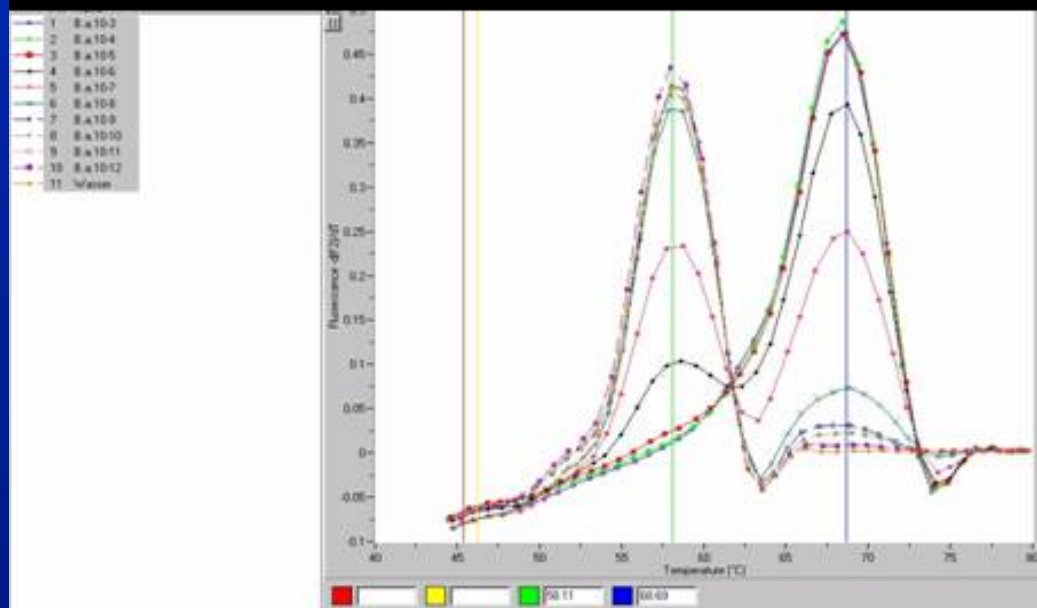
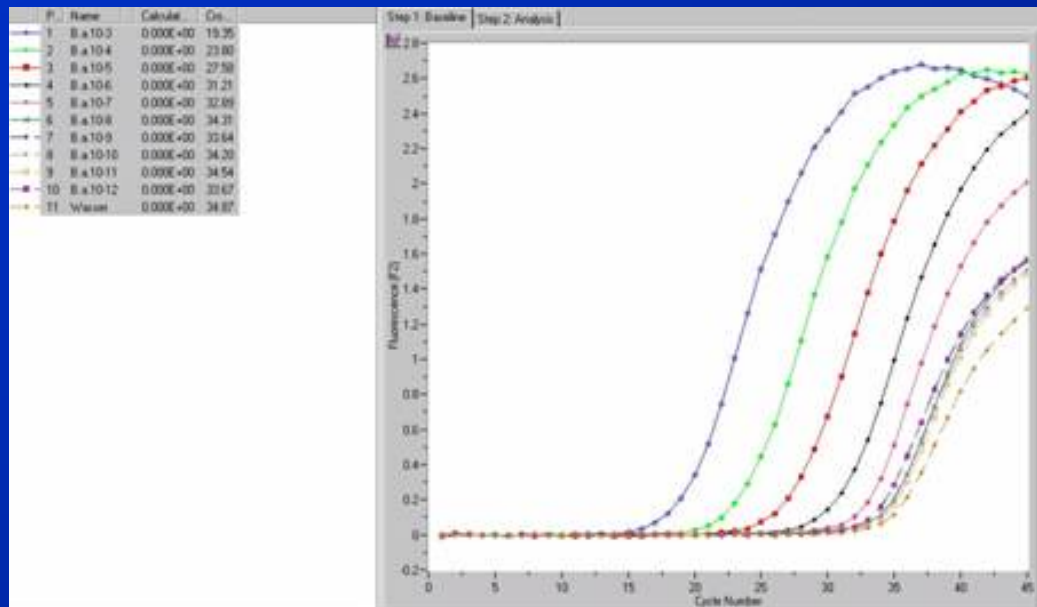
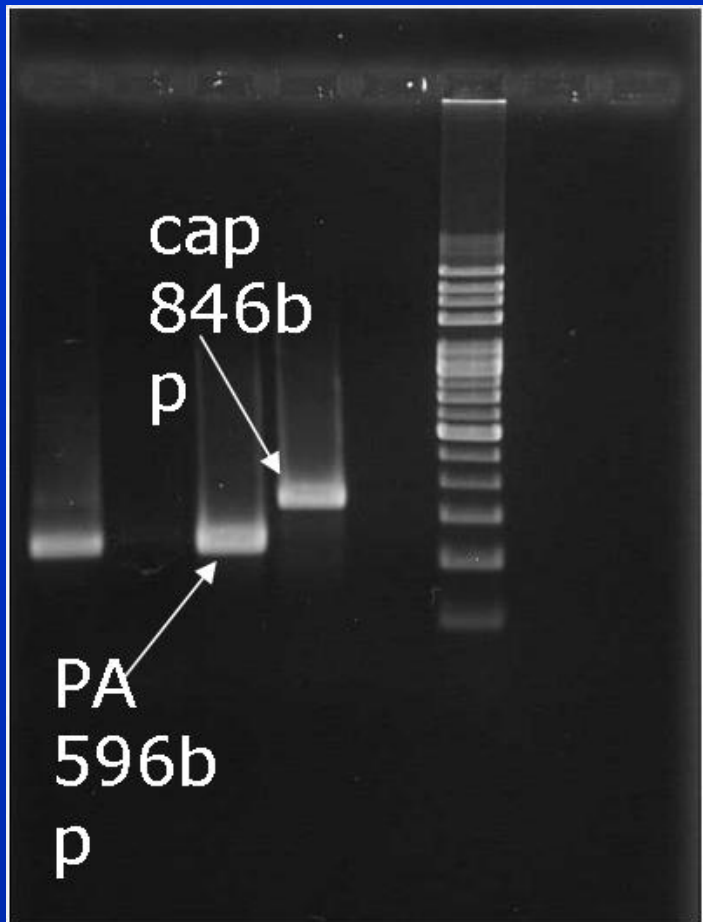
** Capsule: poly- $\gamma\delta$ -Glutamin-
acid

** Toxins



VIRULENCE-FACTORS (plasmid-coded)

- **Poly- γ -D-Glutamyl-capsule: pX01**
- **Exotoxin-complex: pX02**
 - **Lethalfactor (LF)**
 - **Edemafactor (EF)**
 - **protective Antigen (PA)**



Characterization of *Bacillus cereus* Isolates Associated with Fatal Pneumonias: Strains Are Closely Related to *Bacillus anthracis* and Harbor *B. anthracis* Virulence Genes†

Alex R. Hoffmaster,^{1*} Karen K. Hill,⁴ Jay E. Gee,¹ Chung K. Marston,¹ Barun K. De,¹ Tanja Popovic,²
David Sue,^{1‡} Patricia P. Wilkins,¹ Swati B. Avashia,^{3,4§} Rahsaan Drumgoole,⁴ Charles H. Helma,⁵
Lawrence O. Ticknor,⁶ Richard T. Okinaka,⁵ and Paul J. Jackson⁷

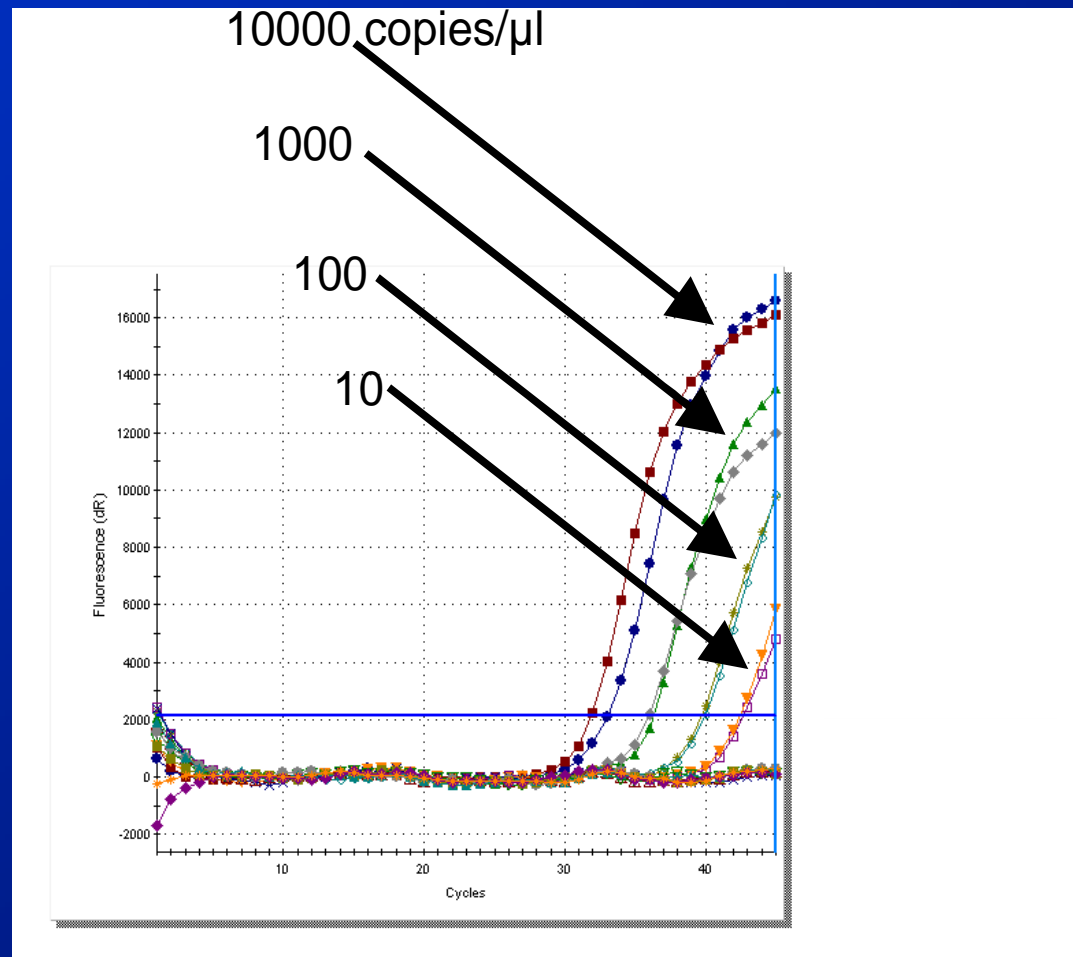
Marker	Number of strains testes versus amplified		Reference
	<i>B. anthracis</i>	<i>B. cereus</i>	
Ba831	47/47	4/60	Ramisse et al 1996, JAM, 87, 224ff
rpoB	14/14	5/36	Ellerbrock et al FEMS Microb Lett. 2002, 214, 51ff
BA5510	4/4	2/289	Olsen et al JMM 2007, 71, 265ff
gyrB	1/1	2/23	La Duc et al JMM 2004, 56, 383ff.
saspB	392/392	0/56	Hoffmaster et al EID 2002, 8, 1178ff
plcR	89/89	0/29	Easterday et al JCM 2005, 43, 1995ff
gyrA	43/43	0/49	Hurtle JCM 2004, 42, 179ff
BA5345	142/142	0/236	Antwerpen et al, MCP 2008, 22, 313ff



„Dhp61-PCR“



The detection limit was calculated using Probit-analysis to be 12.7 copies per μl (95% confidence interval, 10.2 - 17.5 copies).





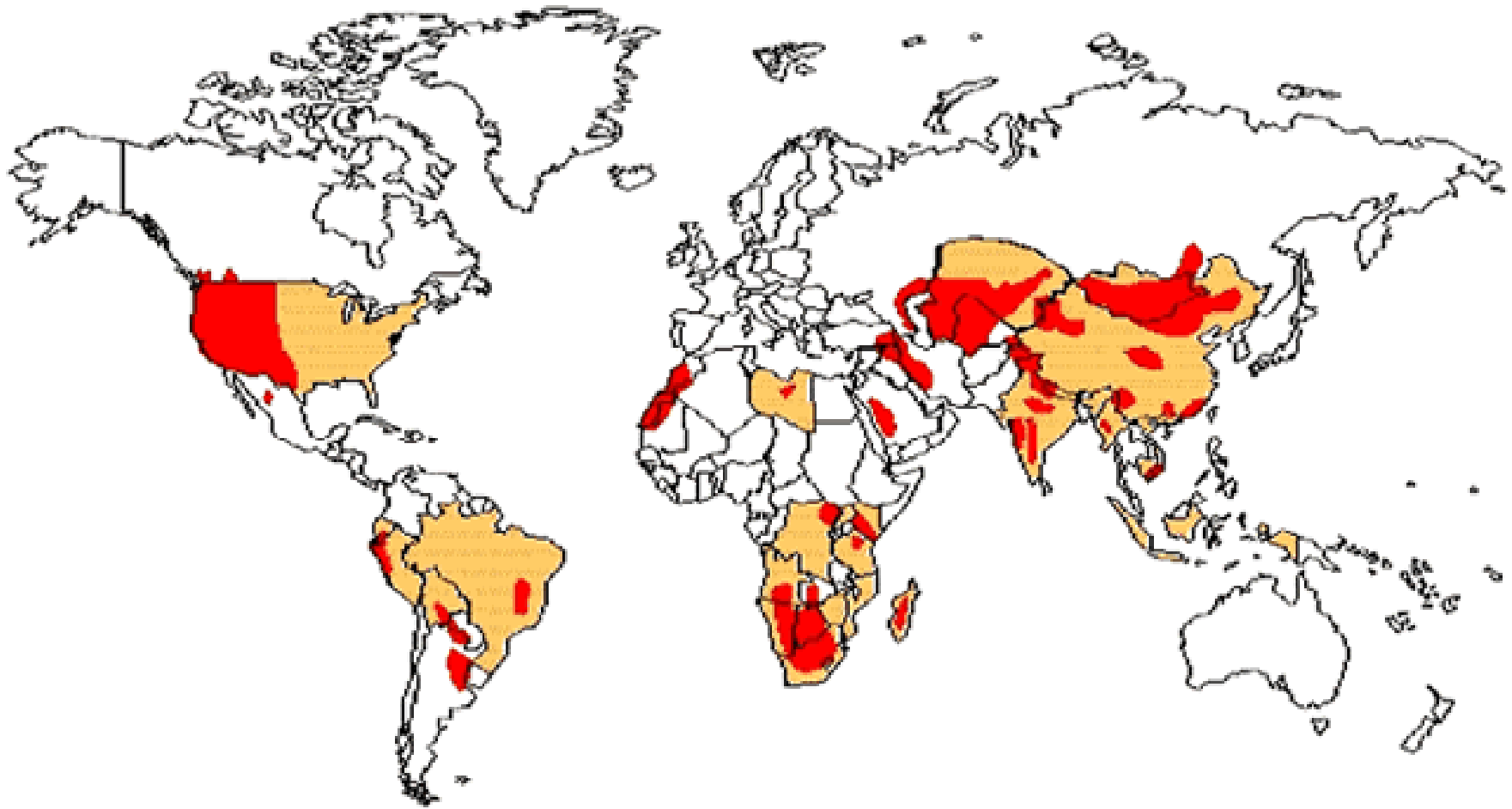
Yersinia pestis



1. Gram-negative, short rod
2. Family: Enterobacteriaceae
3. 11 *Yersinia*-Species,
3 human-pathogen
 - *Y. pestis*
 - *Y. pseudotuberculosis*
 - *Y. enterocolitica*



World Distribution of Plague, 1998



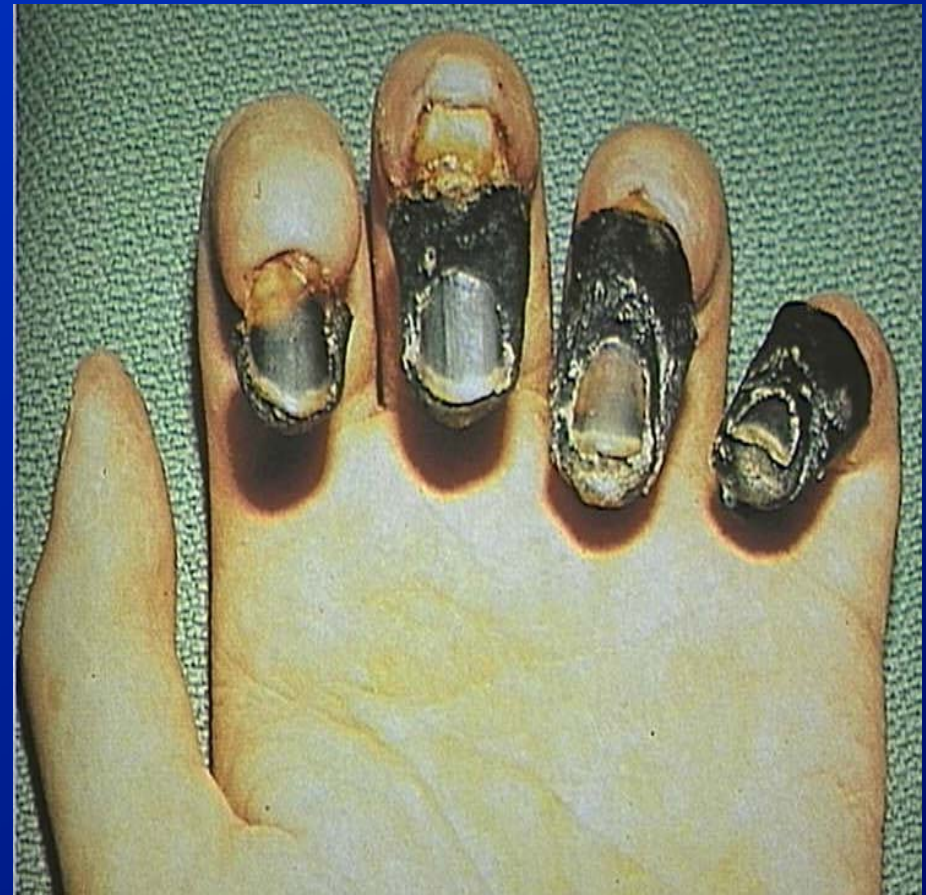
■ Countries reported plague, 1970-1998.

■ Regions where plague occurs in animals.

WHO, Geneva, Switzerland



Clinical aspects



CDC, Atlanta, USA

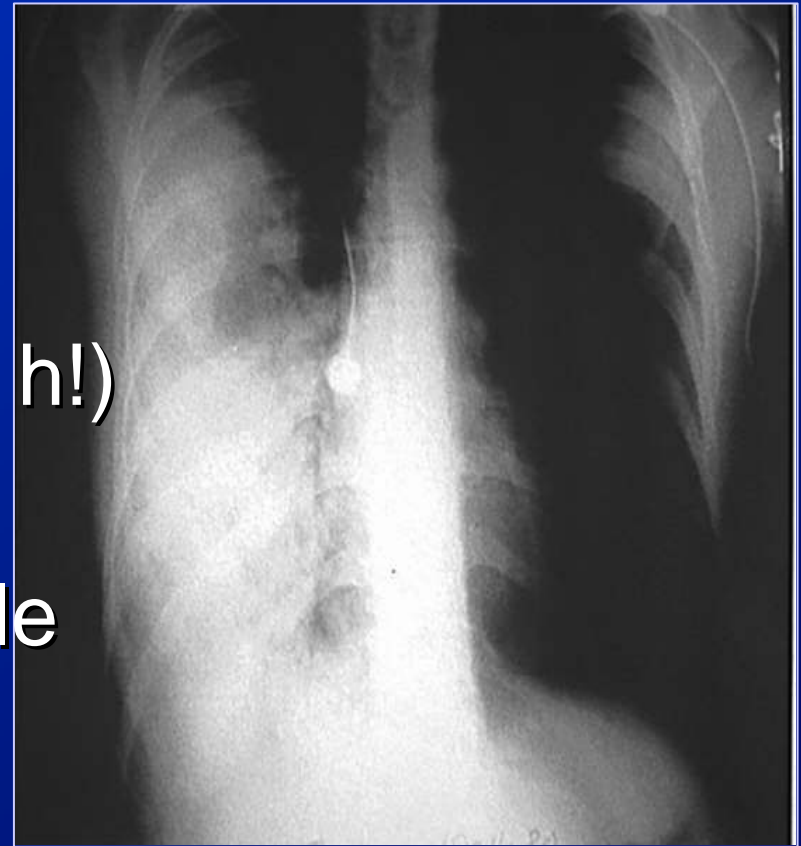




Pulmonary Plague – „The Pox of Bacteria“



- aerogen Infection
(person-to-person)
- less than 100 bacteria
- short Incubationtime (6 h!)
- high letality (>90%)
- no safe vaccine available

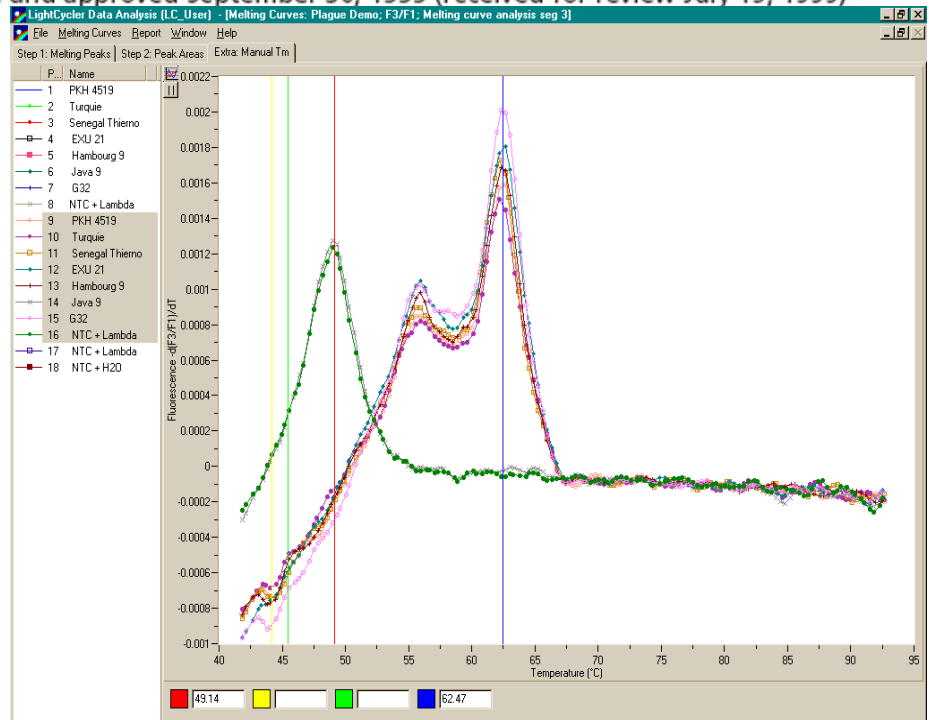


Yersinia pestis, the cause of plague, is a recently emerged clone of *Yersinia pseudotuberculosis*

Mark Achtman^{*†}, Kerstin Zurth^{*}, Giovanna Morelli^{*}, Gabriela Torrea[‡], Annie Guiyoule[‡], and Elisabeth Carniel[‡]

^{*}Max-Planck-Institut für molekulare Genetik, Ihnestr. 73, 14195 Berlin, Germany; and [‡]Unité de Bactériologie Moléculaire et Médicale, Laboratoire des Yersinia, Institut Pasteur, 75724 Paris Cedex 15, France

Edited by Richard M. Krause, National Institutes of Health, Bethesda, MD, and approved September 30, 1999 (received for review July 19, 1999)





Multiplex LC-PCR assays for the specific detection of *Yersinia pestis*



	Targets		Internal amplification control
Multi A	16S rRNA gene	<i>pla</i> (pPla)	Bacteriophage Lambda DNA
Multi B	<i>ymt</i> (pMT1)	<i>caf1</i> (pMT1)	Bacteriophage Lambda DNA

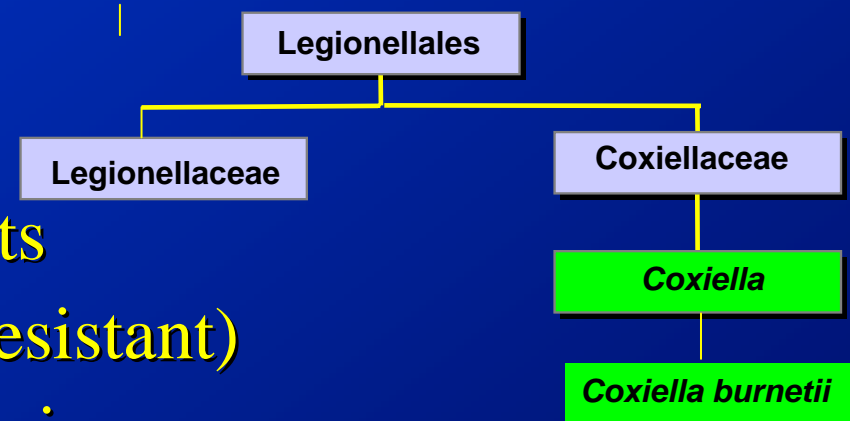




Properties *Coxiella burnetii*



- Pathogen of the Zoonosis Q fever
- gram-negative
- obligate intracellular
- small- u. large cell variants
- „Spore-like“ –particles (resistant)
- extrem low ID: 1-10 organism
- Family of Legionellales
- CDC-Classification 2 (Aerosol, extreme low ID <10 Bacteria)

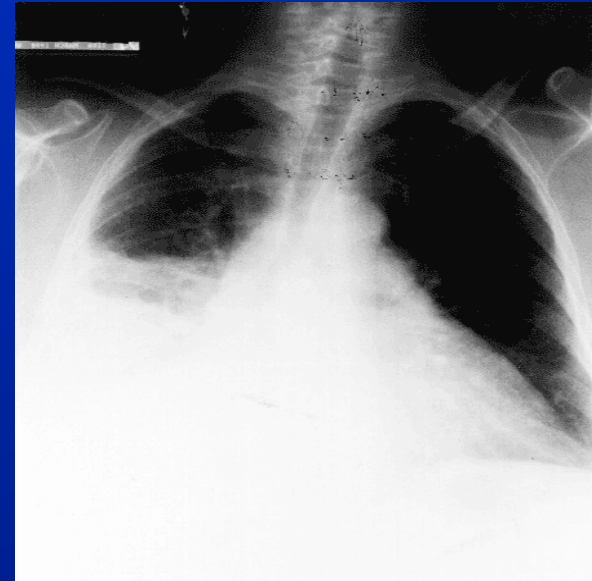




Coxiella burnetii – Q-Fever



- periodical Outbreaks worldwide incl. Germany (e.g.: Dillenburg, Aschaffenburg 2008, NL 2008 - >600 cases)
- 98%: Acute (Pneumonia), 2% : Chronic (Endocarditis)



Diagnostic Gold-Standard: Serology
(ELISA, IFA)





Q Fever in Germany



Year	Location	Cases	Transmission
1993	Sontra	84	birth products, sheep
1997	Gießen	68	lambing
2003	Soest	299	lambing
2005	Jena	322	sheep flock





Q Fever - PCR indications?



- acute Q-fever: serum samples, sputum, BAL,...(low DNA amount)
- chronic Q-fever: blood, placenta/amnion fluid, heart valve (low to moderate DNA)
- „Bioterrorism“ samples (moderate to high DNA)
- veterinary medicine (Milk, feces, birth products – moderate to high DNA)

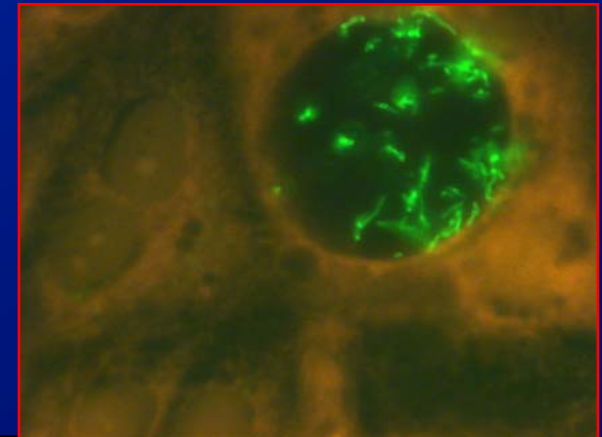




PCR requirements



- Sensitivity (high?!)
- Specificity (high!!)
- Controls (pos., neg., internal), multiple targets?
- Stability of reagents/robustness (RT, 4°/-20°C)
- „Stability“ of results, reproducible?
- Hands-on-time/ „easy-to-use“
- request „frequency“
- Costs
- availability guarantee



„I have a dream.....“

+ 100% Specificity

+ High Sensitivity (<10 cp/μl)

+ Competitive Internal Control

**+ „real“ Quantification with pre-diluted standards (4x)
single use or stabil after solution**

+ Melting-Curve-Analysis

+ „Ready-to-use“ reagents

+ (multi-target Assay)

+ availability guarantee