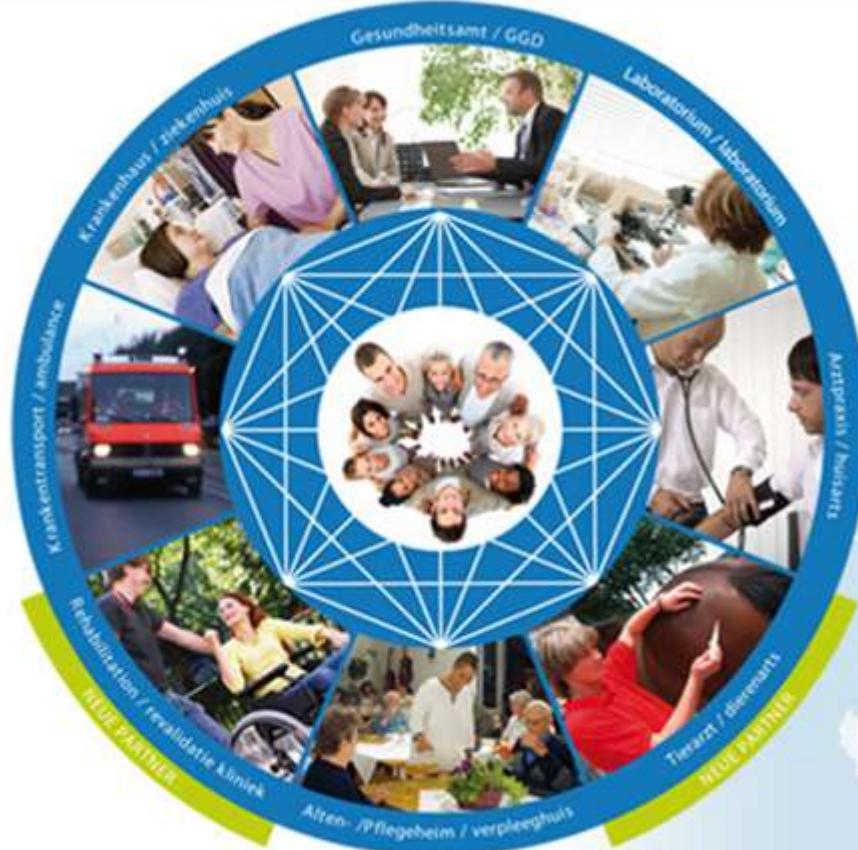


Molecular Epidemiology of MRSA in Europe



Alex W. Friedrich, University of Muenster, Germany

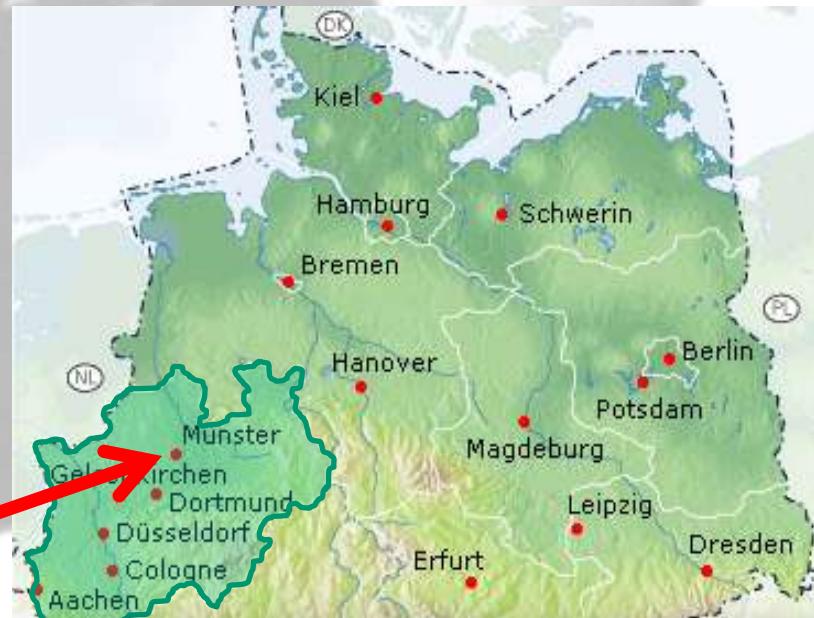
State of North Rhein-Westfalia

18 Mio inhabitants

2nd largest university in
Germany



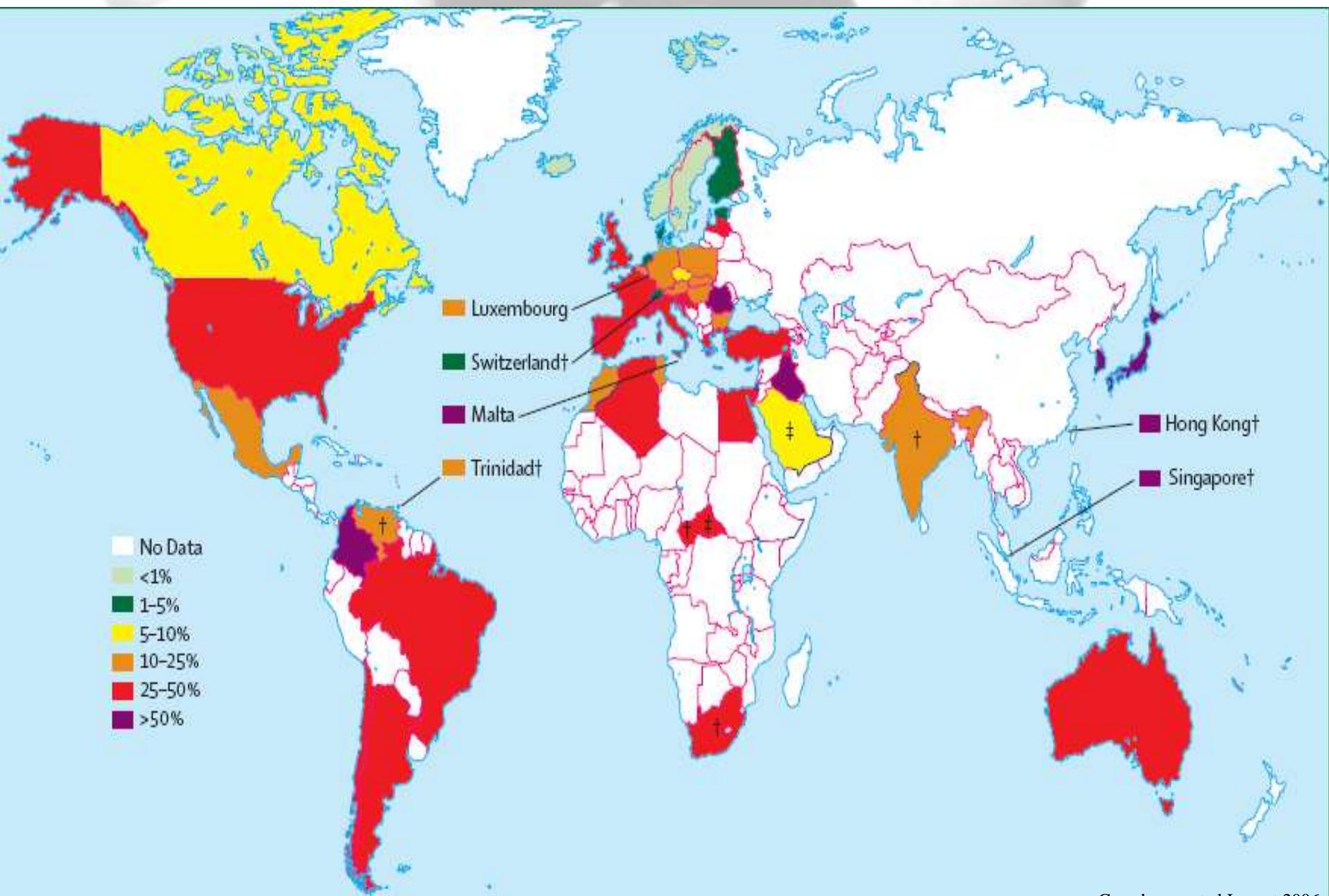
Townhall of the Westfalian Peace Treaty
City of Münster





Münsterland

MRSA prevalence in *S. aureus* bacteraemia worldwide



MRSA in Europe

- Affects **150.000 patients** per year in Europe
- **MRSA-infection:**
Up to 2,4 times significantly higher Mortality
- Issue of **patient safety**
- **Reason for emerging MRSA**
 - insufficient hygiene in hospitals
 - too many antibiotics given
 - lack of trans-mural case management
- **Nasal carriage** origin of infection
-> Finding and treating carriers avoids infection !
- **MRSA additional costs:** Per patient: 5.000 € to 20.000 €
EU healthcare system: 380 Mio € per year
- **Differences in MRSA-Prevalence** between countries



Methicillin-resistant *Staphylococcus aureus* (MRSA): burden of disease and control challenges in Europe

R Köck^{1,2}, K Becker², B Cookson³, J E van Gemert-Pijnen⁴, S Harbarth^{5,6}, J Kluytmans^{7,8}, M Mielke⁹, G Peters², R L Skov¹⁰, M J Struelens^{11,12}, E Tacconelli¹³, A Navarro Torné¹², W Witte¹⁴, A W Friedrich (alexander.friedrich@ukmuenster.de)¹

1. Institute of Hygiene, University Hospital Muenster, Muenster, Germany

2. Institute of Medical Microbiology, University Hospital Muenster, Muenster, Germany

3. Laboratory of Healthcare Associated Infections, Centre for Infections, Health Protection Agency, London, United Kingdom

4. Faculty of Behavioural Sciences, University of Twente, Enschede, the Netherlands

5. Infection Control Program, University Hospitals of Geneva, Geneva, Switzerland

6. University of Geneva Medical School, Geneva, Switzerland

7. VU University Medical Centre, Amsterdam

8. Amphia Hospital,

9. Robert Koch Institut

10. Statens Serum Ins

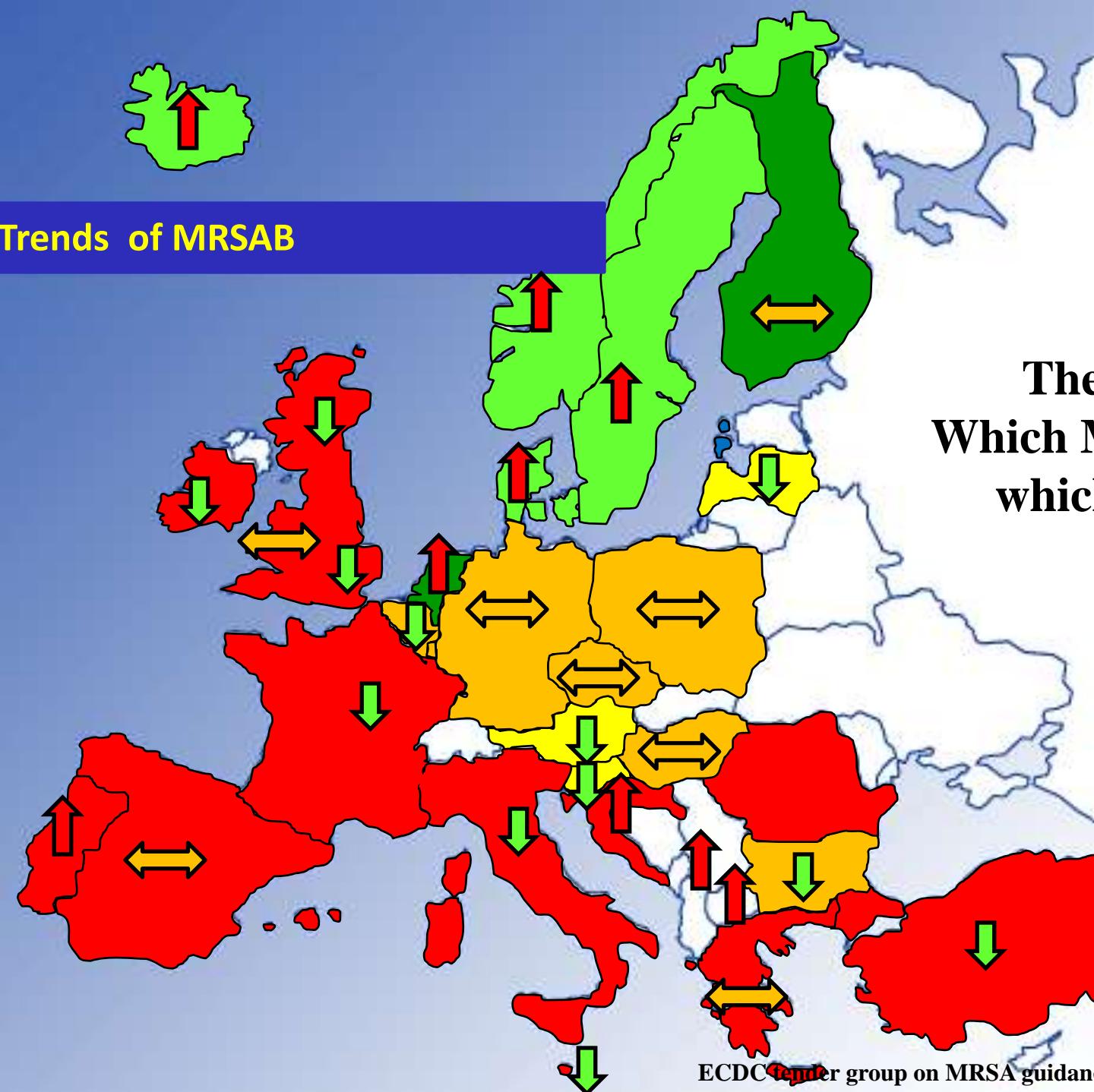
11. Department of Microbiology (not yet started)

12. Scientific Advice Unit

13. Department of Infectious Diseases

14. National Reference Laboratory

Type, place and period of study	Odds ratio/hazard ratio for MRSA-associated mortality (95% CI)	Percentage mortality in MRSA patients
Single-centre, university hospital, Taiwan, 1990–2004	1.78 (1.3–2.44)	50% ^a
Single-centre, university hospital, Belgium, 1992–1998	1.93 (1.18–3.18)	64% ^b
Single-centre, teaching hospital, UK, 1995–2000	1.72 (0.92–3.20)	12% ^c
Veterans affairs health care system, USA, 1995–2003	1.8 (1.2–3.0)	34% ^d
Single-centre, university hospital, USA, 1996–2001	5.4 (1.5–18.7)	35% ^d
Single-centre, university hospital, France, 1997–1998	2.97 (1.12–7.88)	43% ^e
Single-centre, tertiary-care teaching hospital, USA, 1997–2000	1.2 (0.68–2.12)	23% ^f
Multi-centre, Germany, 1997–2002	3.84 (1.51–10.2)	17% ^c
Two centres, teaching hospital UK, 1997–2004	1.49 (0.99–2.26)	34% ^a
Single-centre, teaching hospital, USA, 1999–2001	1.4 (0.7–3.0)	31% ^a
Single-centre, teaching hospital, Brazil, 2000–2001	2.52 (0.96–6.6)	55% ^a
Single-centre, university hospital, Taiwan, 2001–2006	0.73 (0.21–2.60)	10% ^a
Single-centre, university hospital, Belgium, 2002–2004	3.04 (1.15–8.04)	42% ^g
Single-centre, university hospital, Germany, 2002–2007	2.6 (1.4–4.9)	42% ^d
Single-centre, tertiary care, USA, 2004–2005	5.1 (1.1–22.9) ⁱ	47% ^h



The question is: Which MRSA are rising, which are falling ?

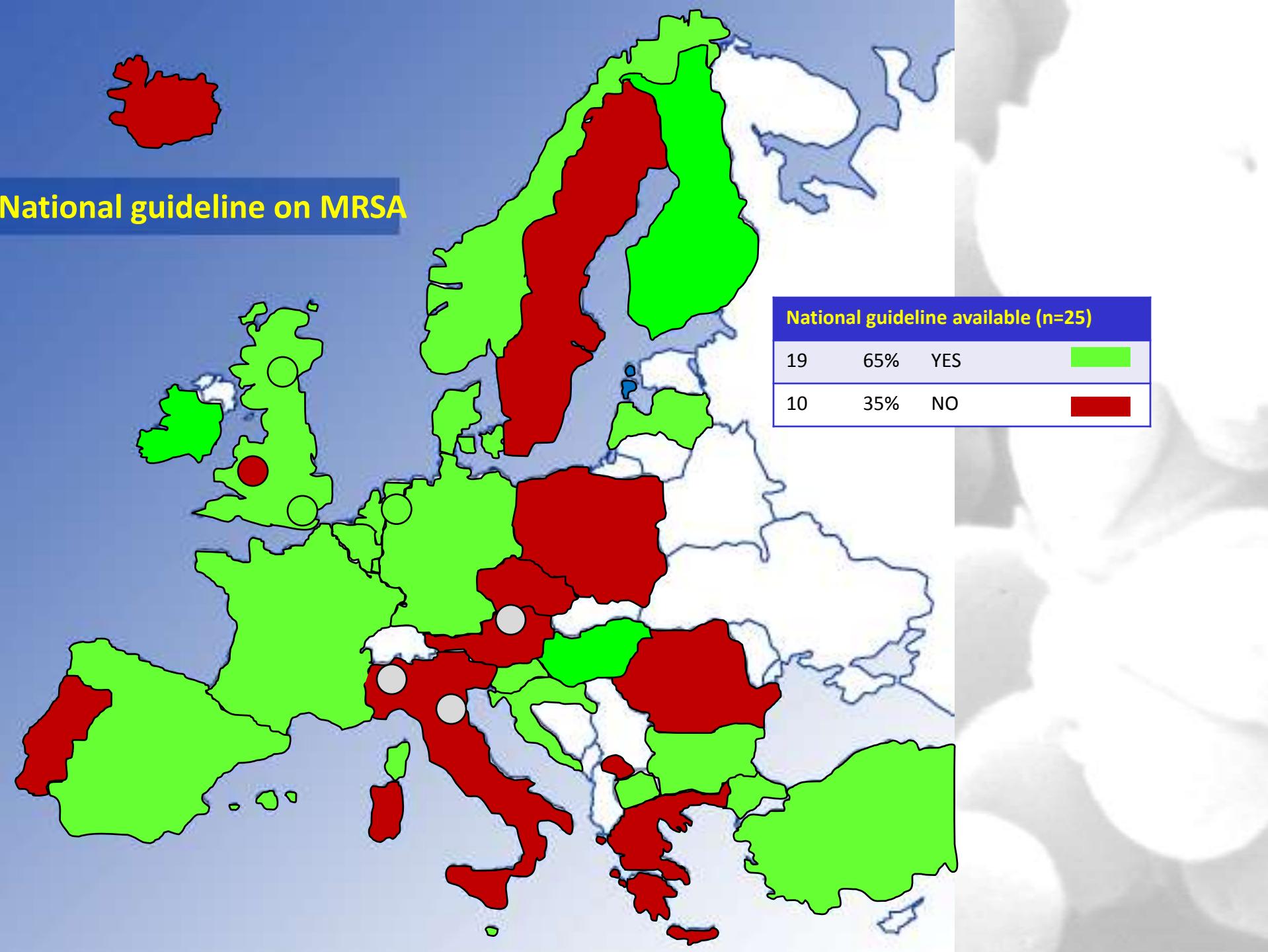
	No data
	<1%
	1-5%
	5-10%
	10-25%
	25-50%

National guideline on MRSA

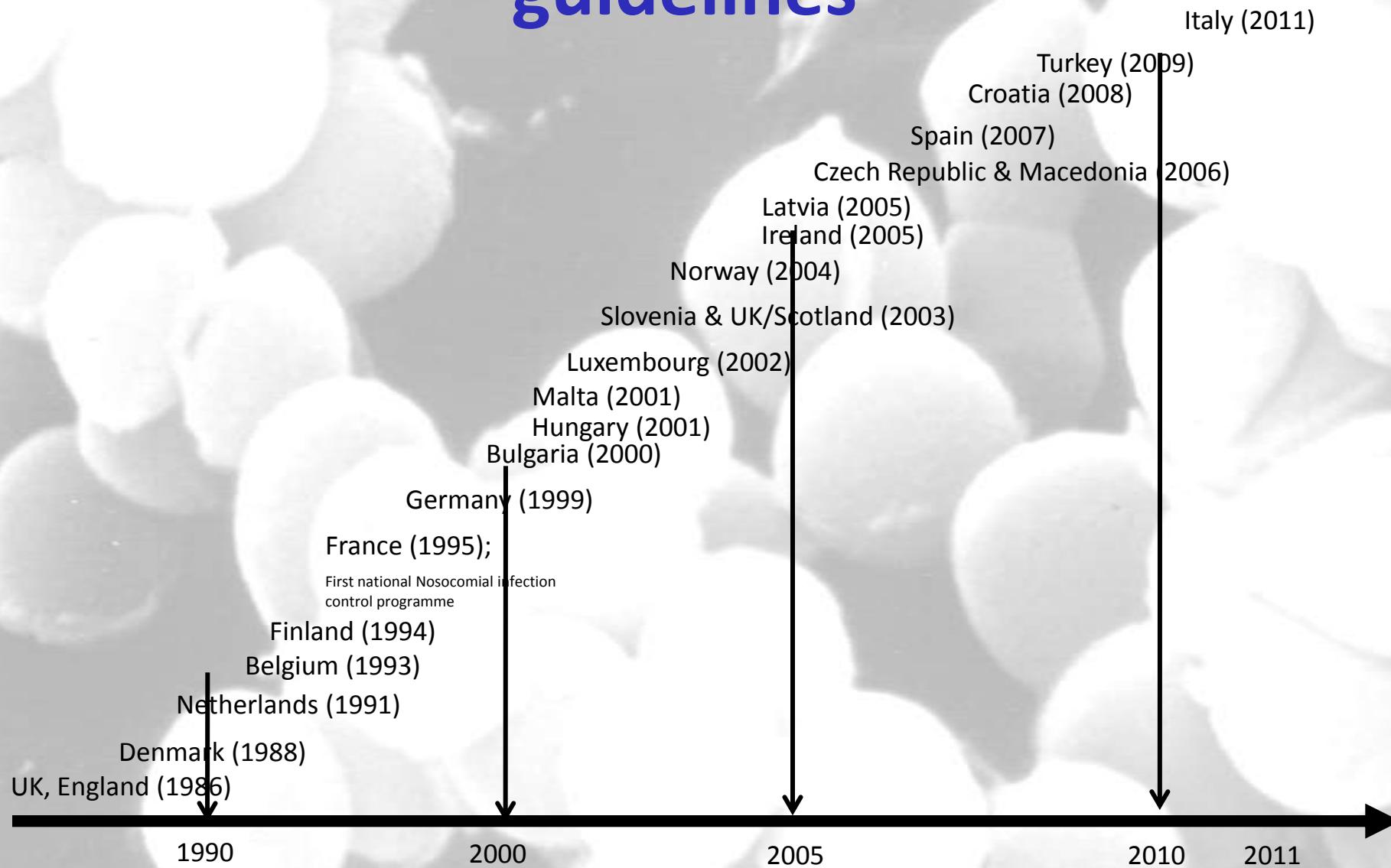
National guideline available (n=25)

19 65% YES

10 35% NO



First edition of national MRSA guidelines





Preventive Microbiology

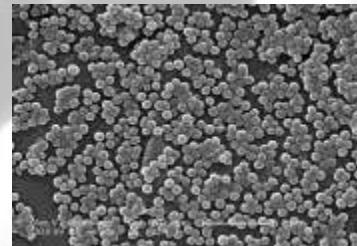
=

infection control

Robert Koch,
founder of modern bacteriology



Vibrio cholerae,



MRSA,

Cholera-infection control measures (1908):

„Everything depends on the fact that we have valid detection methods for the microorganisms.

This counts especially for the human carriers who contribute most for the spread of the disease“

Speech of Robert Koch on 11.2.1908 in front of the Medical order of Berlin
Deutsche Medizinische Wochenschrift, 1908, Nr. 8

MRSA „search&follow-Strategy“ (2010)

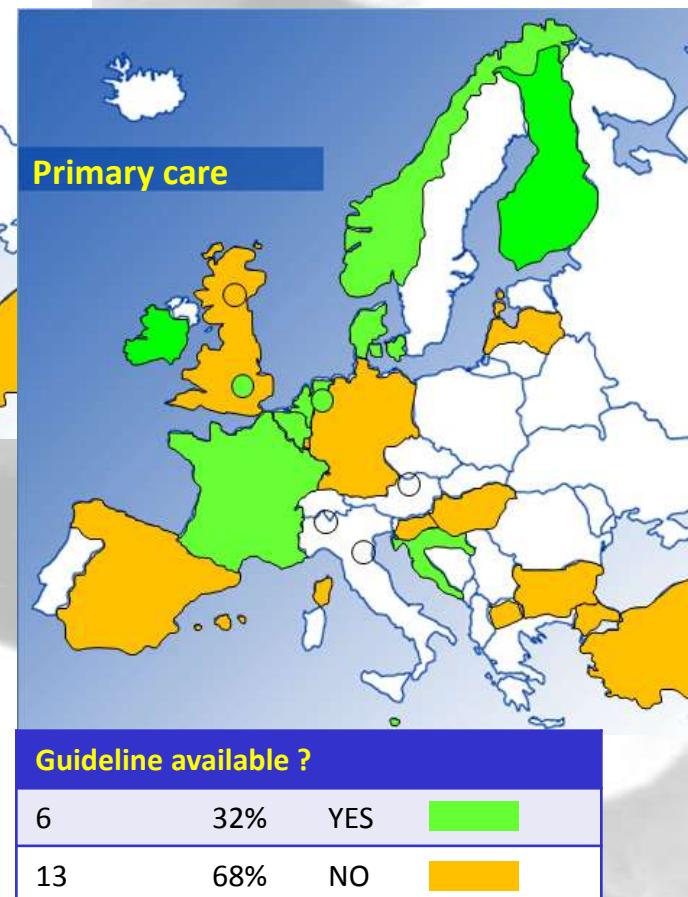
Colonisation comes before infection

- A: Rapid identification of MRSA-carriers
- B: Decolonization of MRSA-carriers,

In order to:

- a.) prevent infection in the patient
- b.) prevent spread to other patients
- c.) start early an optimal antibiotic therapy

Health-care settings covered by MRSA national guideline



SeqNet.org does...

A. Capacity building for sequence based typing in European laboratories

- International workshops since 2004, next 5-8 of October 2010
Münster, Zagreb, Sophia

B. Internal quality control

- SOP for internal quality control

C. External quality control

- SeqNet.org certification
- Annual proficiency testing
- Continuous quality control

D. Curatorship for spa server

- Synchronization of typing data
- Access via StaphType and Bionumerics
- Study groups within spa server
(Euregio study area, SRL-study group)

The screenshot shows the SeqNet.org website homepage. At the top, there is a navigation bar with links for 'Advertise' and 'http://www.seqnet.org'. On the right side of the header, there are links for 'Home | Membership | Contact'. Below the header, there is a large logo featuring a stylized orange 'S' and 'N' intertwined with a yellow ribbon-like shape, set against a background of blue stars. The main content area has a white background. At the top of this area, it says 'European Network of Laboratories for Sequence Based Typing of Microbial Pathogens'. Below this, a section titled 'Welcome to SeqNet.org' is visible. It includes a brief description of the network's purpose: 'SeqNet.org is an initiative of currently 57 laboratories from 29 European countries. The main objective is to establish a European network of excellence for sequence based typing of microbial pathogens.' There is also a link to 'How to participate?'. Further down, there is a 'SpaServer Summary' section with a table of statistics:

Spa-types:	5920
Repeats:	342
Total strains:	102522
Strain records:	73961
Strain countries:	70
Registered users:	320
User countries:	38

Next to the summary table, there is a list of links:

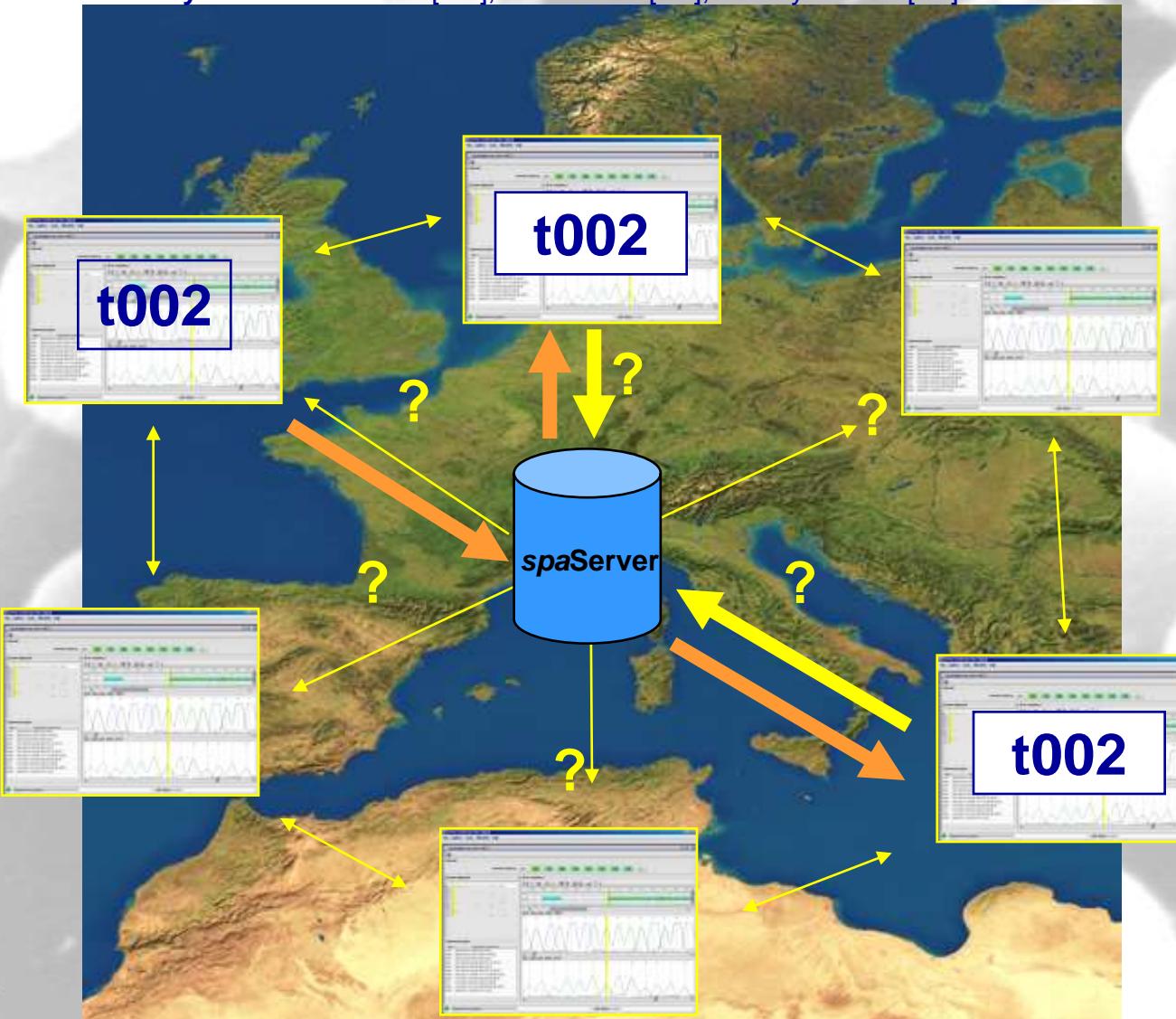
- Data submissions and activities [card of europe]
- The Top20 of Spa-Types in countries [members only]
- Spa distribution in countries [members only]

At the bottom of the page, there is a footer with links: 'Home | Aims | Objectives | Participants | Management Board | Activities | Back-ground | Downloads | Discussion List | Press releases | Publications | Declaration | Rules for procedure | European Guidelines | Cooperation with INICC | Disclaimer'.

Ensuring a common typing language...

Coordinators: A.W. Friedrich [Münster] & W. Witte [Robert Koch Institute]

Advisory board: H. Westh [DK], J. Scheres [NL], W. Hryniwicz [PL] & H. de Lencastre [USA]



Friedrich A.W. et al. 2006. Euro Surveill; 11

de Sousa et al. 2006. J Clin Microbiol; 44

Friedrich A. W. et al. 2008 Euro Surveill; 13

Frénay et al., Eur. J. Clin Microbiol. Infect. Dis. 1996

Harmsen et al., J Clin Microbiol 2003

Strommenger et al J Clin Microbiol 2006

Austria

Belgium

Bulgaria

Canada

China

Croatia

Czech Republic

Denmark

Finland

France

Germany

Greece

Hungary

India

Italy

Island

Latvia

Lebanon, Jordania

Norway

Malaysia

Netherlands

Poland

Portugal

Romania

Slovenia

Sweden

Switzerland

Thailand

Turkey

UK

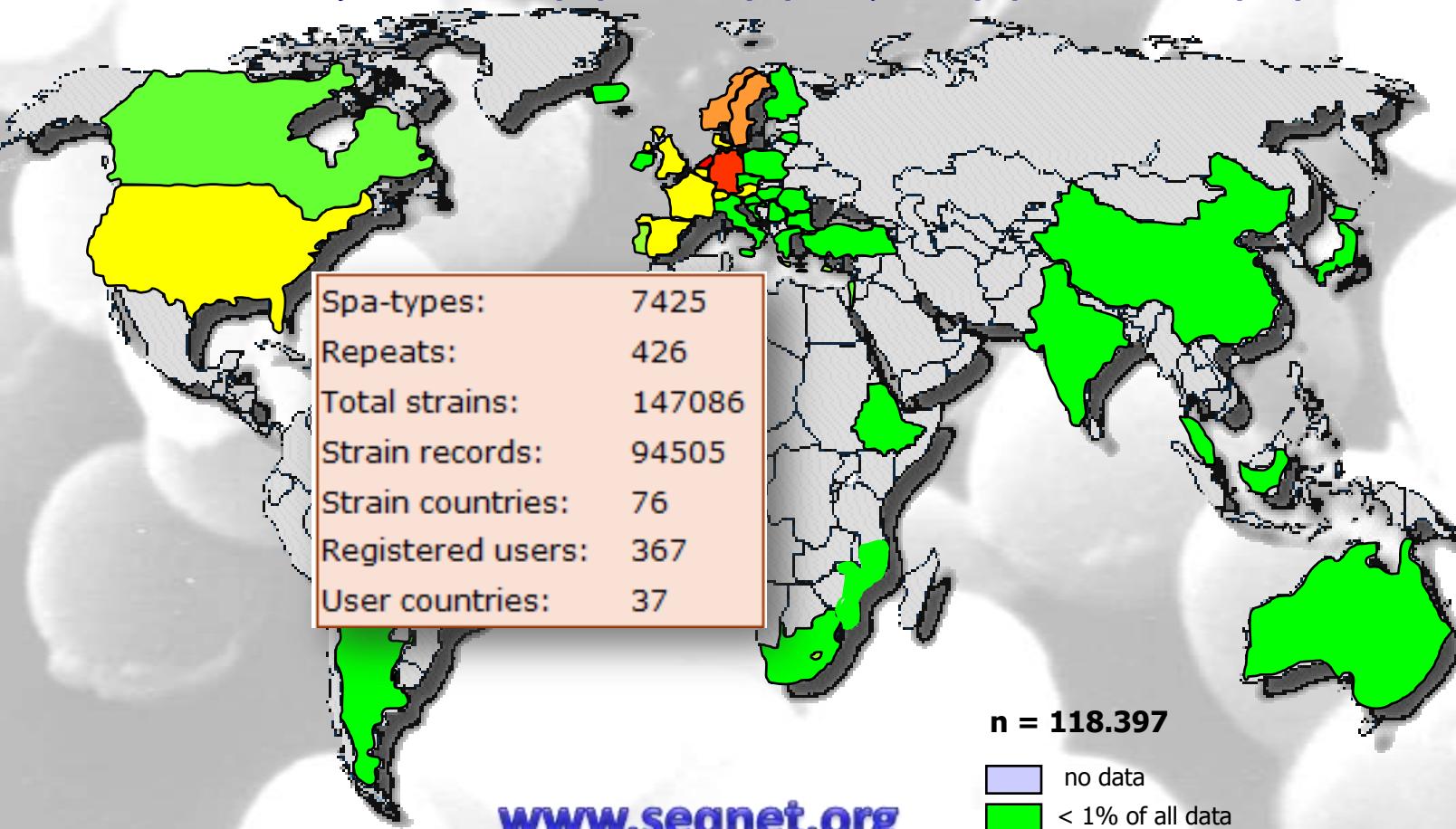
USA

SeqNet.org Network Laboratory-Esperanto for *S. aureus*



Coordinators: A.W. Friedrich [Münster] & W. Witte [Robert Koch Institut]

Advisory board: H. Westh [DK], J. Scheres [NL], W. Hryniewicz [PL] & H. de Lencastre [USA]



www.seqnet.org



spa typing



n = 105443

- National Staphylococcal Reference Laboratory
- University Laboratory , other laboratory
- Veterinarian reference laboratories

■	no data
■	< 1%
■	1% to 4%
■	5% to 10%
■	> 10%

Friedrich et al. 2006 Eurosurveillance
 Friedrich et.al. 2009 Eurosurveillance

Geographic Distribution of *Staphylococcus aureus* Causing Invasive Infections in Europe: A Molecular-Epidemiological Analysis

Hajo Grundmann^{1,2*}, David M. Aanensen³, Cees C. van den Wijngaard¹, Brian G. Spratt³, Dag Harmsen⁴, Alexander W. Friedrich⁵, the European Staphylococcal Reference Laboratory Working Group¹

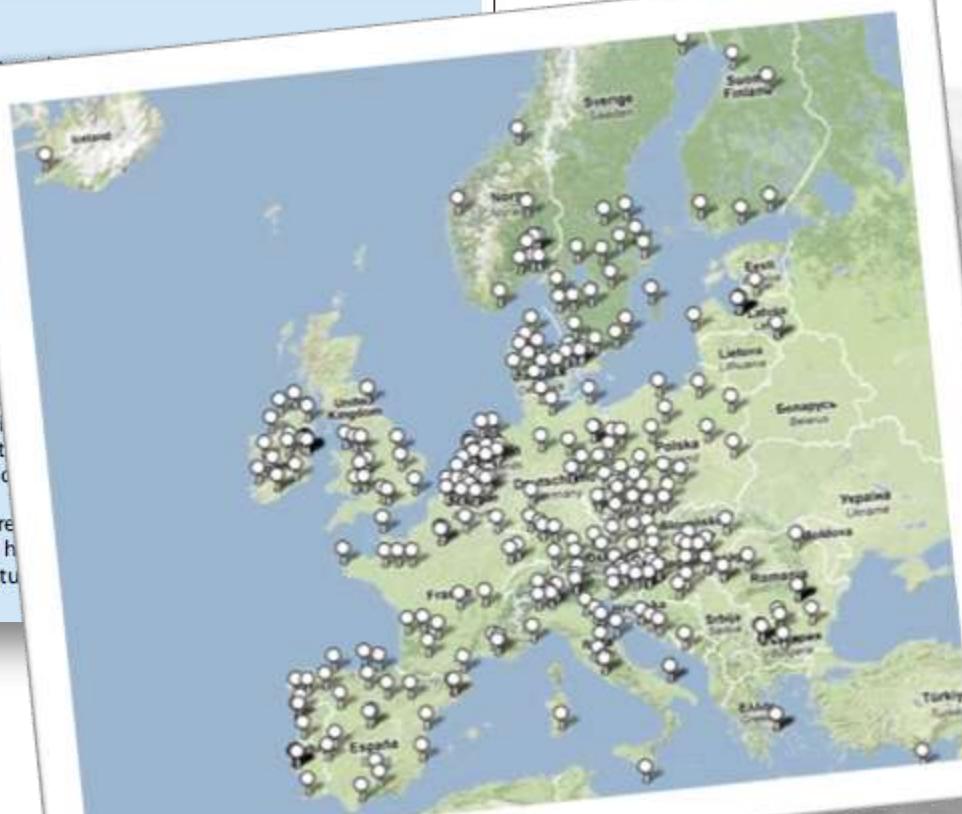
1 National Institute for Public Health and the Environment, Bilthoven, The Netherlands, **2** Department of Medical Microbiology, University Medical Centre, Groningen, The Netherlands, **3** Department of Infectious Disease Epidemiology, Imperial College London, London, United Kingdom, **4** Department of Periodontology, University Hospital Münster, Germany, **5** Institute of Hygiene, University Hospital Münster, Germany

Abstract

Background: *Staphylococcus aureus* is one of the most important human pathogens. Methicillin-resistant *S. aureus* (MRSA) are a major cause of hospital and community-acquired infection. We performed a molecular epidemiological analysis to determine the dominant clones that cause invasive infections in Europe.

Methods and Findings: In each country, staphylococcal reference laboratories collected isolates from a representative number of hospital laboratories to achieve national geo-demographic representation. Isolates were collected from consecutive methicillin-susceptible (MSSA) and MRSA isolates from patients with invasive infections according to a standard protocol. All isolates were sent to the respective national reference laboratory for pulsed-field gel electrophoresis (PFGE) and sequence typing of the variable region of the staphylococcal *spa* gene (*spa* STRT). The data were entered into a central database. Relevant genetic and phenotypic information was assembled for each isolate and entered into a Web-based mapping application. Between September 2006 and February 2008, 15 European countries collected 2,890 MSSA and MRSA isolates from patients with invasive infections. The distribution of *spa* types was found with some prevalent in all European countries. Genetic diversity of MRSA differed considerably between countries with distinct geographical clusters. We provide evidence that a network approach consisting of aggregated data using an interactive mapping tool can provide important information for populations such as early signalling of emerging strains, cross border spread, and transmission.

Conclusions: In contrast to MSSA, MRSA *spa* types have a predominantly regional distribution, which is indicative of the selection and spread of a limited number of clones within health care facilities. Efforts aimed at interrupting the spread within and between health care institutions have been successful and should therefore be strongly encouraged.



General results

Statistics	n ^a	MSSA	MRSA	Total/Overall	p-Value*
Frequency (%)	2,890	1,923 (66.5)	967 (33.5)	2,890 (100%)	—
Median age (IQR)	2,836	63 (46–75)	69 (55–78)	66 (49–76)	<0.0001
Male gender (%)	2,862	1,159 (60.8)	606 (63.3)	1,765 (61.7)	0.2
All-cause mortality after 14 d (%)	1,838	153 (13.2)	141(20.8)	294 (16.0)	<0.0001
Hospital acquisition (%)	2,322	777 (51.6)	585 (71.7)	1,362 (58.7)	<0.0001
N spa types	2,850	565	155	660 ^b	—
N not typeable	2,850	27 (1.4)	13 (1.3)	40 (1.4)	0.9
Index of diversity (95% CI)	2,850	0.985 (0.983–0.987)	0.940 (0.933–0.947)	0.983 (0.982–0.984)	<0.05 ^c
Mean distance in kilometres between laboratories that isolated identical spa types (95% CI)	1,614 ^d	1,046.2 (1109.5–983.0)	786.8 (975.7–597.9)	—	0.03 ^d

*p-Value for the comparison of MSSA versus MRSA.

^aNumber of isolates for which data were available.

^bTotal number of spa types includes 60 spa types that contain both MSSA and MRSA.

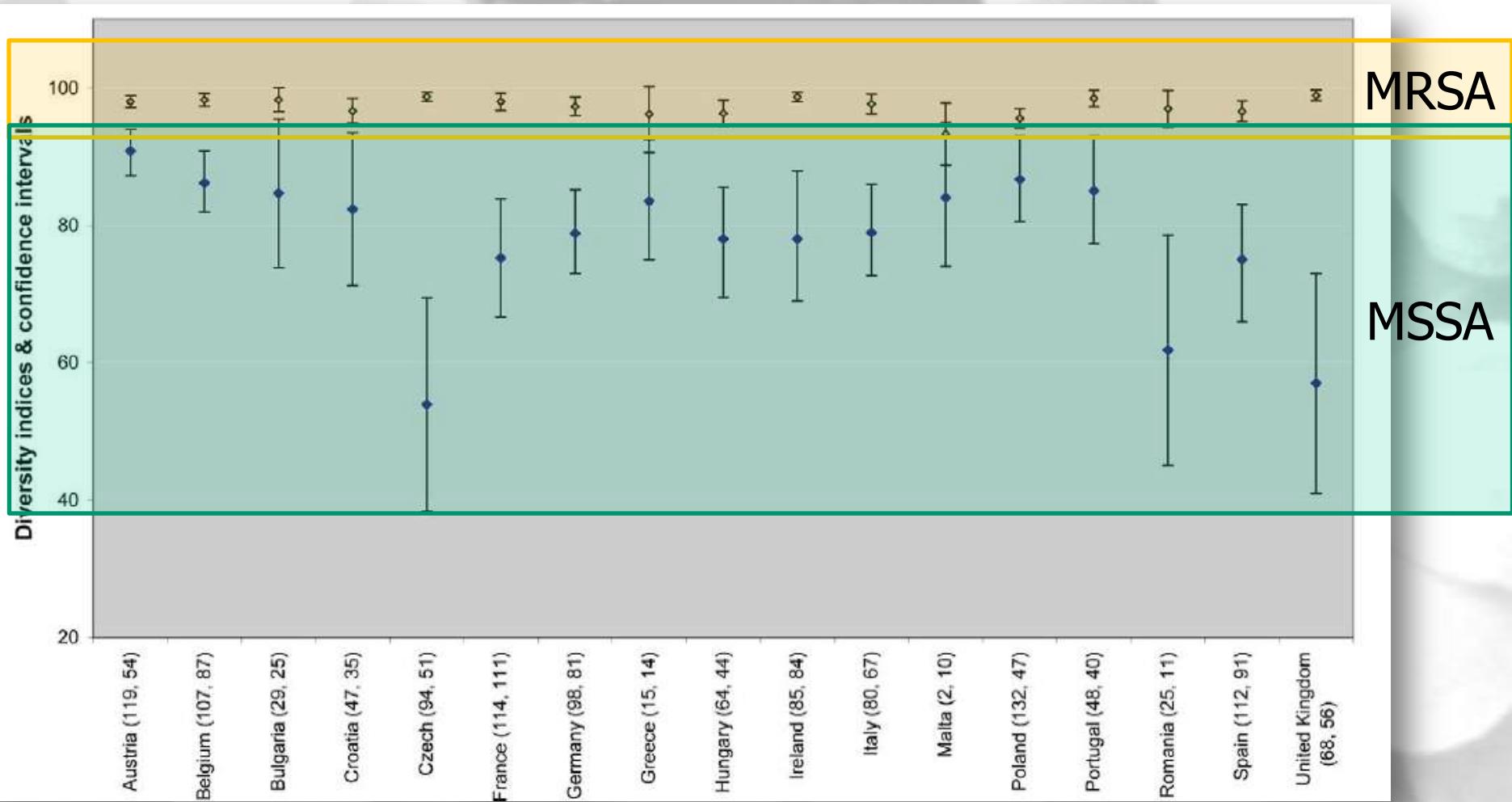
^cDeduced from non-overlapping 95% confidence intervals.

^dIncludes only MRSA and MSSA with more than ten isolates per spa type.

IQR, interquartile range.

doi:10.1371/journal.pmed.1000215.t002

MRSA vs. MSSA



S. aureus blood culture isolates in Europe

Rank	MSSA		Frequency	Percent	Cumulative Percent	Rank	MRSA		Frequency	Percent	Cumulative Percent
	spa	Type					spa	Type			
		MLST						MLST			
1	t002	ST-5 ^a , S-231 ^a	93	4.8	4.8	1	t032	ST-22 ^a	140	14.5	14.4
2	t084	ST-15 ^a (ST-18)	89	4.6	9.5	2	t008	ST-8 ^a (ST-247, ST-250, ST-254)	120	12.4	26.8
3	t015	ST-45 ^a	84	4.4	13.8	3	t041	ST-111 ^a , ST-228 ^a	72	7.4	34.2
4	t091	ST-7 ^a	82	4.3	18.1	4	t003	(ST-5) ST-225 ^a	71	7.3	41.6
5	t012	ST-30 ^a	77	4.0	22.1	5	t002	ST-5 ^a , ST-231 ^a	62	6.4	48.1
6	t127	ST-1 ^a	57	3.0	25.1	6	t067	ST-5 ^a , ST-125 ^a	50	5.2	53.3
7	t008	ST-8 ^a (ST-247, ST-250, ST-254)	55	2.9	27.9	7	t001	(ST-5, ST-222) ST-228 ^a	30	3.1	56.4
8	t021	ST-30 ^a (ST-33, ST-55)	49	2.5	30.5	8	t037	ST-239 ^a (ST-240), ST-241 ^a	27	2.8	59.2
9	t005	ST-22 ^a (ST-23, ST-60)	42	2.2	32.7	9	t030	ST-239 ^a (ST-246)	20	2.1	61.2
10	t026	(ST-45, ST-47)	27	1.4	34.1	10	t024	ST-8 ^a	14	1.4	62.7
11	t065	(ST-45, ST-46)	26	1.4	35.4	11	t190	ST-8 ^a	14	1.4	64.1
12	t160	(ST-12, ST-13)	26	1.4	36.8	12	t515	ST-22 ^a	12	1.3	65.5
13	t056	(ST-101)	25	1.3	38.1	13	t038	ST-45 ^a	12	1.2	66.7
14	t050	ST-45 ^a	21	1.1	39.2	14	t022	ST-22 ^a	11	1.1	67.8
15	t078	(ST-26)	21	1.1	40.2	15	t740	ST-45 ^a	11	1.1	69.0
16	t164	(ST-20)	19	1.0	41.2	16	t012	ST-30 ^a	9	0.9	69.9
17	t346	(ST-15, ST-620)	18	0.9	42.2	17	t015	ST-45 ^a	9	0.9	70.8
18	t024	ST-8 ^a	17	0.9	43.1	18	t044	ST-80 ^a	9	0.9	71.8
19	t230	ST-45 ^a	17	0.9	43.9	19	t045	ST-5 ^a (ST-225)	8	0.8	72.6
20	t166	(ST-34)	16	0.8	44.8	20	t127	ST-1 ^a	8	0.8	73.4
—	Other	—	1,062	55.2	100.0	—	Other	—	258	26.6	100.0
Total	—	—	1,923	100	—	—	—	—	967	100	—

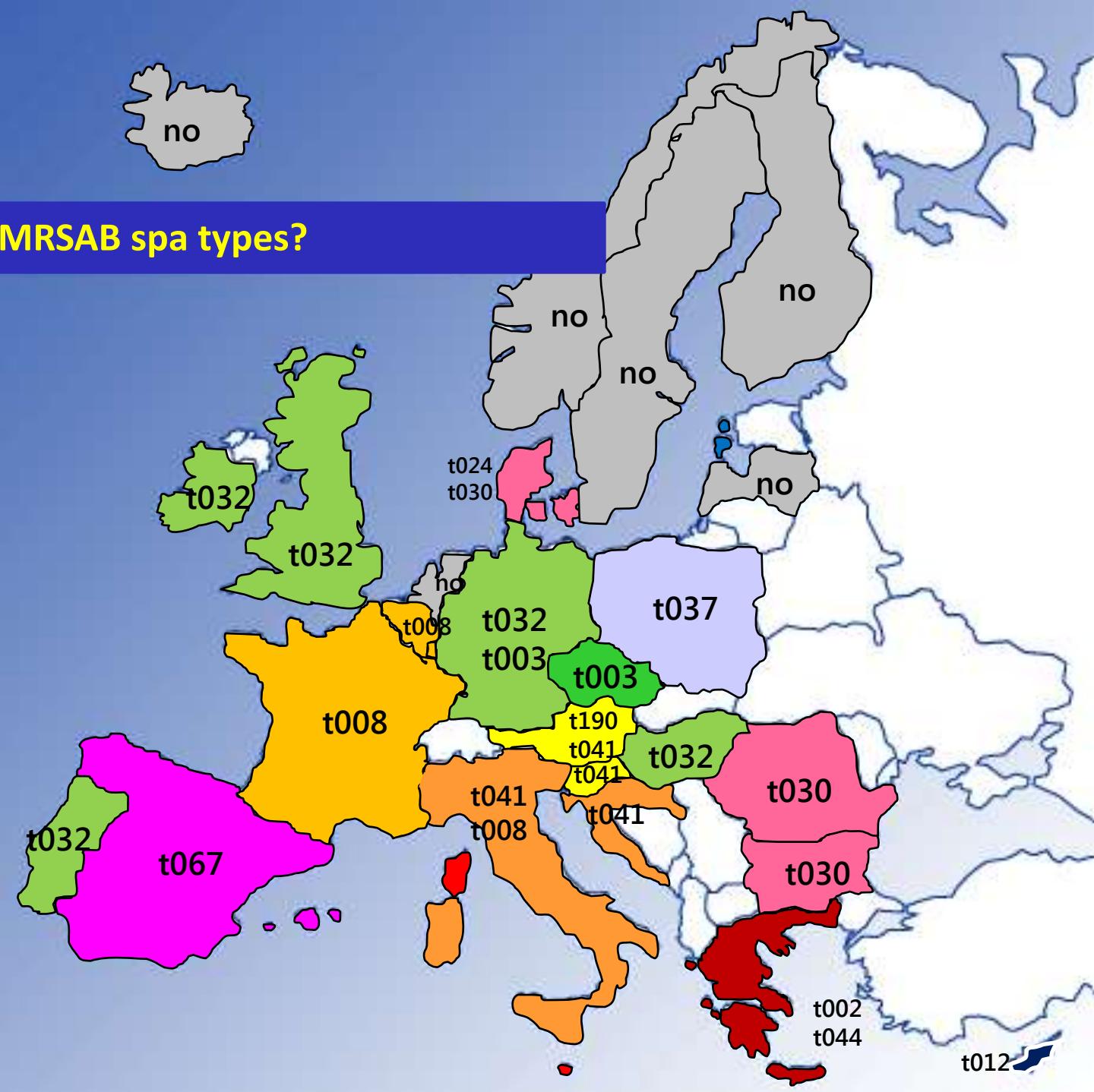
STs in parentheses are those associated with the spa type in the SeqNet.org Spa typing data base.

^aMLST as determined by SRLs.

doi:10.1371/journal.pmed.1000215.t003

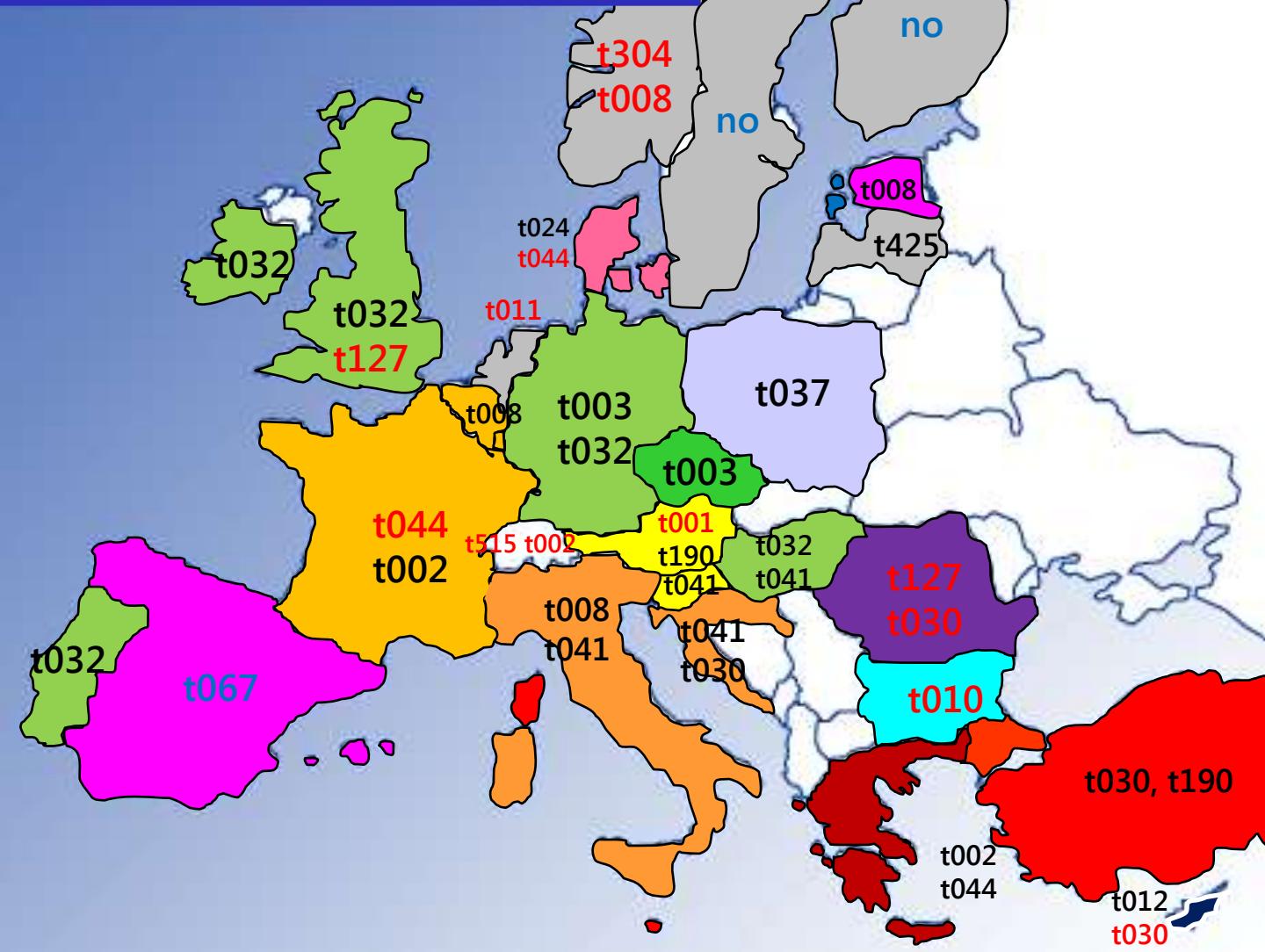


MRSAB spa types?



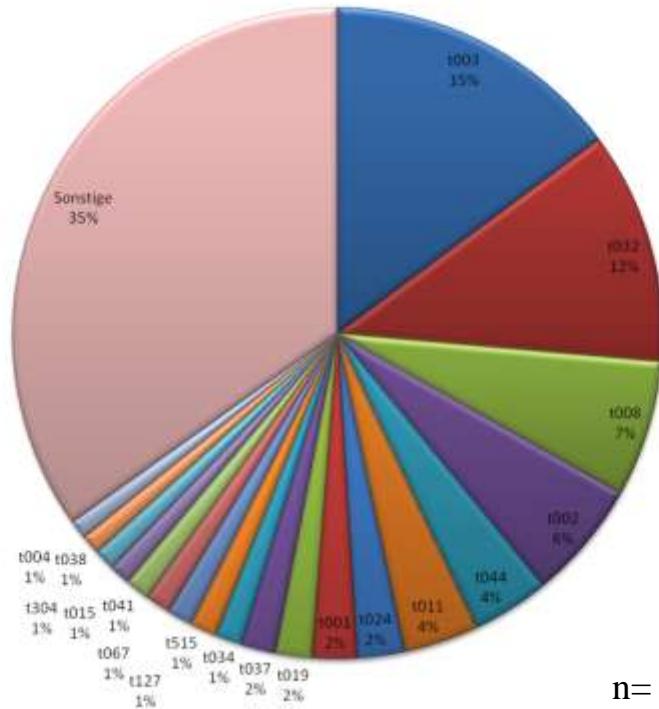


MRSA spa types on the spa server

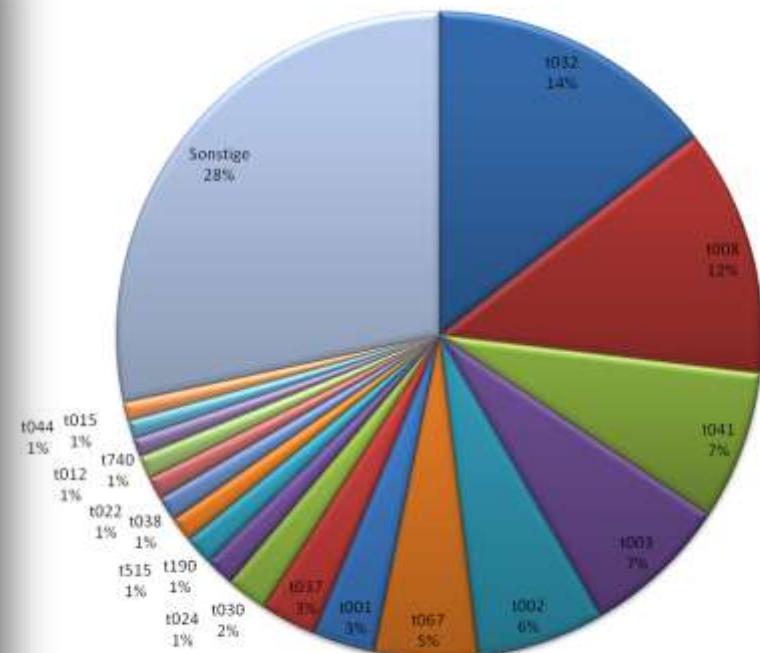


TOP 30 MRSA in Europe

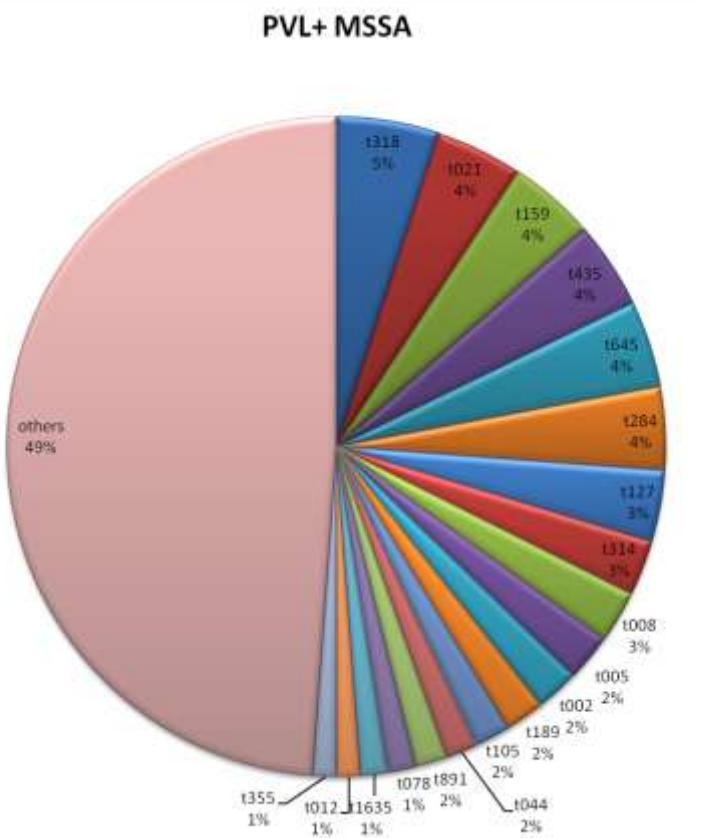
MRSA on the spa server



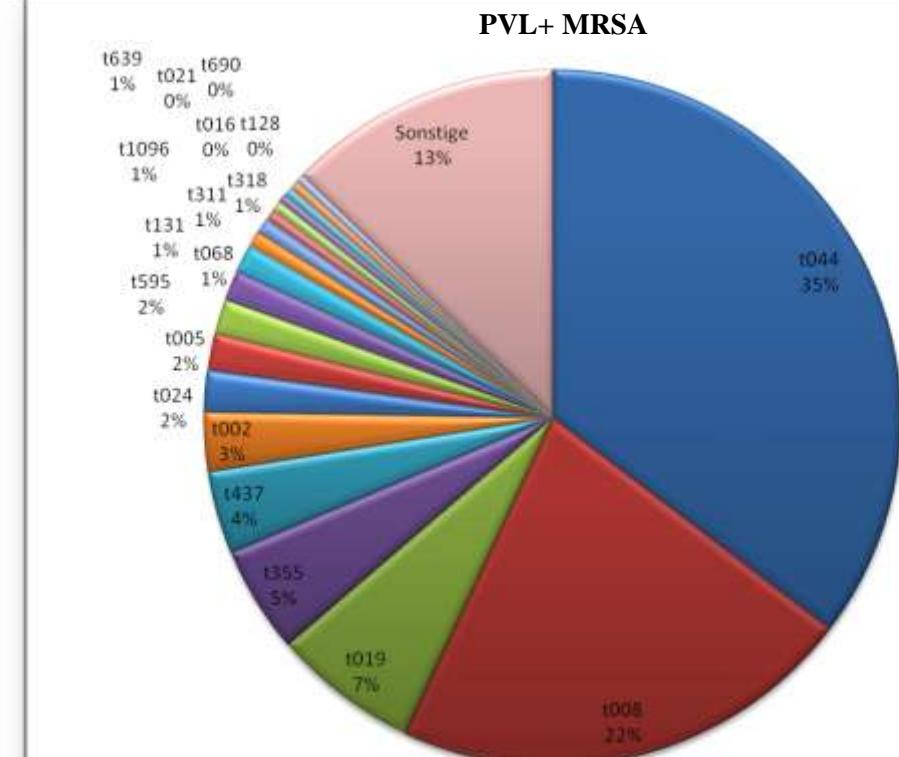
MRSAB from SRL-study



PVL-positive CA-MRSA

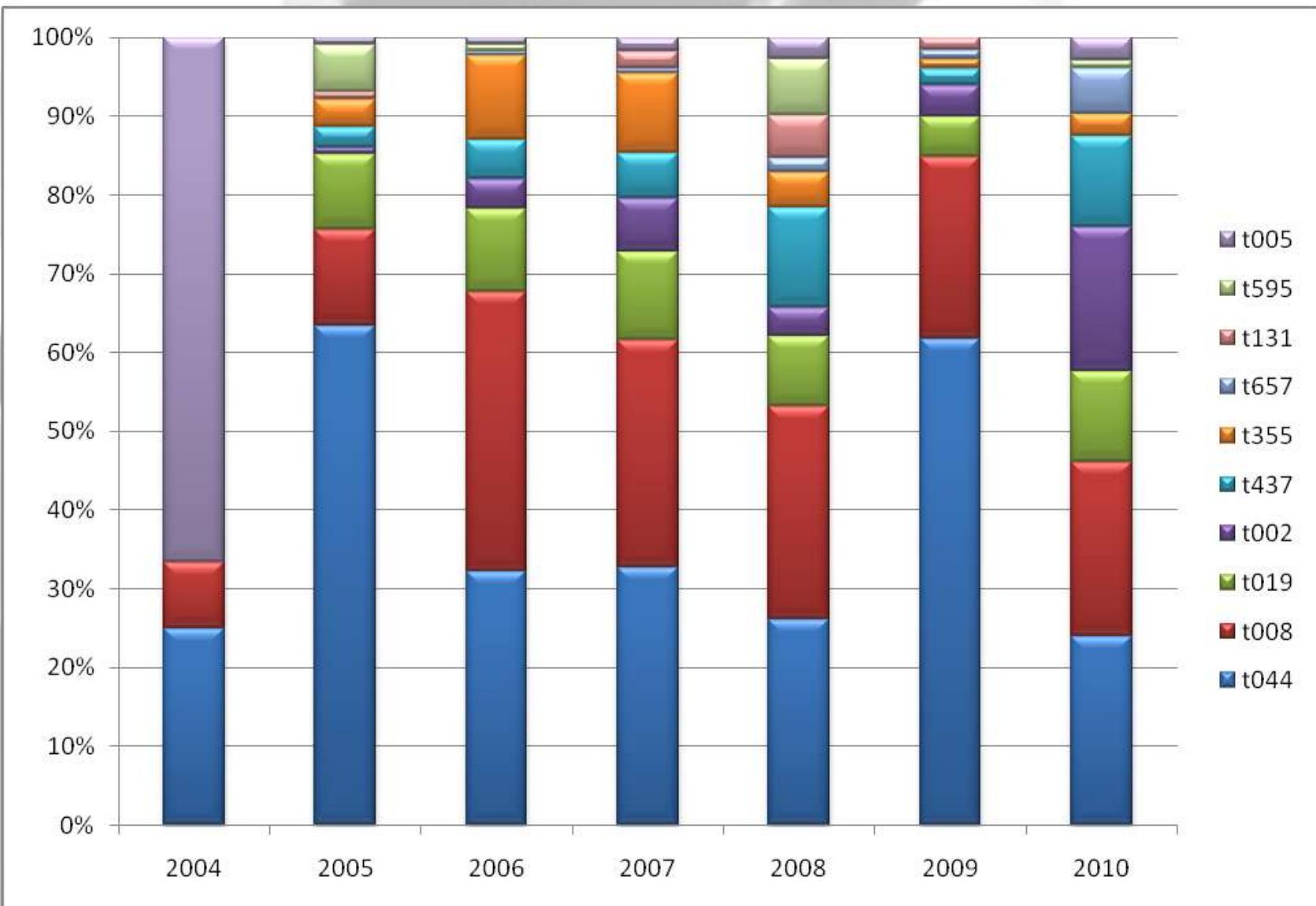


194 different *spa* types



75 different *spa* types

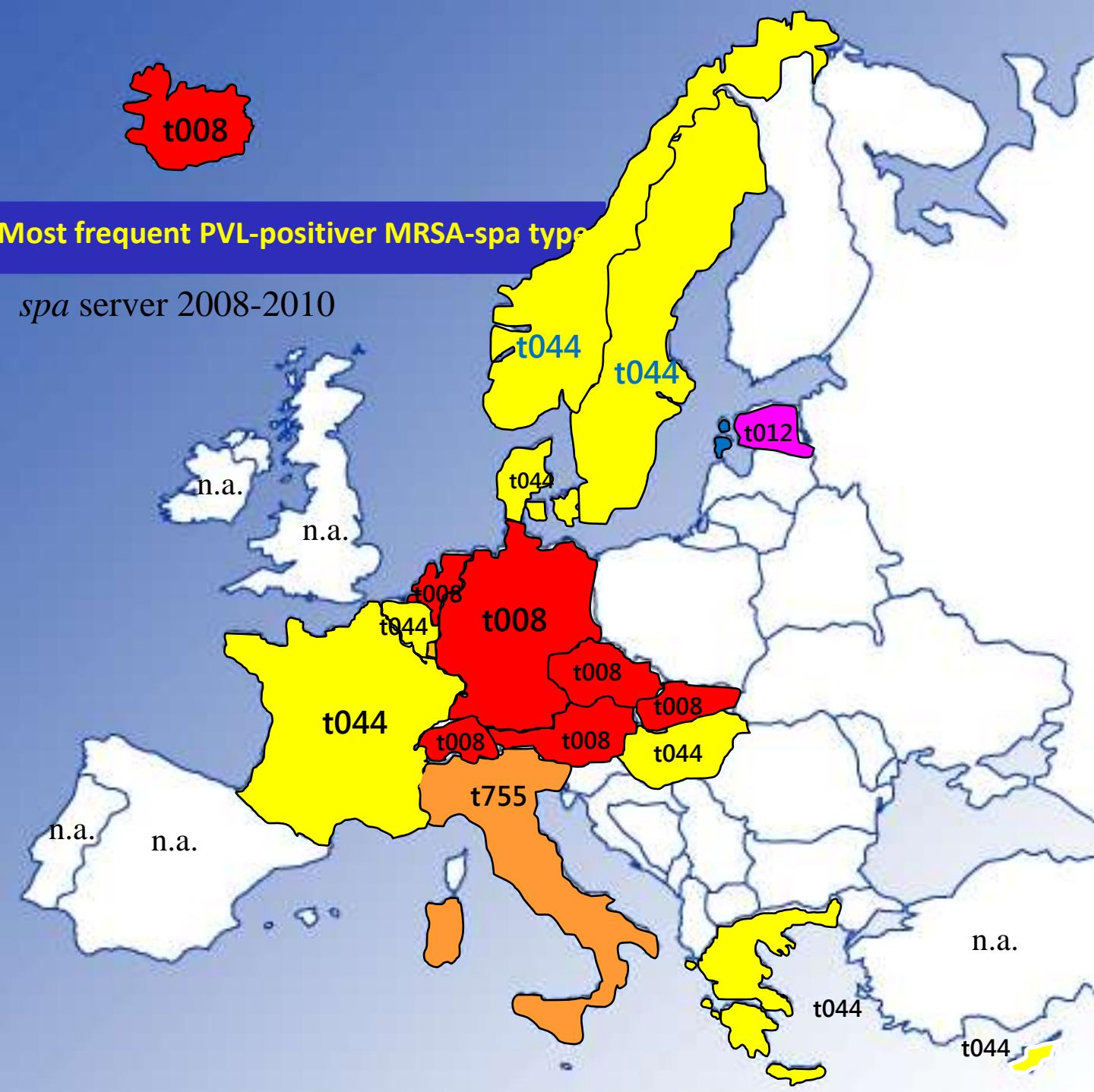
Change over time PVL+ MRSA (n= 2035)





Most frequent PVL-positiver MRSA-spa type

spa server 2008-2010



Regional Cluster

Regional Cluster Number	spa Type	spa Complex ^a	ST ^b	Window Centre	Window Radius (km)	Countries reporting Clustered spa Type within Window			Log Likelihood Ratio	n MRSA among Observed Isolates	Percent MRSA among Observed Isolates
						n Isolates Observed	n Isolates Expected ^c				
1	t067	2	5 & 125	Alicante, Spain	716	ES, FR	55	4.7	126.9	46	84
2	t041	1	228	Split, Croatia	522	AT, HR, HU, SI, IT	59	10.6	84.76	59	100
3	t032	32	22	Belfast, Northern Ireland (UK)	596	IE, UK	77	15.9	84.74	75	97
4	t003	45	225	Leipzig, Germany	386	CZ, DE, PL	58	10.0	82.26	54	93
5	t008	8	8	Perpignan, France	931	AT, BE, DE, ES, FR, HR, PT, SI	119	45.5	72.96	105	88
6	t740	740	45	Goes, Netherlands	81	BE, NL	15	0.6	50.36	11	73
7	t030	12	239	Pleven, Bulgaria	331	BG, RO	16	0.7	45.24	15	94
8	t037	12	239	Plock, Poland	330	PL	21	2.2	36.02	18	86
9	t038	740	45	Wilrijk, Belgium	92	BE	12	0.8	32.71	12	100
10	t190	190	8	St Pölten, Austria	56	AT	10	0.3	29.63	9	90
11	t001	1	228	Sibenik, Croatia	885	AT, DE, HU, IT, MT	29	10.2	27.55	29	100
12	t435	435	427	Daugavpils, Latvia	189	LV	8	0.2	25.82	0	0
13	t425	239	368	Riga, Latvia	0	LV	5	0.1	22.76	5	100
14	t127	127	1	Inowroclaw, Poland	131	PL	12	0.9	22.65	0	0
15	t032	32	22	Berlin, Germany	128	DE	19	3.2	22.2	19	100
16	t015	15	45	Lomza, Poland	581	CZ, LV, PL	35	11	20.72	7	20
17	t091	91	7	Ried, Austria	771	AT, BE, CZ, DE, FR, HR, HU, IT, NL, PL, SI	71	42.8	20.68	4	6
18	t777	777	5	Laval, France	277	FR	7	0.3	20.61	7	100
19	t081	78	25	Maków Mazowiecki, Poland	0	PL	4	0.0	19.37	3	75
20	t515	32	22	Mullingar, Ireland	196	IE, Northern Ireland (UK)	9	0.7	18.94	9	100
21	t002	2	5, 231	Coimbra, Portugal	484	ES, PT	33	10.6	18.37	20	61
22	t230	728	45	Hillerød, Denmark	235	DE, DK, SE	11	1.4	17.19	0	0
23	t044	44	80	Nicosia, Cyprus	916	GR, CY	6	0.2	17.09	6	100
24	t2054	8	8	St Mande, France	226	BE, FR4	5	0.2	16.64	3	60
25	t062	singleton	5	Szolnok, Hungary	145	HU	6	0.3	15.24	6	100
Total	—	—	—	—	368 (mean)	339	702	173.1	—	522	74

SatScan Analysis for border independent cluster



Map Satellite Hybrid Terrain

Total-62 isolates MRSA/MSSA

Mort 14d

Age Distribution (%)

Age Group	Percentage
80+	10
70-79	10
60-69	10
50-59	10
40-49	10
30-39	10
20-29	10
10-19	10
0-9	10
<1	10

Legend: Lab Selected (blue dot), MSSA only (green dot), MSSA&MRSA (yellow dot), MRSA only (red dot).

Map data ©2009 Basksoft, A&D Geodata Consulting, PPK, Tele Atlas, TomTom, Esri, DigitalGlobe, TerraMetrics, Google.

ES056 spa Type Summary

Click to return to European view

No. isolates	spa Type	Pattern	other laboratories (no. isolates elsewhere with same spa Type)	Action
4	1067	r29r23r17r34r17r29r17r12r17	25(58)	click to view on map
1	1073	r04r21r12r41r20r17r12r12r17	15(20)	click to view on map
1	1630	r08r16r02r16r34r17r34r16r34	6(6)	click to view on map
1	1012	r15r12r19r02r16r02r25r17r24r24r24	68(85)	click to view on map
1	r129	r15r12r24	0	-
1	11051	r15r25r02r12r16r02r15r02r25r17r24r24r24	0	-
1	1837	r26r17r20r17r12r17	0	-

Contact Us | Imperial College London | Funded by The Wellcome Trust



22% of European population lives close borders

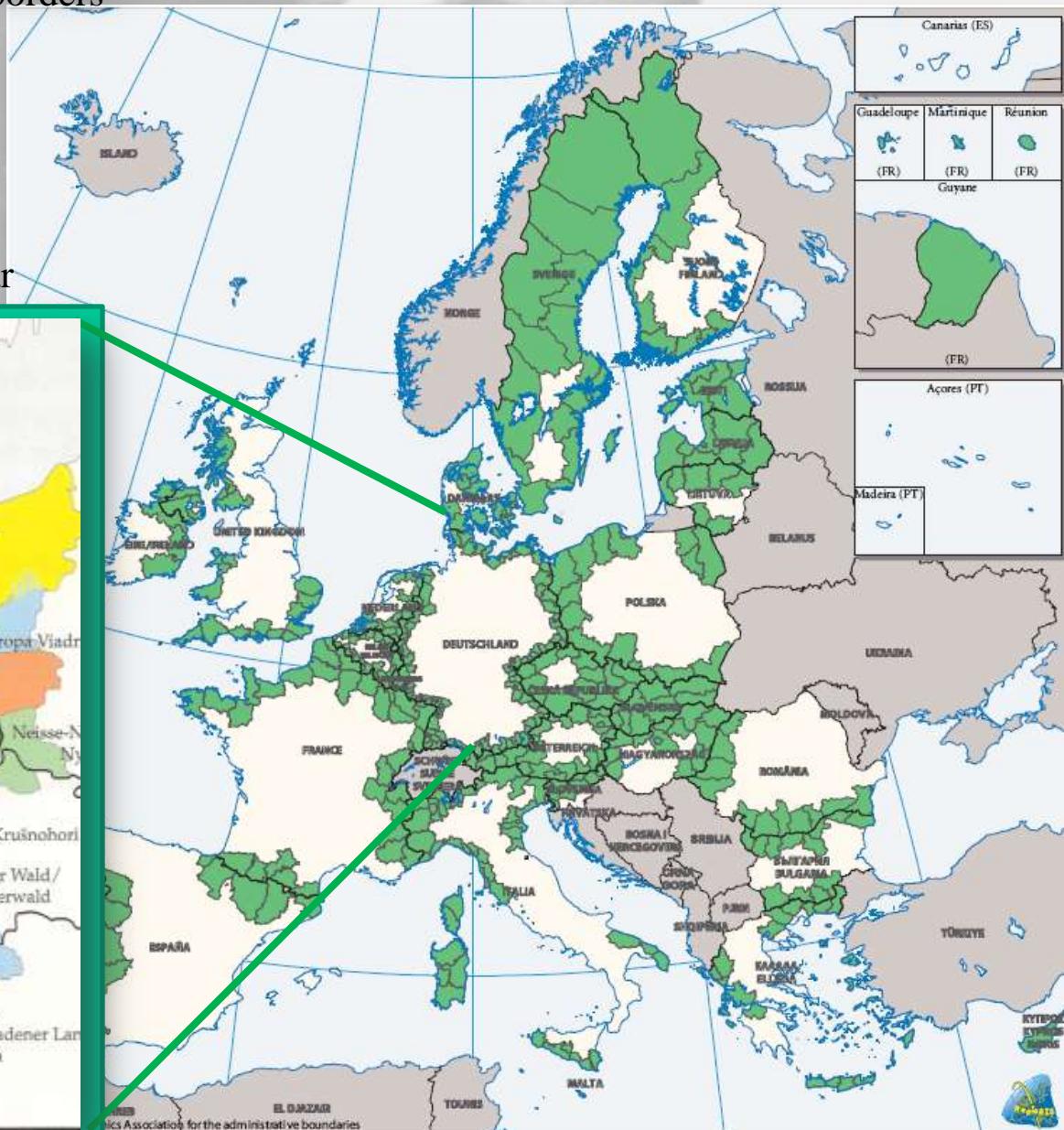
Dutch-German Euregios

Since 1958

12,4 Mio inhabitants

12.000 crossborder commuters/day

500 to 30.000 crossborder patient per year



INTERREG
Deutschland
Nederland

 **INTERREG - Grenzregionen gestalten Europa**
Europäischer Fonds für Regionale Entwicklung der Europäischen Union

INTERREG - Grensregio's bouwen aan Europa
Europees Fonds voor Regionale Ontwikkeling van de Europese Unie



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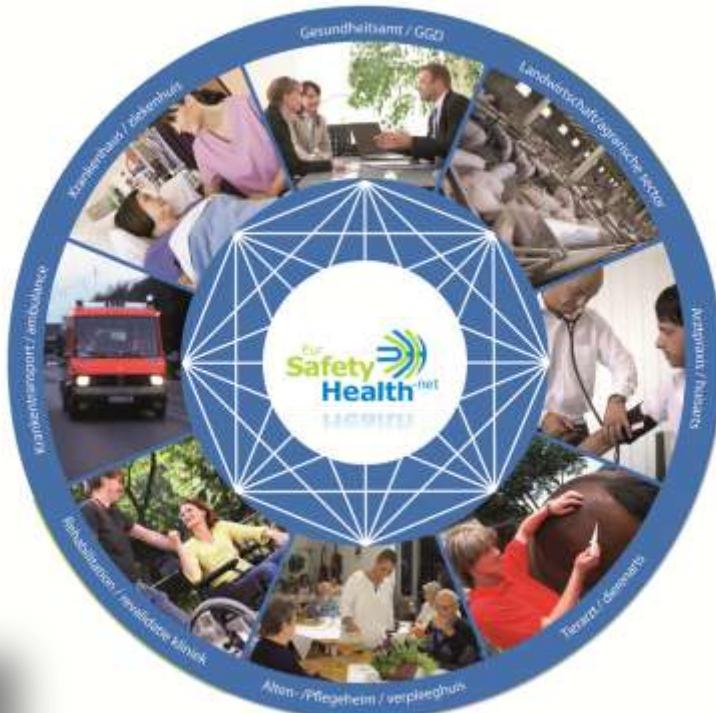
Belique-België

Euregio-Projekte:



EURSAFETY HEALTH-NET

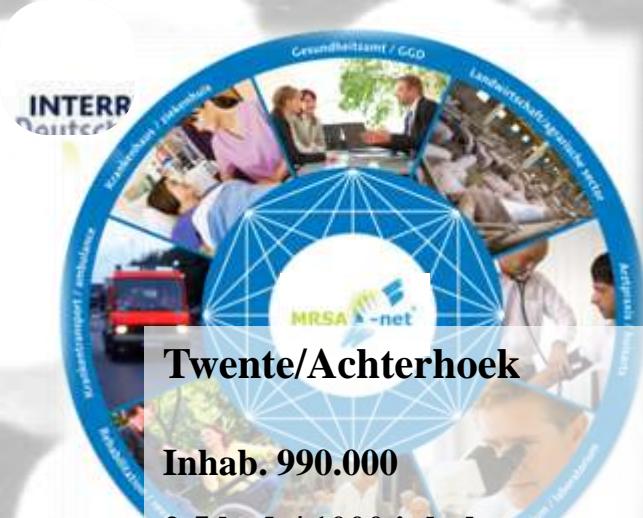
Euregionales Netzwerk
für Patientensicherheit
und Infektionsprävention



www.eursafety.eu

www.mrsa-net.eu

EUREGIO MRSA-net (2009)



Twente/Achterhoek

Inhab. 990.000

2,5 beds/ 1000 inhab

36 GPs/100.000 inhab

Microbiologists: 2,9 per 1000 patients

MRSA : 78 (7,8/100.000 inhab)

0,1 per 100 admissions

MRSA/ *S. aureus*: 1%

MRSAB: 1 (0,1/100.000)

EUREGIO mrsa-net Twente/Münsterland



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INTERREG - Grensregio's bouwen aan Europa
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- Crossborder network of 44 German and Dutch hospitals
- Integrated network with GPs
- Common quality standards (search&follow)

- Prevalence screening of 23566 (D) and 3544 (NL) patients
- Each new MRSA isolate from all patients is spa typed by hospitals
- Hospital and laboratory data collected on EUREGIO server

Münsterland

Inhab: 1.566.000

6,9 beds/1000 Einw.

146 GPs+Specialists/100.000 inhab

Microbiologists: 1,0 per 1000 patients

MRSA: 2953 (189/100.000 inhab)

1,6/100 admissions

MRSA/ *S. aureus*: 3%-30%

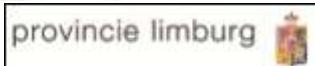
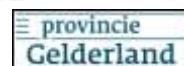
MRSAB: 41 (2,6/100.000)



Ministerium für Wirtschaft,
Mittelstand und Energie
des Landes Nordrhein-Westfalen



Friedrich et al. 2008. Eurosurveillance
Köck et al. 2009, JHI
Böcher et al. JAC 2010



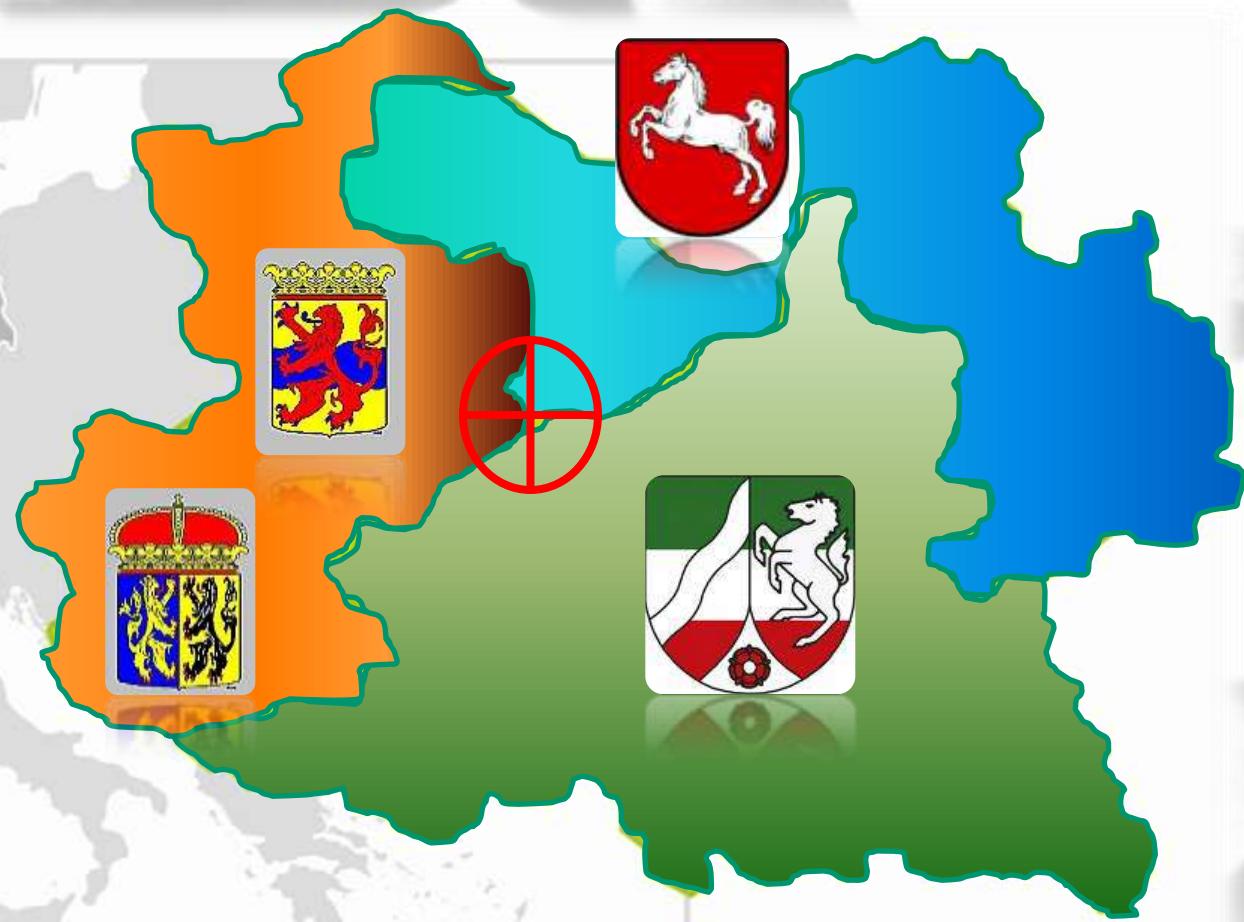


The Netherlands



Germany

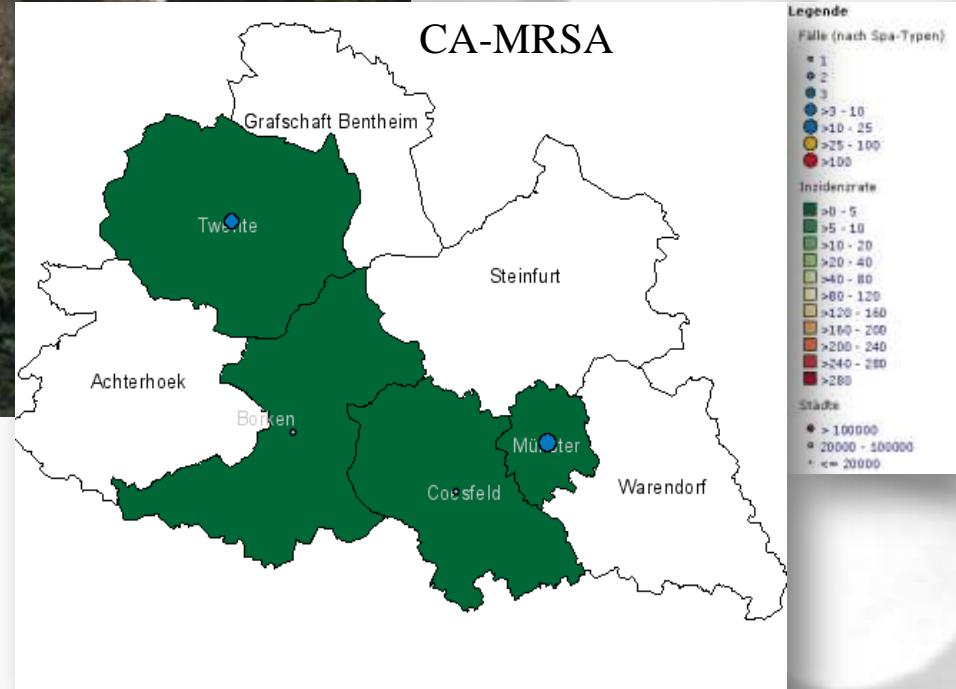
EUREGIO mrsa-net Twente/Münsterland



Friedrich et al. 2008. Eurosurveillance

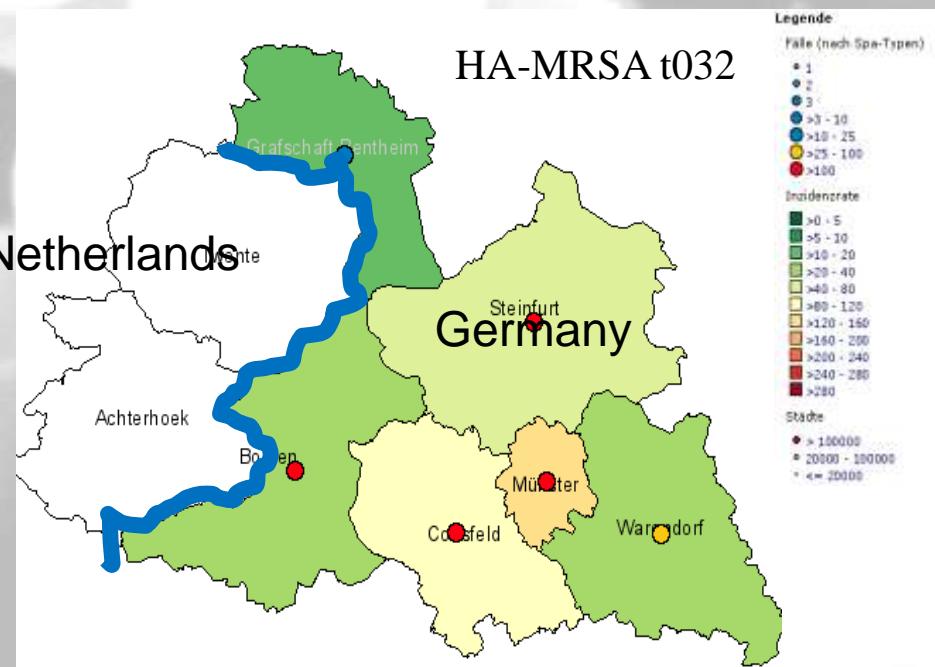


Dreiländereck: NL, NRW, Nds



Real-time Surveillance of human MRSA isolates in the EUREGIO

Healthcare-associated (HA-) MRSA



Livestock-associated (LA-)MRSA



- HA-MRSA respect the borders
- **Patients coming from German hospitals are put into isolation**
- **HA-MRSA not transmitted via social contacts**

Incidence of *spa* types indicative for MLST CC398 ([t011](#), [t034](#), [t108](#), [t567](#), [t571](#), [t588](#), [t753](#), [t898](#), [t899](#), [t1184](#), [t125](#), [t1451](#), [t1456](#), [t1457](#), [t2123](#), [t2330](#), [t2383](#), [t2582](#), [t3013](#)):
Equal incidence of LA-MRSA genotypes on both sides of the border

Prevalence screening in the EUREGIO (n= 23,566)

- Admission screening and risk factor analysis of all patients in all 44 hospitals in the EUREGIO
- All MRSA spa typed
- 1,6 / 100 patients colonized at admission
- 33% of all patients with MRSA risk factor
- 89% of patients with MRSA show up with classical HA risk factors

Risk factor	MRSA (n=354)	No MRSA (n=20836)	Beta Coefficient	OR	95% CI	P
home	89	2742	0.29	1.3	1.0 - 1.8	<0.001

22% of MRSA carriers without classical risk factors

home							
Antibiotics	89	2742	0.29	1.3	1.0 - 1.8	<0.001	0.0776
Haemodialysis	3	179	-§	-§	-§ -	-	-§
Burning trauma	0	39	-§	-§	-§ -§	-§	-§
Chron. wounds	48	415	4.0	54.5	26.2 - 113.6	<0.001	
Katheter	49	541	2.09	8.1	4.1 - 15.9	<0.001	
Chron. nursing	55	662	1.22	3.4	1.8 - 6.4	<0.001	

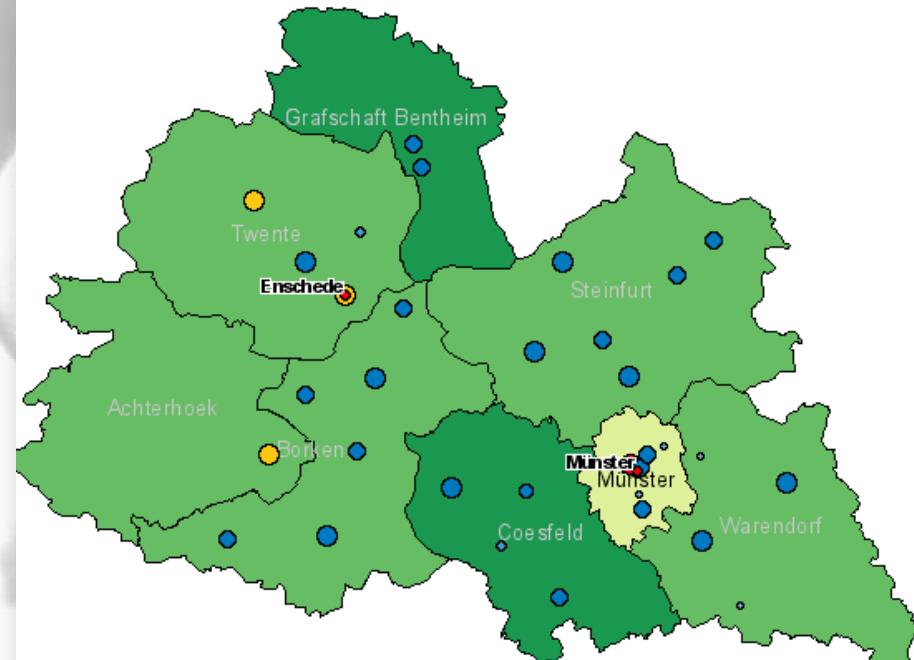
Real-time Surveillance of human MRSA isolates in the EUREGIO

Healthcare-associated (HA-) MRSA



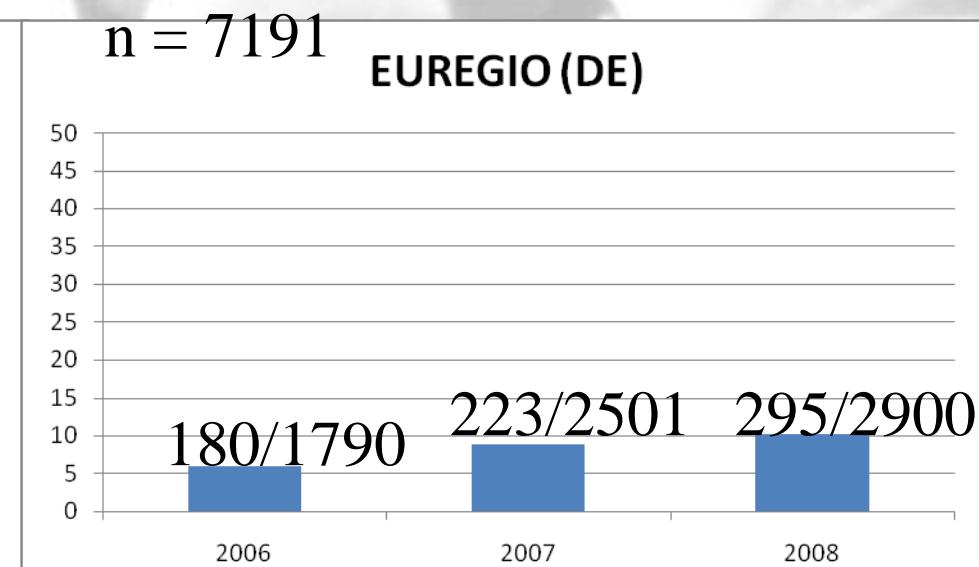
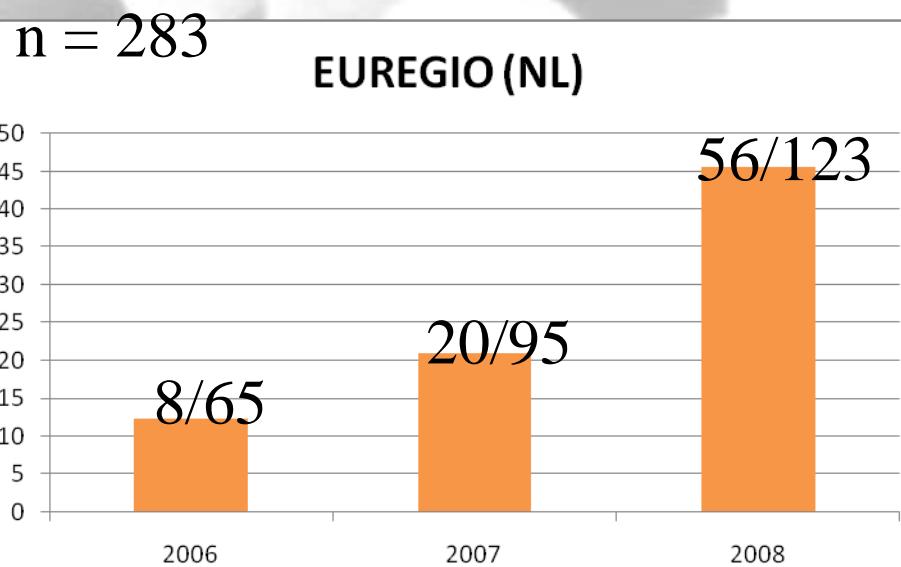
Incidence of all MRSA cases per 100.000 inhabitants:
HA-MRSA respect the borders

Livestock-associated (LA-)MRSA



Incidence of *spa* types indicative for MLST CC398 (**t011, t034, t108, t567, t571, t588, t753, t898, t899, t1184, t125, t1451, t1456, t1457, t2123, t2330, t2383, t2582, t3013**):
Equal incidence of LA-MRSA genotypes on both sides of the border

LA-MRSA in human MRSA-isolates of the EUREGIO



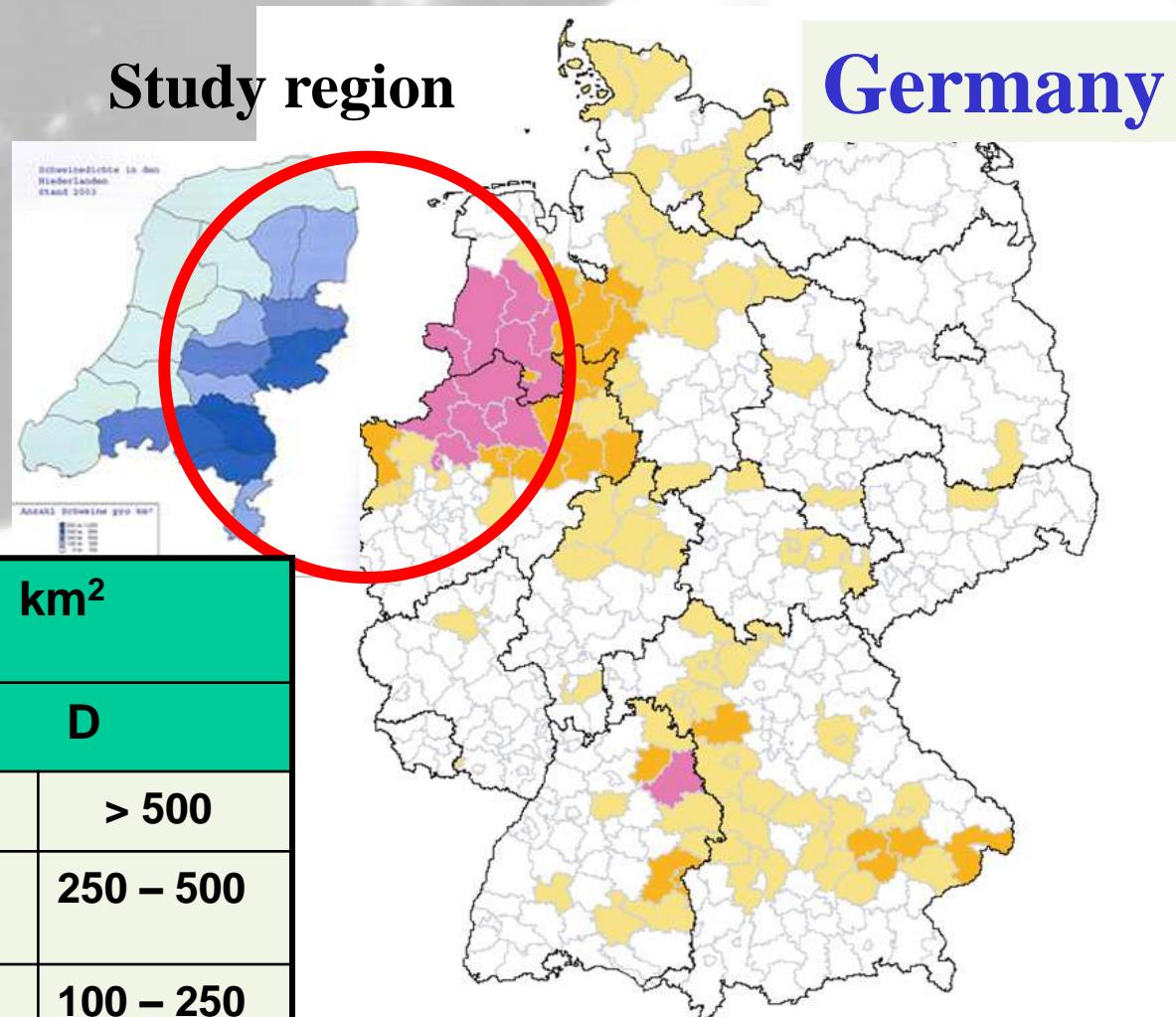
Pig density in the Dutch-German border region

Netherlands

Pigs / 100 ha or 1 km ²			
NL		D	
> 500	■	■	> 500
300 – 500	■	■	250 – 500
100 – 300	■	■	100 – 250
0 – 100	■	□	0 – 100

Study region

Germany



Trading networks are important for spreading

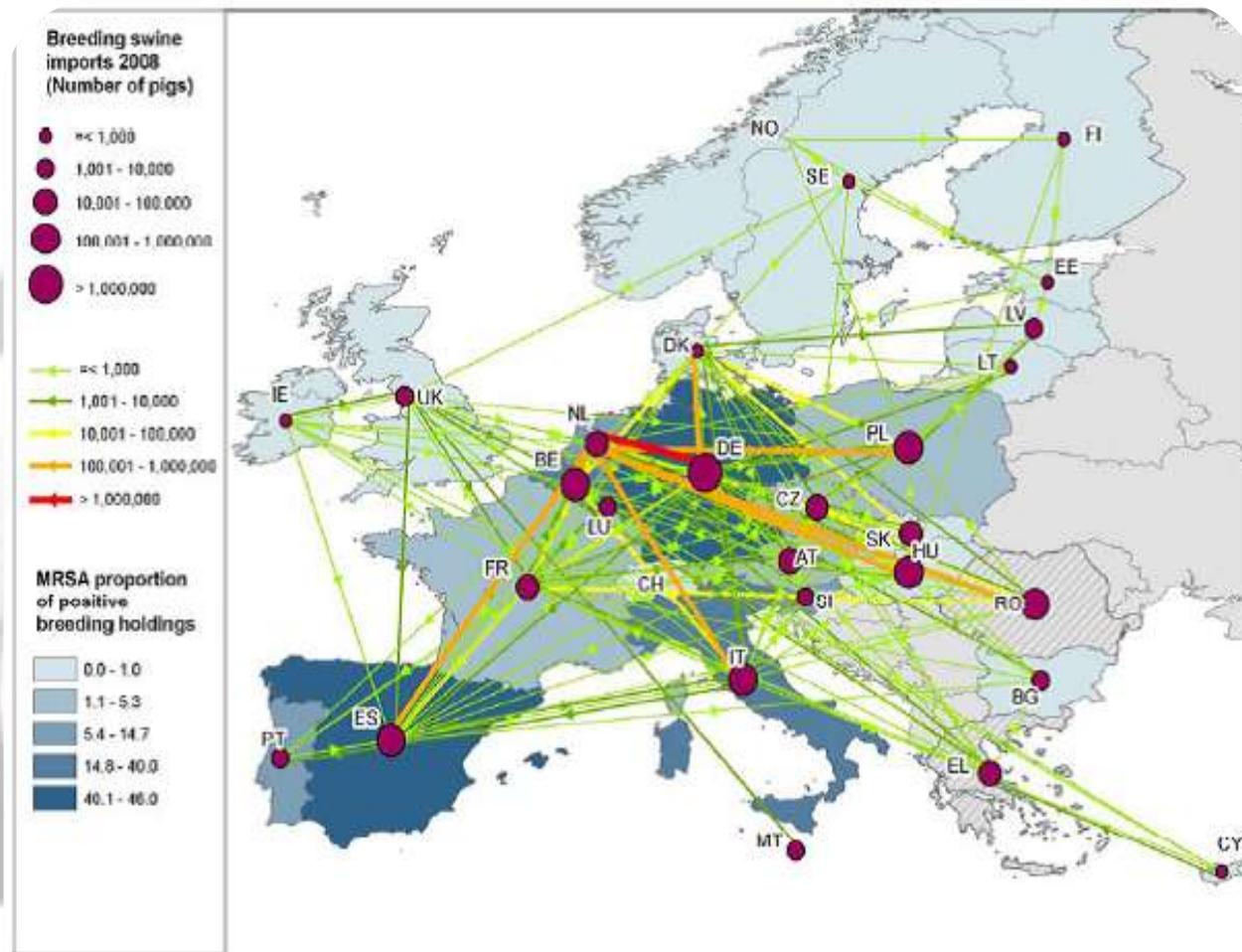


Figure 14: Prevalence of MRSA-positive breeding holdings in 2008 (EFSA, 2009) and within community trades of breeding pigs in 2008^{(a),15}

EFSA Scientific Report 2009, 8(1), 11. Prevalence of MRSA-positive breeding holdings in 2008^{(a),15}

Case-control study: Risk factors for the acquisition of LA-MRSA

Risk factor	Number of patients				Univariate analysis <i>P</i> value ^a	Final logistic regression model Odds ratio (CI 95%) ^b	<i>P</i> value ^b			
	Case group		Control group							
	Yes	No	Yes	No						
Residence in a nursing home	0	100	2	98	0.497	NA	NA			
Prescribing of antibiotics ^c	35	65	39	61	0.7	NA	NA			
Haemodialysis	0	100	2	98	0.3	NA	NA			
Male gender	77	23	62	38	0.03	NA	NA			
Age 0–2 years	3	97	2	98	1.0	NA	NA			
Age 3–18 years	9	91	2	98	0.0588	9.629 (1.374–67.476)	0.023			
Age 19–65 years	61	39	60	40	0.885	NA	NA			
Age 66–80 years	24	76	33	67	0.159	NA	NA			
Age >81 years	3	97	3	97	1.0	NA	NA			
Indwelling devices	12	88	28	72	0.008	0.327 (0.105–1.024)	0.055			
Hospitalization ^c	46	54	70	30	<0.001	NA	NA			
Contact with human MRSA carriers	8	92	13	87	0.356	NA	NA			
Chronic need for nursing care	3	97	23	77	<0.001	0.065 (0.008–0.524)	0.010			
Skin lesions	5	95	23	77	0.001	0.247 (0.060–1.009)	0.052			
Contact with pets	78	22	48	52	<0.001	NA	NA			
Contact with horses	19	81	5	95	0.005	2.955 (0.834–10.473)	0.093			
Contact with cattle	25	75	3	97	<0.001	8.607 (1.729–42.854)	0.009			
Contact with pigs	62	38	6	94	<0.001	20.455 (7.831–64.386)	<0.001			
Contact with sheep	2	98	2	98	0.689	NA	NA			

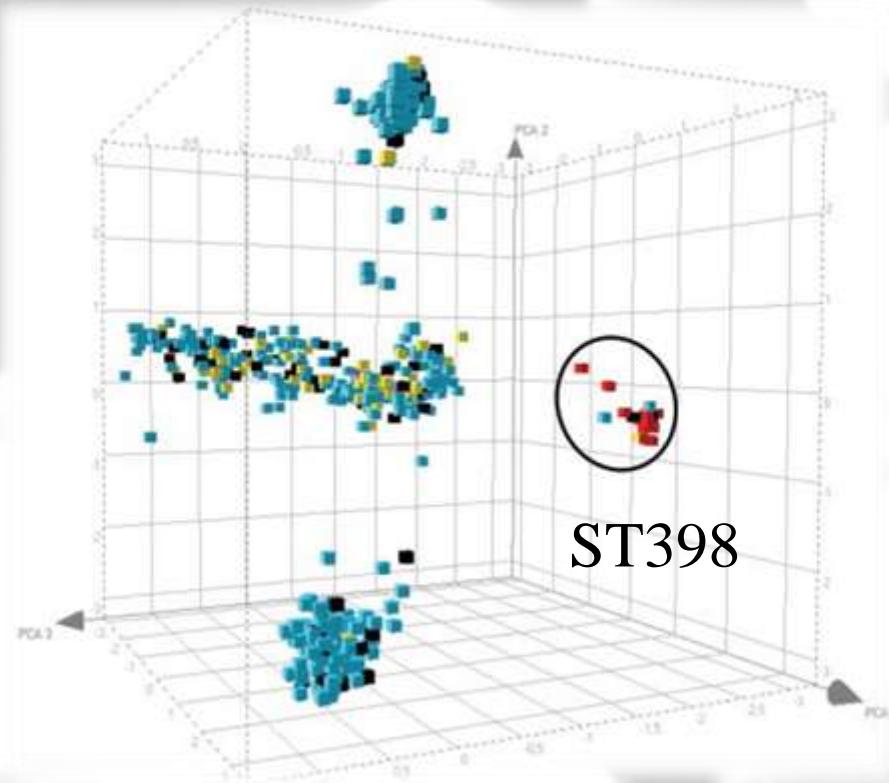
NA not available

^a *P* value resulting from univariate (Chi-square or Fisher Exact test) analysis; all factors with *P*<0.2 were included in the initial regression model

^b Odds ratio (95% confidence interval, CI) and *P* value of variables included in the final logistic regression model

^c During the previous 6 months

AFLP-analysis of clinical MSSA/MRSA-isolates



Blue, carriage isolates (n = 829);
black, bacteremia isolates (n = 146);
yellow, animal isolates (n = 77);
red, ST398 MRSA isolates (n = 46);
pink, reference strains (Mu50/N315).

Other reservoirs for MRSA?



- MRSA in horses, cattle
- MRSA in cats and dogs
- MRSA in rabbits
- MRSA in rats

Walther et al. Berl Munch Tierarztl Wochenschr 2006

Weese et al. J Am Vet Med Assoc 2007

Moodley et al. J Am Vet Med Assoc 2007

MRSA is not the same as MRSA

t011, t034, t108, t567, t571, t588, t753, t898, t899, t1184, t1254, t1255, t1451, t1456, t1457, t2123, t2330, t2383, t2582, t3013, t1197, t1344, t1793, t1939, t2329, t2922, t2346, t2370, t3479, t4659, t4838, t4854, t4872, t6291, t6292, t6293



PVL-negative (2 exceptions in SE, NL)
Livestock-associated (LA)-MRSA



t044, t019, t131, t376, t639, t237, t1199, t1201, t1206, t1200, t008, t437, t216, t005, t310, t1276, t355, t595, t311, t1277, t450, t019, t021, t318, t1273, t914, t202, t128, t125, t558, t175, t1274, t1272

PVL-positive
Community-acquired (CA)-MRSA

Community-MRSA

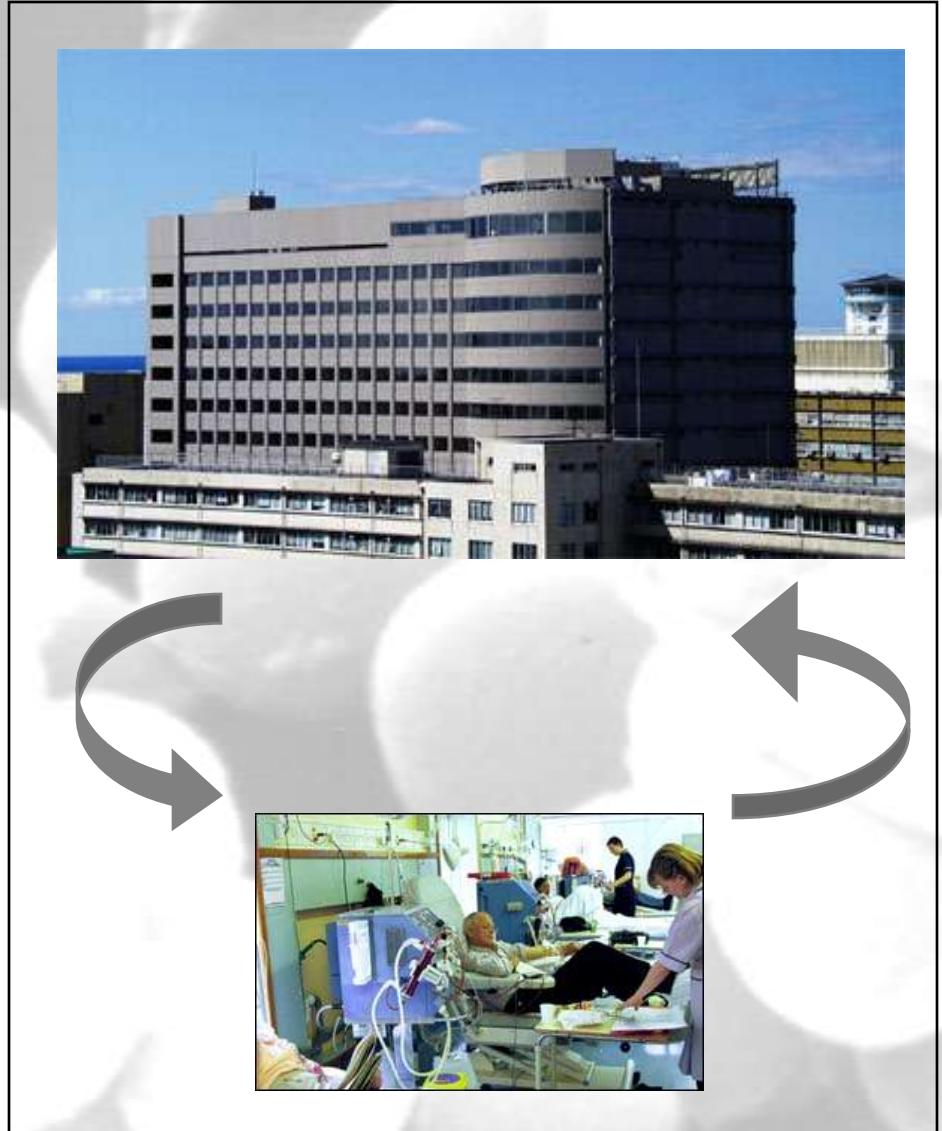


Hospital-acquired (HA)-MRSA

t003, t032, t008, t002, t037, t044, t001, t004, t015, t024, t084, t012, t045, t091, t127, t437, t067, t041, t019, t005, t030, t021, t038, t065, t026, t018, t022, t172

Hospital-MRSA

Extended MRSA-Epidemiology



Conclusions

1. MRSA spread **different** from MSSA
2. Resistances diffuse locally und regionally across the border
Transmission of **HA-MRSA** via health care networks
of **LA-MRSA** via production ways
of **CA-MRSA** via social networks
3. MRSA-prevention needs to be implemented at the critical points
4. **Preventive Microbiology**, infection control and antibiotic stewardship is the **future**
5. Molecular typing is of **epidemiological** and **clinical** interest
6. Collaboration between human medicine and veterinary medicine necessary (One-health)

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