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# Surveillance of ICU - Acquired Infections National Feedback Report Infection Indicators Year 2015

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Table 1: Participation

	#
unit quarters	30
hospital quarters	26
unit years	13
hospital years	11
level 1	10
level 2 basic	20
level 2a (risk factors / invasive devices)	20
level 2b (cvc utilization)	12
level 2c (antibiotic utilization)	14
Pneumonia	30
Bloodstream infections	30
Urinary tract infections	15
Catheter-related infections	15

<sup>a</sup> unit quarters = number of IC units that participated during a particular 3-month quarter; hospital quarters = number of hospitals that participated during a particular 3-month quarter; unit years = number of IC units that participated during at least one 3-month quarter ; hospital years = number of hospitals that participated during at least one 3-month quarter

<sup>b</sup> level = number of IC units that participated during a particular 3-month quarter; level 1 = unit-based surveillance; level 2 = patient-based surveillance

Table 2: General Indicators

	#					
	agg	p10	p25	p50	p75	p90
admissions, total	2968	46	76	146	327	526
patientdays, total	23490	360	593	1306	2464	4971
patientdays, mean	7.9	6.5	6.9	7.8	9.5	10.9
patientdays, median	5	4	5	5	6	8



Table 3: Pneumonia

	# <sup>a</sup>	rate <sup>a</sup>						
		agg.	agg.	p10	p25	p50	p75	p90
patients with P (all)	134	4.51	1.33	2.33	4.48	8.38	8.70	/100 admissions
patients with P $\geq$ D3 <sup>b</sup> (NP <sup>b</sup> )	125	4.21	1.33	2.33	4.48	7.94	8.70	/100 admissions
patients with NP, ID2 <sup>b</sup>	81	2.73	0.95	1.22	2.38	3.95	5.99	/100 admissions
P (all)	167	7.11	2.01	2.78	5.06	11.49	16.95	/1000 patientdays
P $\geq$ D3 (NP)	157	6.68	2.01	2.78	5.06	11.49	16.95	/1000 patientdays
NP, 1st <sup>b</sup>	124	5.28	1.62	2.78	4.72	11.49	13.56	/1000 patientdays
NP, ID2	87	8.47	1.02	3.72	10.46	13.33	14.68	/1000 IDdays <sup>b</sup>
NP, ID2, 1st	34	10.00	4.27	4.85	13.82	16.30	22.65	/1000 IDdays bef 1st NP

<sup>a</sup> #=total; rate=# divided by indicated denominator; agg.=aggregated over all participating units; p10-90=percentiles for distribution of participating units

<sup>b</sup> P=pneumonia, NP(s)=Nosocomial P,  $\geq$ D3=occurring after D2 (nosocomial), IDdays=Invasive Device (intubation for P) exposure days, ID2=Invasive Device (intubation for P) exposure in 2 days before onset of NP, 1st=only 1st infection per patient



Table 4: Bloodstream Infections

	# <sup>a</sup>	rate <sup>a</sup>					
	agg.	agg.	p10	p25	p50	p75	p90
patients with B (all)	54	1.82	0.79	1.32	1.71	2.65	2.99
patients with B $\geq$ D3 <sup>b</sup> (NB <sup>b</sup> )	47	1.58	0.00	0.00	1.32	1.83	2.12
patients with NB, ID2 <sup>b</sup>	35	1.18	0.00	0.00	1.14	1.80	1.83
B (all)	58	2.47	0.73	1.69	2.21	2.78	3.83
B $\geq$ D3 (NBs)	51	2.17	0.00	0.00	1.69	2.44	3.06
NB, 1st <sup>b</sup>	47	2.00	0.00	0.00	1.69	2.44	3.06
NB, ID2	36	2.10	0.00	0.92	1.79	2.44	6.59
NB, ID2, 1st	20	2.93	0.00	0.99	2.61	3.45	11.90
NB, ori=cat/unk <sup>b,c</sup>	31	1.32	0.00	0.00	1.22	1.69	1.82
NB, (ori=cat/unk, ID2)/(ori=cat) <sup>d</sup>	24	1.40	0.00	0.00	1.45	1.80	2.57
NB, ori=cat <sup>e</sup>	7	0.41	0.00	0.00	0.00	0.80	1.20
NB, ori=unk	24	1.02	0.00	0.00	0.77	1.42	1.69
NB, ori=cat or unk, 1st	29	1.23	0.00	0.00	1.22	1.69	1.81
NB, (ori=cat or unk, ID2)/(ori=cat), 1st	14	2.05	0.00	0.00	2.33	2.64	3.97
NB, ori=cat, 1st	3	0.44	0.00	0.00	0.00	1.15	1.98
NB, ori=unk, 1st	22	0.94	0.00	0.00	0.77	1.42	1.61
NBs, ID2, $\geq$ 1HC path or $\geq$ 2HCS with sc <sup>b</sup>	21	3.07	0.00	0.99	2.61	3.45	13.89
NBs, ID2, $\geq$ 1HC path or $\geq$ 2HCS with sc, ori=cat	3	0.44	0.00	0.00	0.00	1.15	1.98
NBs, ID2, $\geq$ 1HC path or $\geq$ 2HCS with sc, ori=cat, 1 st	3	0.44	0.00	0.00	0.00	1.15	1.98

<sup>a</sup> #=total; rate=# divided by indicated denominator; agg.=aggregated over all participating units; p10-90=percentiles for distribution of participating units

<sup>b</sup> B=Bloodstream Infection episodes; NB(s)=Nosocomial B;  $\geq$ D3=occurring after D2 (nosocomial); IDdays=Invasive Device (central vascular catheter for B) days; ID2=Invasive Device (central vascular catheter for B) in 2 days before onset of B; 1st=only 1st infection per patient;

ori=cat/unk=NB origin is either catheter or unknown; ori=cat=NB origin is catheter;  $\geq$ 1HC path=at least one hemoculture from a pathogen;  $\geq$ 2HCS with sc=at least 2 hemocultures from a skincontaminant

<sup>c</sup> Primary BSI  
<sup>d</sup> Catheter Associated Primary BSI  
<sup>e</sup> Definite Catheter Associated Primary BSI



Table 5: Urinary Tract Infections

	# <sup>a</sup>					rate <sup>a</sup>
	agg.	agg.	p10	p25	p50	
patients with U (all)	59	6.40	0.71	1.80	2.33	4.35
patients with U $\geq$ D3 (NU) <sup>b</sup>	53	5.75	0.71	1.80	2.33	4.35
patients with NU, ID2 <sup>b</sup>	41	4.45	0.00	0.71	1.80	4.35
U (all)	79	8.93	0.91	1.89	2.78	6.78
U $\geq$ D3 (NU)	73	8.26	0.91	1.89	2.78	6.78
NU, 1st <sup>b</sup>	53	5.99	0.91	1.42	2.78	6.78
NU, ID2	52	7.23	1.32	1.32	2.00	10.61
NU, ID2, 1st	39	5.25	1.83	1.83	2.06	10.35

<sup>a</sup> #=total; rate=# divided by indicated denominator; agg.=aggregated over all participating units; p10-90=percentiles for distribution of participating units

<sup>b</sup> U=Urinary Tract Infection; NU=Nosocomial U;  $\geq$ D3=occurring after D2 (nosocomial); IDdays=Invasive Device (urinary catheter for U) days; ID2=Invasive Device (urinary catheter for U) in 2 days before onset of U; 1st=only 1st infection per patient



Table 6: Catheter Related Infections

	# <sup>a</sup>	rate <sup>a</sup>					
	agg.	agg.	p10	p25	p50	p75	p90
patients with CRI (all)							
patients with CRI $\geq$ D3 (NCRI) <sup>b</sup>	4	0.39	0.00	0.00	0.31	0.57	0.57
CRI (all)	4	0.39	0.00	0.00	0.31	0.57	0.57
CRI $\geq$ D3 (NCRI)	4	0.42	0.00	0.00	0.41	0.60	0.60
NCRI, 1st <sup>b</sup>	4	0.42	0.00	0.00	0.41	0.60	0.60

<sup>a</sup> #=total; rate=# divided by indicated denominator; agg.=aggregated over all participating units; p10-90=percentiles for distribution of participating units

<sup>b</sup> U=Catheter Related Infection; NCRI=Nosocomial CRI;  $\geq$ D3=occurring after D2 (nosocomial); 1st=only 1st infection per patient



Table 7: Case Definitions: Nosocomial Pneumonia

	# <sup>a</sup>	% <sup>a</sup>					
	agg.	agg.	p10	p25	p50	p75	p90
PN1	25	15.9	0.0	0.0	0.0	0.0	19.6
PN2	10	6.4	0.0	0.0	0.0	20.0	100.0
PN3	1	0.6	0.0	0.0	0.0	0.0	0.0
PN4	111	70.7	0.0	0.0	80.0	100.0	100.0
PN5	10	6.4	0.0	0.0	0.0	0.0	40.0

<sup>a</sup> #=crude number; %=percentage; agg.=aggregated, treating national sample as one unit; p10-90=percentiles for national distribution

<sup>b</sup> PN=clinical definition for Nosocomial Pneumonia; PN1=PN and positive quantitative culture from minimally contaminated lower respiratory tract specimen; PN2=PN and positive quantitative culture from possibly contaminated lower respiratory tract specimen; PN3=PN and alternative microbiology methods; PN4=PN and positive sputum culture or non-quantitative LRT specimen culture; PN5=PN with no positive microbiology

Table 8: Case Definitions: Nosocomial Bloodstream Infections

	# <sup>a</sup>	% <sup>a</sup>					
	agg.	agg.	p10	p25	p50	p75	p90
BSI-A	50	98.0	83.3	100.0	100.0	100.0	100.0
BSI-B	1	2.0	0.0	0.0	0.0	0.0	16.7

<sup>a</sup> #=crude number; %=percentage; agg.=aggregated, treating national sample as one unit; p10-90=percentiles for national distribution

<sup>b</sup> BSI-A=positive blood culture for a recognised pathogen or clinical signs with 2 positive blood cultures for a common skin contaminant; BSI-B=clinical signs and positive blood culture with a skin contaminant or positive blood antigen test

Table 9: Case Definitions: Nosocomial Urinary Tract Infections

	# <sup>a</sup>	% <sup>a</sup>					
	aggregated	aggregated	p10	p25	p50	p75	p90
UTI-A	71	97.3	0.0	100.0	100.0	100.0	100.0
UTI-B	0	0.0	0.0	0.0	0.0	0.0	0.0
UTI-C	2	2.7	0.0	0.0	0.0	0.0	100.0

<sup>a</sup> #=crude number; %=percentage; agg.=aggregated, treating national sample as one unit; p10-90=percentiles for national distribution

<sup>b</sup> UTI-A=microbiologically confirmed symptomatic UTI; UTI-B=not microbiologically confirmed symptomatic UTI; UTI-C=asymptomatic bacteriuria

Table 10: Case Definitions: Nosocomial Catheter Related Infections

	# <sup>a</sup>	% <sup>a</sup>					
	aggregated	aggregated	p10	p25	p50	p75	p90
CRI1	4	100.0	100.0	100.0	100.0	100.0	100.0
CRI2	0	0.0	0.0	0.0	0.0	0.0	0.0
CRI3	0	0.0	0.0	0.0	0.0	0.0	0.0

<sup>a</sup> #=crude number; %=percentage; agg.=aggregated, treating national sample as one unit; p10-90=percentiles for national distribution

<sup>b</sup> CRI1=local CVC-related infection; CRI2=general CVC-related infection; CRI3=CVC-related BSI



Table 11: All reported micro-organisms (nosocomial infections) (a)

microorganism - family	code	NP <sup>a</sup>		NB <sup>a</sup>		NU <sup>a</sup>		tot <sup>a</sup>	
		# <sup>b</sup>	% <sup>b</sup>	#	%	#	%	#	%
<b>Gram-positive cocci</b>									
<i>Enterococcus faecalis</i>	ENCFAE	0	0.0	5	9.3	5	6.9	11	3.5
<i>Enterococcus faecium</i>	ENCFAI	0	0.0	6	11.1	5	6.9	11	3.5
<i>Enterococcus sp., not specified</i>	ENCNSP	1	0.6	0	0.0	0	0.0	1	0.3
<i>Enterococcus sp., other</i>	ENCOTH	0	0.0	0	0.0	1	1.4	1	0.3
<i>Other coagulase-negative staphylococci (cns)</i>	STAOTH	0	0.0	1	1.9	0	0.0	1	0.3
<i>Staphylococcus aureus</i>	STAAUR	24	13.3	6	11.1	0	0.0	31	9.9
<i>Staphylococcus haemolyticus</i>	STAHAE	1	0.6	0	0.0	0	0.0	1	0.3
<i>Staphylococcus sp., not specified</i>	STANSP	0	0.0	1	1.9	0	0.0	1	0.3
<i>Streptococcus pneumoniae</i>	STRPNE	5	2.8	1	1.9	0	0.0	6	1.9
<i>Streptococcus sp., other</i>	STROTH	0	0.0	0	0.0	0	0.0	0	0.0
<b>Gram negative cocci</b>									
<i>Moraxella catharralis</i>	MORCAT	3	1.7	0	0.0	0	0.0	3	1.0
<b>Gram positive bacilli</b>									
<i>Bacillus species</i>	BACSP	0	0.0	0	0.0	0	0.0	1	0.3
<b>Gram-negative bacilli, enterobacteriaceae</b>									
<i>Citrobacter freundii</i>	CITFRE	1	0.6	0	0.0	2	2.8	3	1.0
<i>Citrobacter koseri (ex. diversus)</i>	CITDIV	1	0.6	0	0.0	2	2.8	3	1.0
<i>Citrobacter sp., not specified</i>	CITNSP	1	0.6	0	0.0	0	0.0	1	0.3
<i>Citrobacter sp., other</i>	CITOOTH	1	0.6	0	0.0	0	0.0	1	0.3
<i>Enterobacter aerogenes</i>	ENBAER	3	1.7	0	0.0	1	1.4	4	1.3
<i>Enterobacter cloacae</i>	ENBCLO	14	7.8	3	5.6	1	1.4	18	5.8
<i>Enterobacter sp., other</i>	ENBOTH	1	0.6	0	0.0	0	0.0	1	0.3
<i>Enterobacteriaceae, not specified</i>	ETBNSP	2	1.1	0	0.0	0	0.0	2	0.6
<i>Escherichia coli</i>	ESCCOL	26	14.4	6	11.1	13	18.1	46	14.7
<i>Hafnia species</i>	HAFSP	4	2.2	0	0.0	0	0.0	4	1.3
<i>Klebsiella oxytoca</i>	KLEOXY	6	3.3	3	5.6	0	0.0	9	2.9
<i>Klebsiella pneumoniae</i>	KLEPNE	13	7.2	5	9.3	2	2.8	20	6.4
<i>Klebsiella sp., not specified</i>	KLENSP	0	0.0	1	1.9	0	0.0	1	0.3
<i>Klebsiella sp., other</i>	KLEOTH	2	1.1	0	0.0	0	0.0	2	0.6
<i>Morganella species</i>	MOGSPP	3	1.7	0	0.0	3	4.2	6	1.9
<i>Proteus mirabilis</i>	PRTMIR	4	2.2	1	1.9	3	4.2	8	2.6
<i>Proteus vulgaris</i>	PRTVUL	0	0.0	0	0.0	1	1.4	1	0.3
<i>Serratia liquefaciens</i>	SERLIQ	0	0.0	1	1.9	0	0.0	1	0.3
<i>Serratia marcescens</i>	SERMAR	7	3.9	1	1.9	1	1.4	9	2.9

<sup>a</sup> NP=Nosocomial Pneumonia, NB=Nosocomial Bloodstream Infection, NU=Nosocomial Urinary Tract Infection,  
tot=all Nosocomial Infections

<sup>b</sup> #=number of micro-organisms, % =percentage of total micro-organisms



Table 12: All reported micro-organisms (nosocomial infections) (b)

microorganism - family	code	NP <sup>a</sup>		NB <sup>a</sup>		NU <sup>a</sup>		tot <sup>a</sup>	
		# <sup>b</sup>	% <sup>b</sup>	#	%	#	%	#	%
<b>Gram-negative bacilli, other</b>									
<i>Haemophilus influenzae</i>	HAEINF	39	21.7	6	11.1	11	15.3	56	17.9
<i>Haemophilus sp., not specified</i>	HAENSP	8	4.4	0	0.0	0	0.0	8	2.6
<i>Pseudomonas aeruginosa</i>	PSEAER	1	0.6	0	0.0	0	0.0	1	0.3
<i>Stenotrophomonas maltophilia</i>	STEMAL	25	13.9	6	11.1	11	15.3	42	13.4
		5	2.8	0	0.0	0	0.0	5	1.6
<b>Anaerobes</b>									
<i>Bacteroides fragilis</i>	BATFRA	0	0.0	1	1.9	0	0.0	2	0.6
<i>Clostridium difficile</i>	CLODIF	0	0.0	0	0.0	0	0.0	1	0.3
<b>Other bacteria</b>									
<i>Other bacteria, not specified</i>	BCTNSP	2	1.1	0	0.0	0	0.0	2	0.6
		2	1.1	0	0.0	0	0.0	2	0.6
<b>Parasites</b>									
<i>Aspergillus fumigatus</i>	ASPFUM	4	2.2	5	9.3	21	29.2	32	10.2
<i>Candida albicans</i>	CANALB	1	0.6	0	0.0	0	0.0	1	0.3
<i>Candida glabrata</i>	CANGLA	2	1.1	3	5.6	14	19.4	20	6.4
<i>Candida sp., not specified</i>	CANNSP	1	0.6	2	3.7	3	4.2	7	2.2
<i>Other yeasts</i>	YEAOOTH	0	0.0	0	0.0	2	2.8	2	0.6
<b>Virus</b>									
<i>Herpes simplex virus</i>	VIRHSV	1	0.6	0	0.0	0	0.0	1	0.3
		1	0.6	0	0.0	0	0.0	1	0.3

<sup>a</sup> NP=Nosocomial Pneumonia, NB=Nosocomial Bloodstream Infection, NU=Nosocomial Urinary Tract Infection, tot=all Nosocomial Infections

<sup>b</sup> #=number of micro-organisms, % =percentage of total micro-organisms



Table 13: Anti Microbial Resistance Indicators I: Gram Positive Cocci

	SIR <sup>a</sup>						IR <sup>a</sup>					
	# <sup>a</sup>	% <sup>a</sup>	p10	p25	p50	p75	# <sup>a</sup>	% <sub>o</sub> <sup>a</sup>	p10	p25	p50	p75
<b><i>Staphylococcus aureus</i></b>												
cultures	31											
Fluoroquinolones: Ciprofloxacin/ofloxacin	7	22.6	0.0	0.0	0.0	28.6	100.0	4	57.1	0.0	0.0	0.0
Macrolides/sim.: Clindamycin (lincosamides)	2	6.5	0.0	0.0	0.0	0.0	100.0	1	50.0	0.0	50.0	100.0
Macrolides/sim.: Erythromycin (macrolides)	2	6.5	0.0	0.0	0.0	0.0	100.0	1	50.0	0.0	50.0	100.0
Other antibiotics: Fosfomycin	0	0.0	0.0	0.0	0.0	0.0	0.0					
Other antibiotics: Fusidic acid	0	0.0	0.0	0.0	0.0	0.0	0.0					
Aminoglycosides: Gentamycin	14	45.2	0.0	0.0	0.0	0.0	100.0	0	0.0	0.0	0.0	0.0
Glycopeptides: Vancomycin/teicoplanin (Glycopeptides)	14	45.2	0.0	0.0	28.6	100.0	1	7.1	0.0	0.0	0.0	0.0
Penicillins: Methicillin/Oxacillin (beta-lact.res.penic.)	29	93.5	66.7	100.0	100.0	100.0	100.0	9	31.0	0.0	0.0	50.0
Penicillins: Penicillin	2	6.5	0.0	0.0	0.0	0.0	100.0	2	100.0	100.0	100.0	100.0
<b><i>Streptococcus pneumoniae</i></b>												
cultures	6											
Penicillins: Ampicillin	0	0.0	0.0	0.0	0.0	0.0	0.0					
Cephalosporins: Cefotaxime-ceftiraxone (3rd gen cephalosp.)	3	50.0	0.0	0.0	50.0	100.0	0	0.0	0.0	0.0	0.0	0.0
Macrolides/sim.: Clindamycin (lincosamides)	0	0.0	0.0	0.0	0.0	0.0	0.0					
Macrolides/sim.: Erythromycin (macrolides)	2	33.3	0.0	0.0	0.0	50.0	100.0	2	100.0	100.0	100.0	100.0
Penicillins: Methicillin/Oxacillin (beta-lact.res.penic.)	2	33.3	0.0	0.0	0.0	50.0	100.0	2	100.0	100.0	100.0	100.0
Penicillins: Penicillin	0	0.0	0.0	0.0	0.0	0.0	0.0					
Tetracyclines: Tetra-doxo/minocycline (tetracyclines)	0	0.0	0.0	0.0	0.0	0.0	0.0					
<b><i>Enterococcus faecalis</i></b>												
cultures	11											
Penicillins: Ampicillin	8	72.7	0.0	41.7	91.7	100.0	100.0	0	0.0	0.0	0.0	0.0
Penicillins: Amoxicillin/clavulanate	6	54.5	0.0	16.7	66.7	100.0	100.0	1	16.7	0.0	0.0	50.0
Aminoglycosides: Gentamycin	8	72.7	50.0	50.0	66.7	91.7	100.0	1	12.5	0.0	0.0	10.0
Glycopeptides: Vancomycin/teicoplanin (Glycopeptides)	10	90.9	0.0	50.0	100.0	100.0	100.0	0	0.0	0.0	0.0	0.0

<sup>a</sup> SIR-all Sensitive/Intermediate/Resistant antibiogram results, IR-all Intermediate/Resistant antibiogram results, #=aggregated total, %#=#/100 cultures (SIR) or #/#100 SIR results (IR),



Table 14: Anti Microbial Resistance Indicators II: Enterobacteriaceae (a)

	SIR <sup>a</sup>						IR <sup>a</sup>						
	# <sup>a</sup>	% <sup>a</sup>	p10	p25	p50	p75	p90	# <sup>a</sup>	% <sup>a</sup>	p10	p25	p50	p75
<b><i>Escherichia coli</i></b>													
cultures	46	73.9	0.0	59.1	100.0	100.0	100.0	17	50.0	0.0	33.3	50.0	61.5
Penicillins: Amoxicillin/clavulanate	34	47.8	0.0	12.2	100.0	100.0	100.0	7	31.8	0.0	8.3	50.0	100.0
Cephalosporins: Cefotaxime/ceftriaxone (3rd gen cephalosp.)	22	52.2	0.0	19.9	100.0	100.0	100.0	0	0.0	0.0	0.0	0.0	0.0
Carbapenems: Meropenem/imipenem (carbapenems)	24	13.0	0.0	0.0	100.0	100.0	100.0	1	16.7	0.0	0.0	33.3	33.3
ESBL: Extended Spectrum Beta Lactamase	6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<b><i>Enterobacter aerogenes</i></b>													
cultures	4	100.0	100.0	100.0	100.0	100.0	100.0	4	100.0	100.0	100.0	100.0	100.0
Penicillins: Amoxicillin/clavulanate	1	25.0	0.0	0.0	0.0	100.0	100.0	0	0.0	0.0	0.0	0.0	0.0
Cephalosporins: Cefotaxime/ceftriaxone (3rd gen cephalosp.)	2	50.0	0.0	0.0	100.0	100.0	100.0	0	0.0	0.0	0.0	0.0	0.0
Carbapenems: Meropenem/imipenem (carbapenems)	0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
ESBL: Extended Spectrum Beta Lactamase	0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
<b><i>Enterobacter cloacae</i></b>													
cultures	18	83.3	0.0	33.3	100.0	100.0	100.0	15	100.0	100.0	100.0	100.0	100.0
Penicillins: Amoxicillin/clavulanate	15	61.1	0.0	33.3	50.0	100.0	100.0	8	72.7	0.0	30.0	80.0	100.0
Cephalosporins: Cefotaxime/ceftriaxone (3rd gen cephalosp.)	11	61.1	0.0	33.3	50.0	100.0	100.0	0	0.0	0.0	0.0	0.0	0.0
Carbapenems: Meropenem/imipenem (carbapenems)	11	5.6	0.0	0.0	0.0	100.0	100.0	0	0.0	0.0	0.0	0.0	0.0
ESBL: Extended Spectrum Beta Lactamase	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0

<sup>a</sup> SIR=all Sensitive/Intermediate/Resistant antibiogram results, IR=all Intermediate/Resistant antibiogram results, #=aggregated total, %=#/100 cultures (SIR) or #/#100 SIR results (IR),



Table 15: Anti Microbial Resistance Indicators III: Enterobacteriaceae (b)

	SIR <sup>a</sup>						IR <sup>a</sup>						
	# <sup>a</sup>	% <sup>a</sup>	p10	p25	p50	p75	p90	# <sup>a</sup>	% <sup>a</sup>	p10	p25	p50	p75
<b><i>Klebsiella pneumoniae</i></b>													
cultures	20	85.0	0.0	66.7	100.0	100.0	100.0	6	35.3	0.0	33.3	41.7	50.0
Penicillins: Amoxicillin/clavulanate	17	60.0	0.0	0.0	100.0	100.0	100.0	4	33.3	0.0	0.0	33.3	100.0
Cephalosporins: Cefotaxime/ceftriaxone (3rd gen cephalosp.)	12	60.0	0.0	0.0	100.0	100.0	100.0	0	0.0	0.0	0.0	0.0	100.0
Carbapenems: Meropenem/imipenem (carbapenems)	12	60.0	0.0	0.0	100.0	100.0	100.0	3	37.5	0.0	16.7	66.7	100.0
ESBL: Extended Spectrum Beta Lactamase	8	40.0	0.0	0.0	100.0	100.0	100.0	3	37.5	0.0	16.7	66.7	100.0
<b><i>Klebsiella oxytoca</i></b>													
cultures	9	100.0	100.0	100.0	100.0	100.0	100.0	6	66.7	25.0	25.0	100.0	100.0
Penicillins: Amoxicillin/clavulanate	9	66.7	25.0	25.0	100.0	100.0	100.0	5	83.3	75.0	75.0	100.0	100.0
Cephalosporins: Cefotaxime/ceftriaxone (3rd gen cephalosp.)	6	55.6	0.0	0.0	100.0	100.0	100.0	0	0.0	0.0	0.0	0.0	100.0
Carbapenems: Meropenem/imipenem (carbapenems)	5	11.1	0.0	0.0	0.0	100.0	100.0	1	100.0	100.0	100.0	100.0	100.0
ESBL: Extended Spectrum Beta Lactamase	1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0

<sup>a</sup> SIR=all Sensitive/Intermediate/Resistant antibiogram results, IR=all Intermediate/Resistant antibiogram results, #=aggregated total, %#=#/100 cultures (SIR) or #/#/100 SIR results (IR),



Table 16: Anti Microbial Resistance Indicators IV: Nonfermentative Bacilli

	SIR <sup>a</sup>						IR <sup>a</sup>						
	# <sup>a</sup>	% <sup>a</sup>	p10	p25	p50	p75	p90	# <sup>a</sup>	% <sup>a</sup>	p10	p25	p50	p75
<b>Pseudomonas aeruginosa</b>													
Cultures	42												
Carbapenems: Meropenem/imipenem (carbapenems)	27	64.3	0.0	37.5	100.0	100.0	11	40.7	0.0	7.1	53.3	100.0	
Carbapenems: Ceftazidim (anti-pseudom 3G ceph)	35	83.3	0.0	50.0	100.0	100.0	10	28.6	0.0	18.8	29.2	80.0	
Cephalosporins: Ceftazidim (anti-pseudom 3G ceph)	9	21.4	0.0	0.0	0.0	75.0	100.0	1	11.1	0.0	0.0	20.0	
Polymyx.: Colistin (polymixins)	9	21.4	0.0	0.0	0.0	83.3	100.0	4	44.4	0.0	0.0	30.0	
Penicillins: Piperacillin/ticarcillin (anti-pseudom. penic.)	27	64.3	0.0	0.0	25.0	83.3	100.0	5	18.5	0.0	0.0	12.5	
Penicillins: Piperacillin/ticarcillin + enzyme inhibitor													

**Acinetobacter baumanii**

no data available

<sup>a</sup> SIR=all Sensitive/Intermediate/Resistant antibiogram results, IR=all Intermediate/Resistant antibiogram results, #=aggregated total, %=#/100 cultures (SIR) or #/100 SIR results (IR),



Table 17: Missing Infection variables

variable	L1 <sup>a</sup>	L2 <sup>a</sup>	Ls <sup>a</sup>	Req <sup>a</sup>	# <sup>a</sup>	% <sup>a</sup>	p10	p25	p50	p75	p90
invasive device in 48 hours preceding infection	V	V	0	R	38	13.1	0.0	0.0	0.0	26.3	26.9
origin of bloodstream infection	V	V	0	O	21	36.2	0.0	0.0	15.6	90.9	100.0
antimicrobial treatment	V	V	0	O	119	41.0	0.0	0.0	6.7	100.0	101.4
Microorganism indicator	V	V	0	M	3	1.0	0.0	0.0	0.0	0.0	0.0
total infection variables	V	V	0	O	181	14.4	2.3	3.3	7.5	25.0	25.3

<sup>a</sup> l1= level 1 variable (denominator-based surveillance) ; l2= level 2 variable (patient-based surveillance) ; L2s= level 2 suboption (0=basic level 2 variables a=extra patient admission and day by day variables, b=central vascular catheter follow-up, c=antimicrobial treatment follow-up); Req: M=mandatory, R=Required, O=Optional #=number of missing variables; %=# / 100 eligible records



Hospital (all) :  
2968 patients

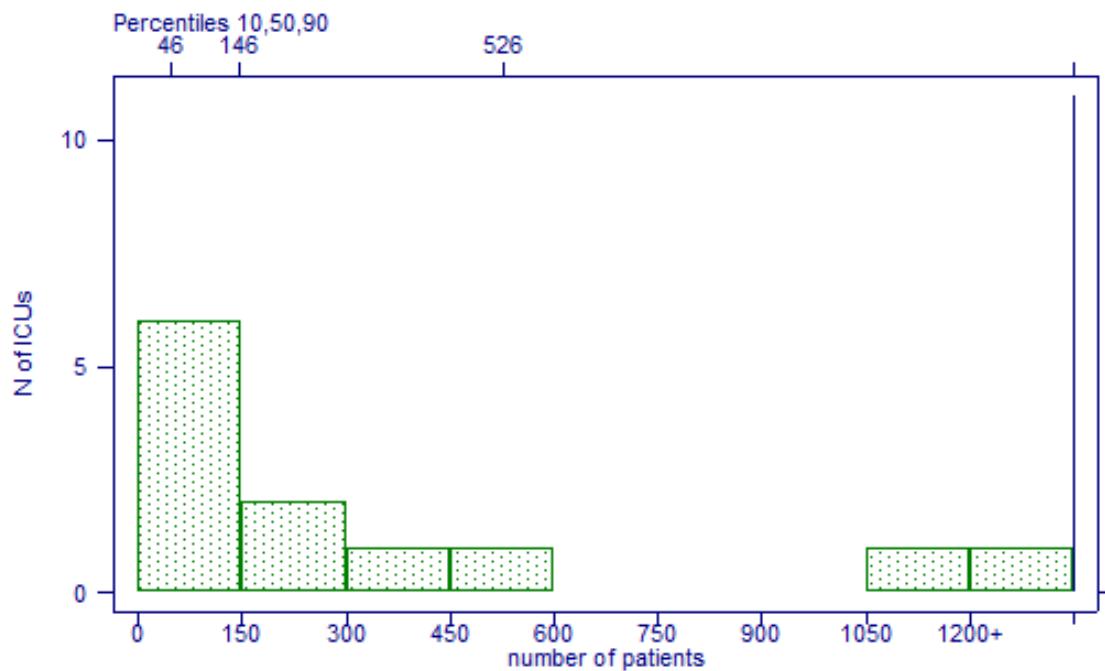


Figure 1: Distribution: number of patients

Hospital (all) :  
7.9 days

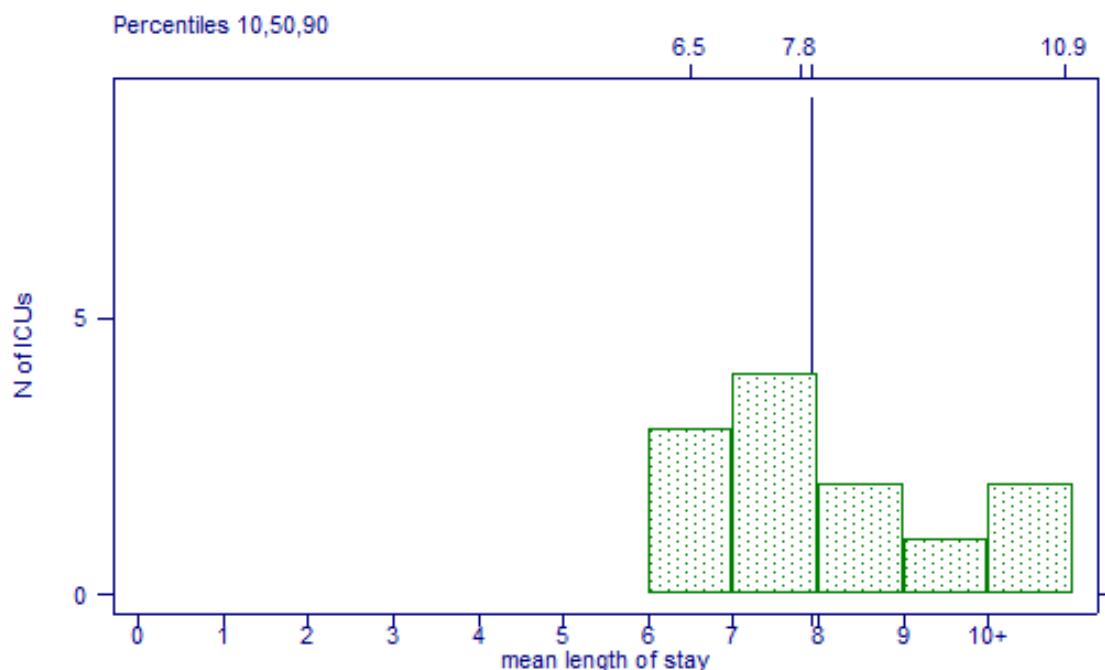


Figure 2: Distribution: mean length of stay



Hospital (all) :  
6.7 NP/1000 patientdays (95% CI 5.7 - 7.8)

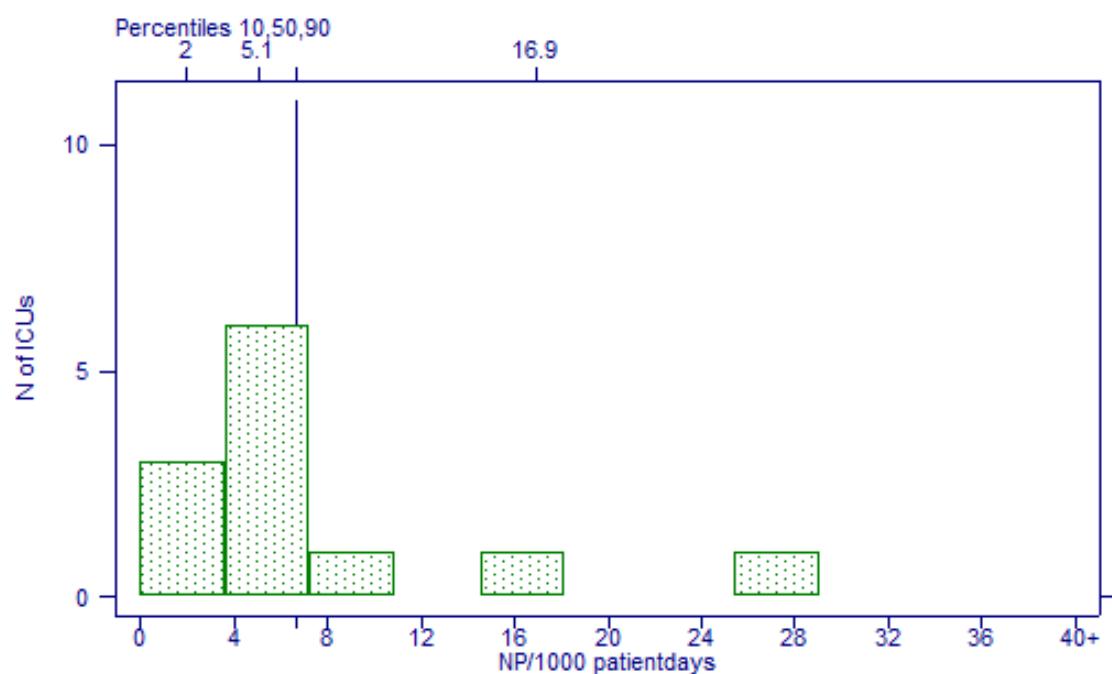


Figure 3: Distribution:  $P \geq D3$  (NP) / 1000 patientdays

Hospital (all) :  
8.5 NPs, ID2, all / 1000 IDdays (95% CI 6.8 - 10.5)

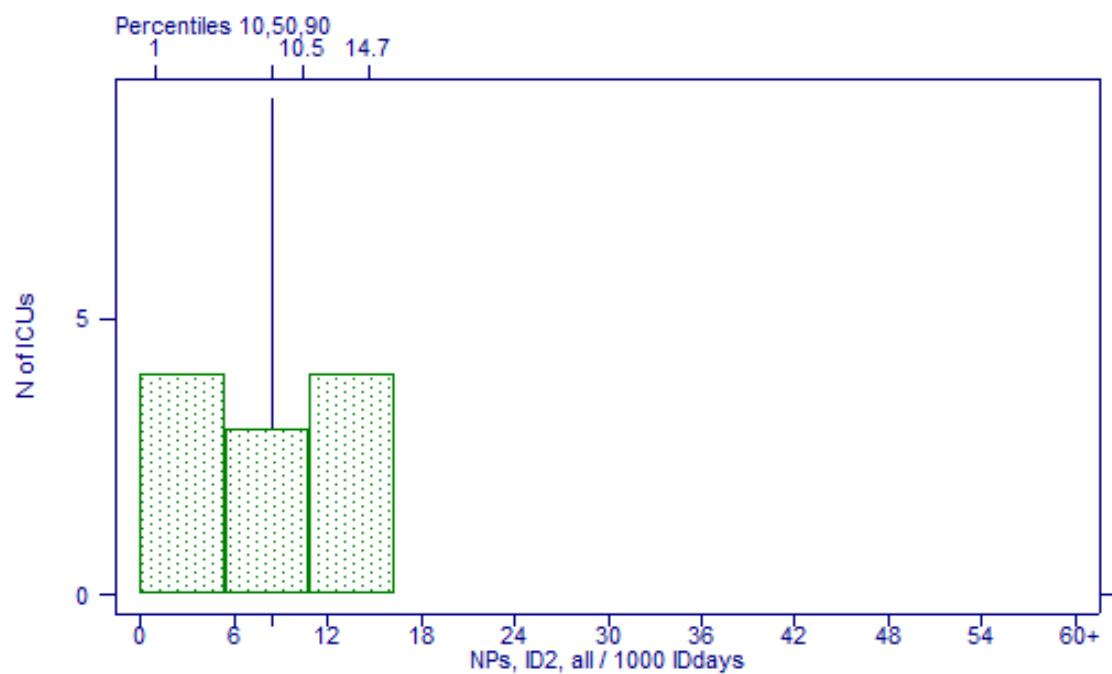


Figure 4: Distribution: NPs, ID2, all / 1000 IDdays ; NPs, with intubation in 2 days before onset of NP, all episodes / 1000 intubation days

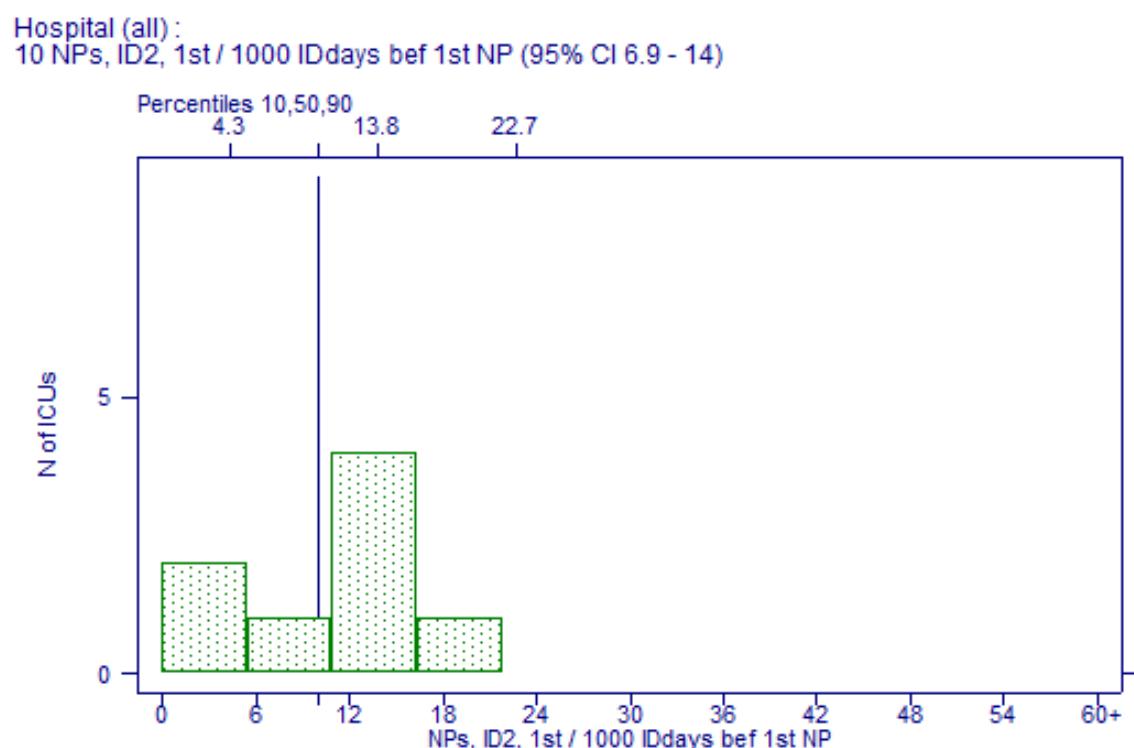


Figure 5: Distribution: NPs, ID2, 1st / 1000 IDdays bef 1st NP ; NPs, 1st episode only, with intubation in 2 days before onset of NP / 1000 intubation days



Hospital (all) :  
2.2 Bs $\geq$ D3 (NBs) / 1000 patientdays (95% CI 1.6 - 2.9)

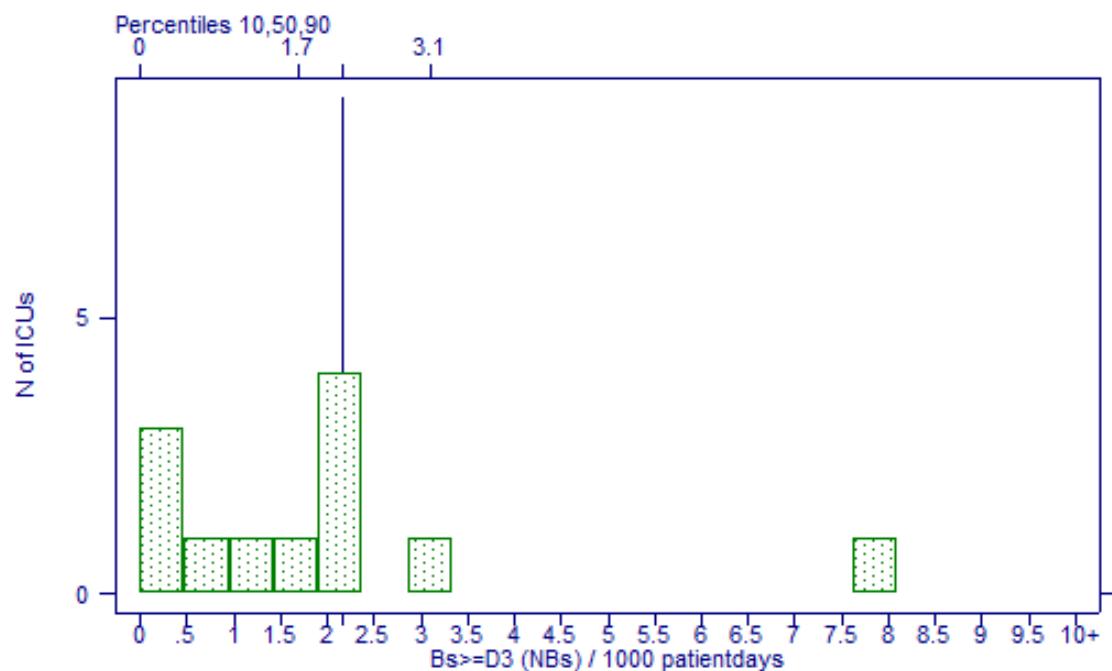


Figure 6: Distribution: Bs $\geq$ D3 (NBs) / 1000 patientdays

Hospital (all) :  
2.1 NBs, ID2, all / 1000 IDdays (95% CI 1.5 - 2.9)

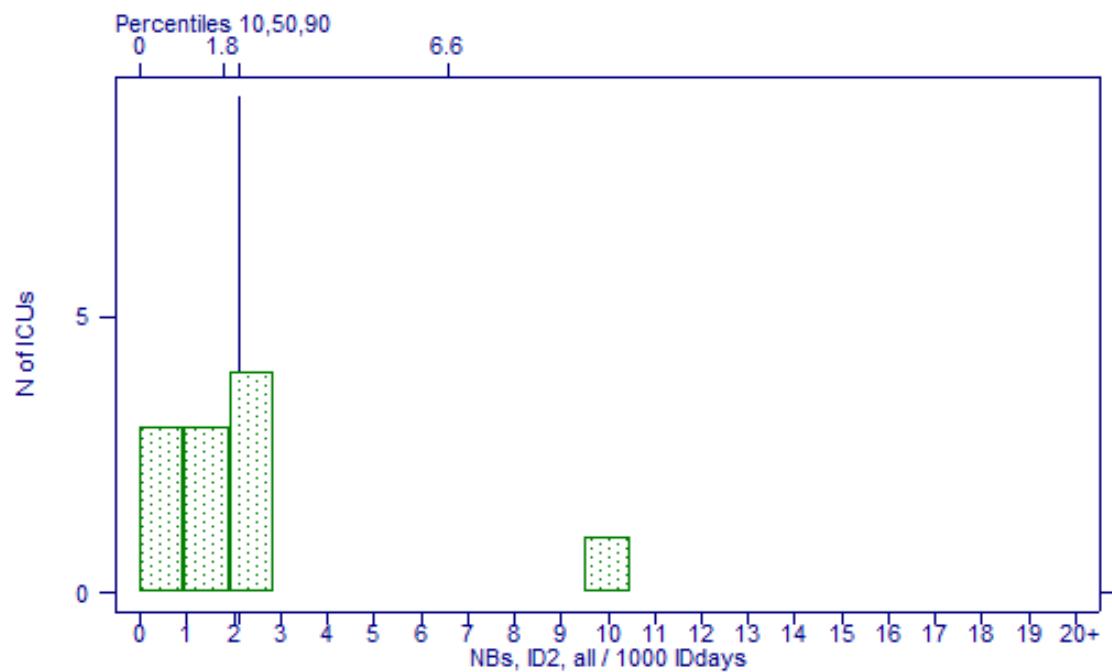


Figure 7: Distribution: NBs, with catheter use in 2days preceding onset of infection, all / 1000 IDdays ; NBs, with catheter use in 2days preceding onset of infection, all episodes / 1000 catheterdays



Hospital (all) :  
1.3 NBs, ori=(cat/un) / 1000 patientdays (95% CI .9 - 1.9)

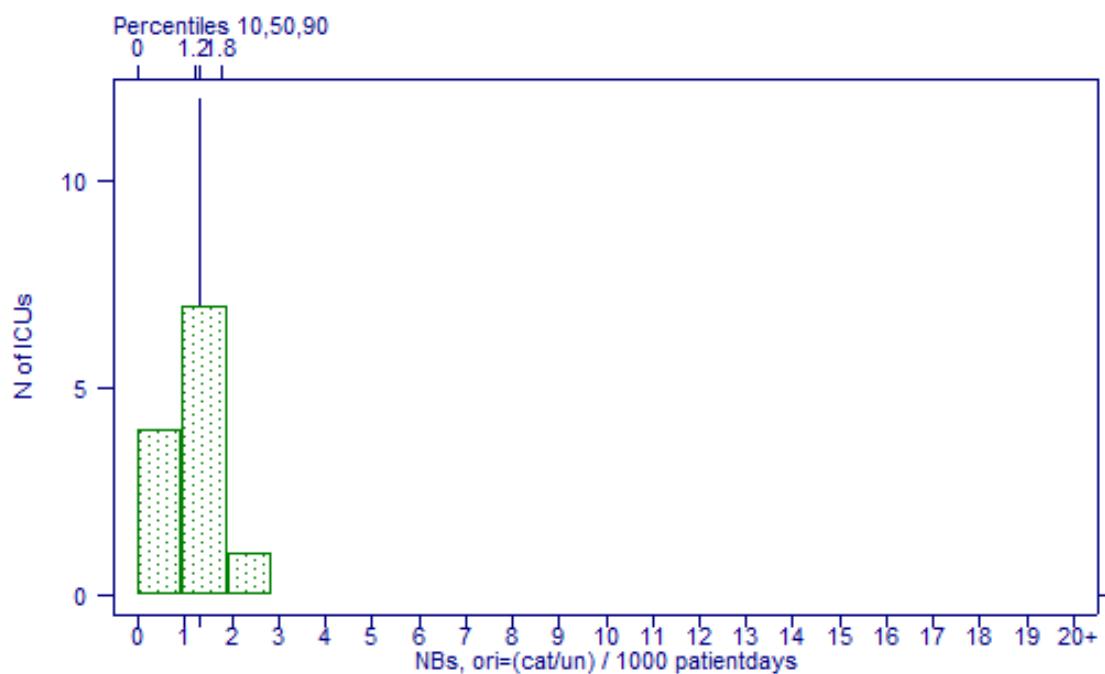


Figure 8: Distribution: NBs, ori=cat/unk / 1000 patientdays ; primary NBs / 1000 patientdays

Hospital (all) :  
1.4 NBs, (ori=cat/un, ID2) / 1000 IDdays (95% CI .9 - 2.1)

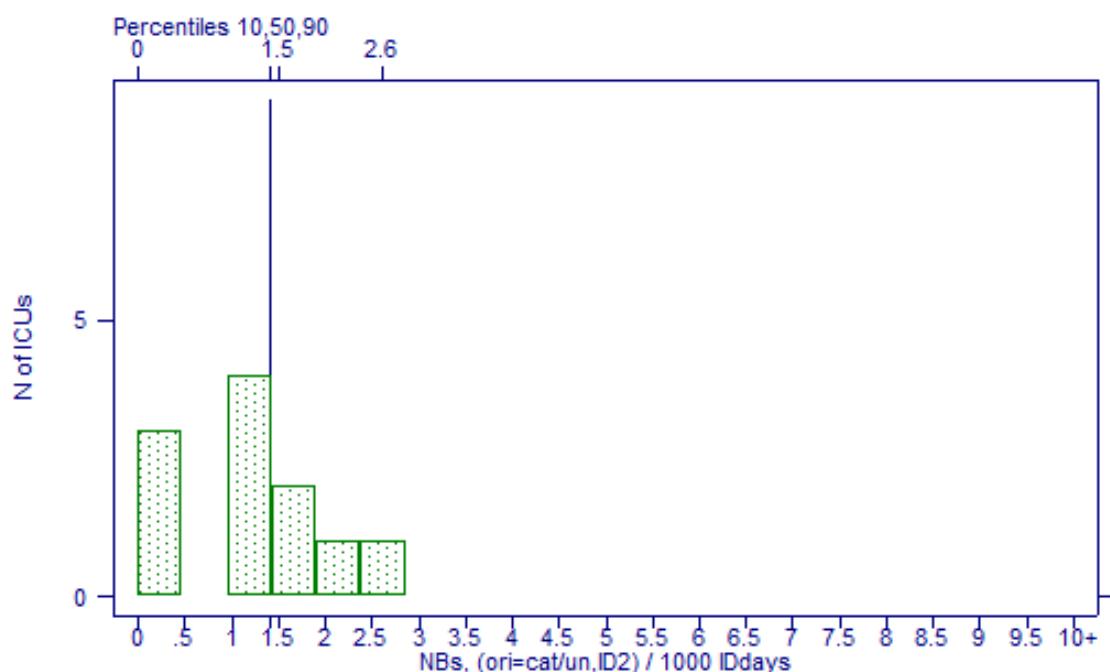


Figure 9: Distribution: NBs, (ori=cat/un, ID2) OR (ori=cat) / 1000 IDdays ; catheter associated NBs / 1000 catheterdays (CDC)



Hospital (all) :  
.4 NBs, ori=cat, 1st / 1000 IDdays bef 1st NB (95% CI .1 - 1.3)

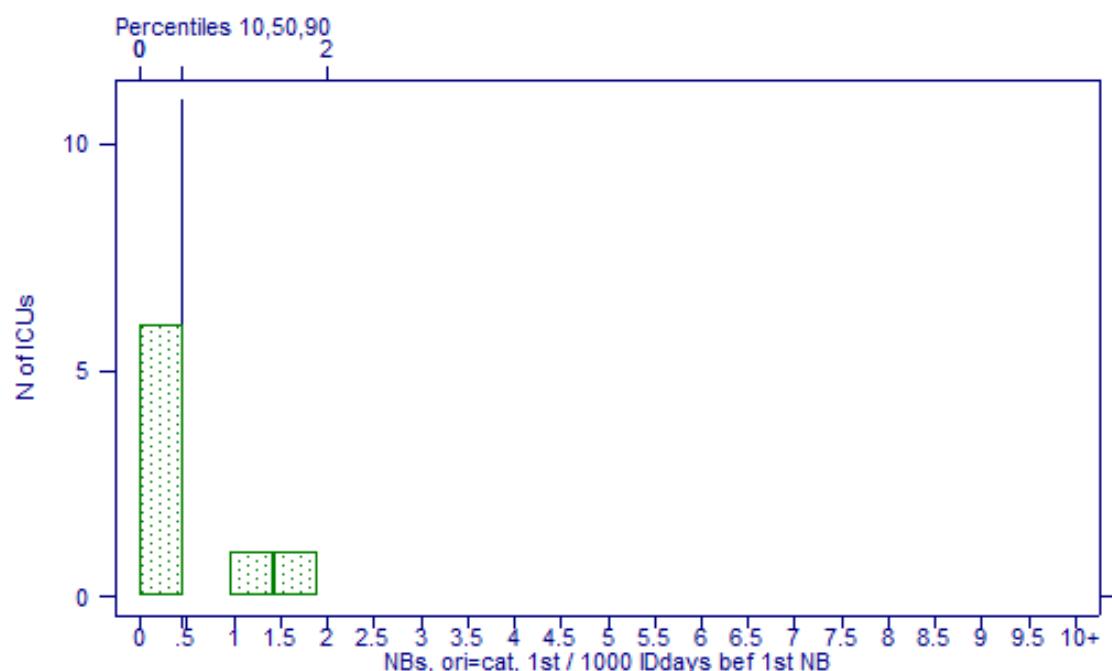


Figure 10: Distribution: NBs, ori=cat, 1st / 1000 IDdays bef 1st NB ; definite catheter associated NBs, 1st episodes only / 1000 catheterdays



Hospital (all) :  
3.1 NUs / 1000 patientdays (95% CI 2.4 - 3.9)

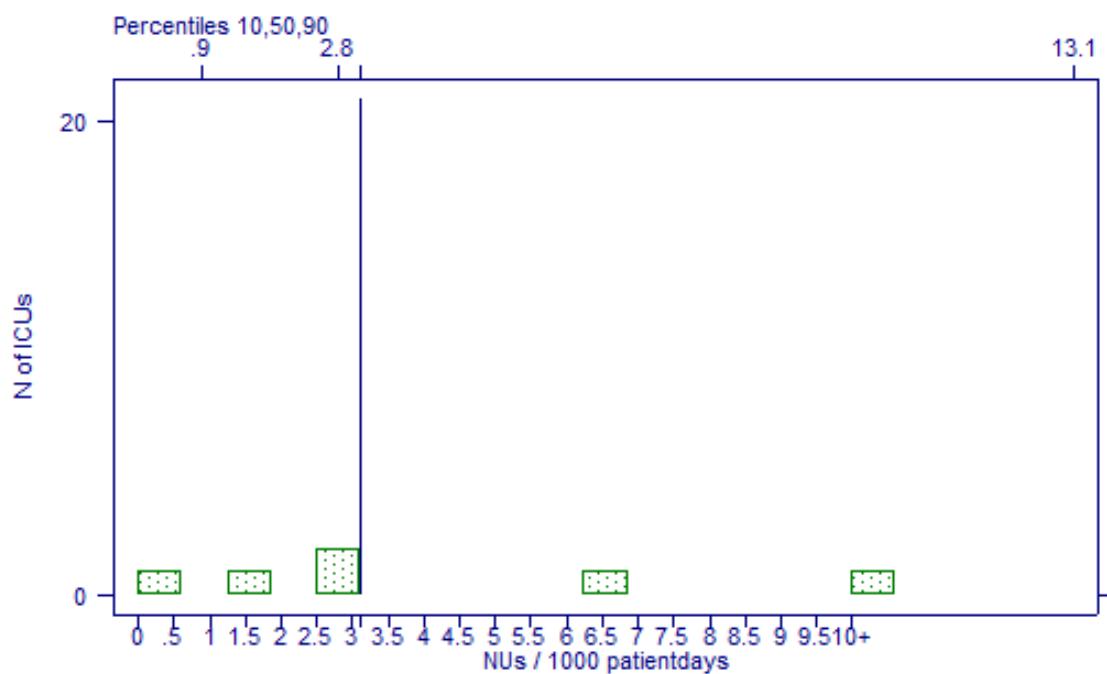


Figure 11: Distribution: Us $\geq$ D3 (NU) / 1000 patientdays ; NUs / 1000 patientdays

Hospital (all) :  
5.3 NUs, ID2 / 1000 ID days (95% CI 3.9 - 6.9)

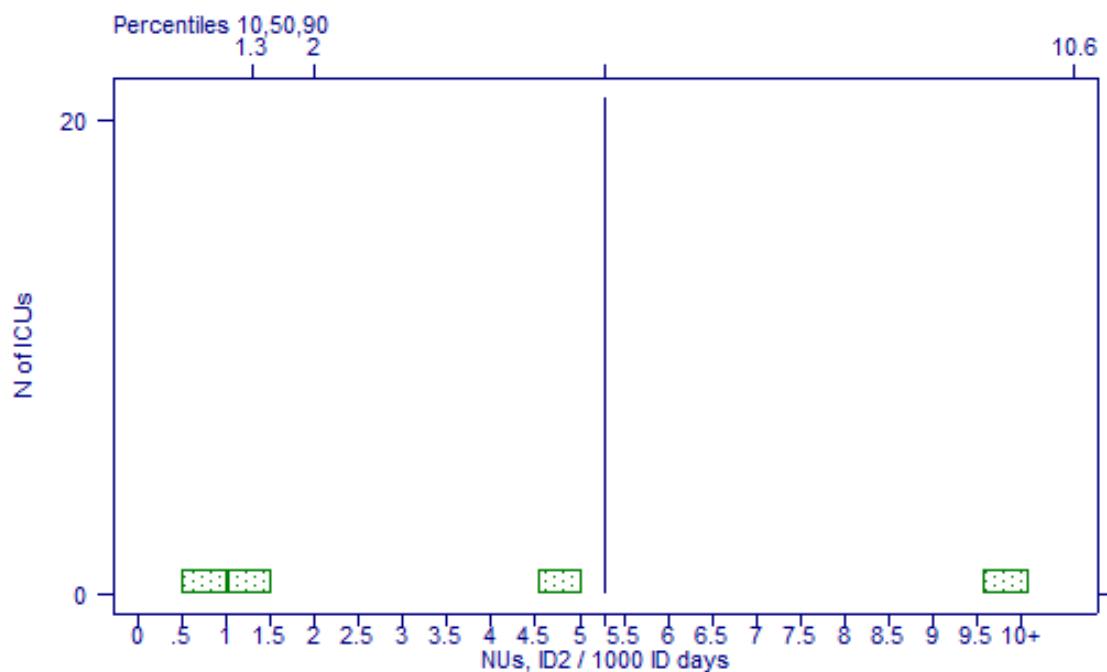


Figure 12: Distribution: NUs, ID2 / 1000 ID days ; catheter associated NUs / 1000 urinary catheterdays

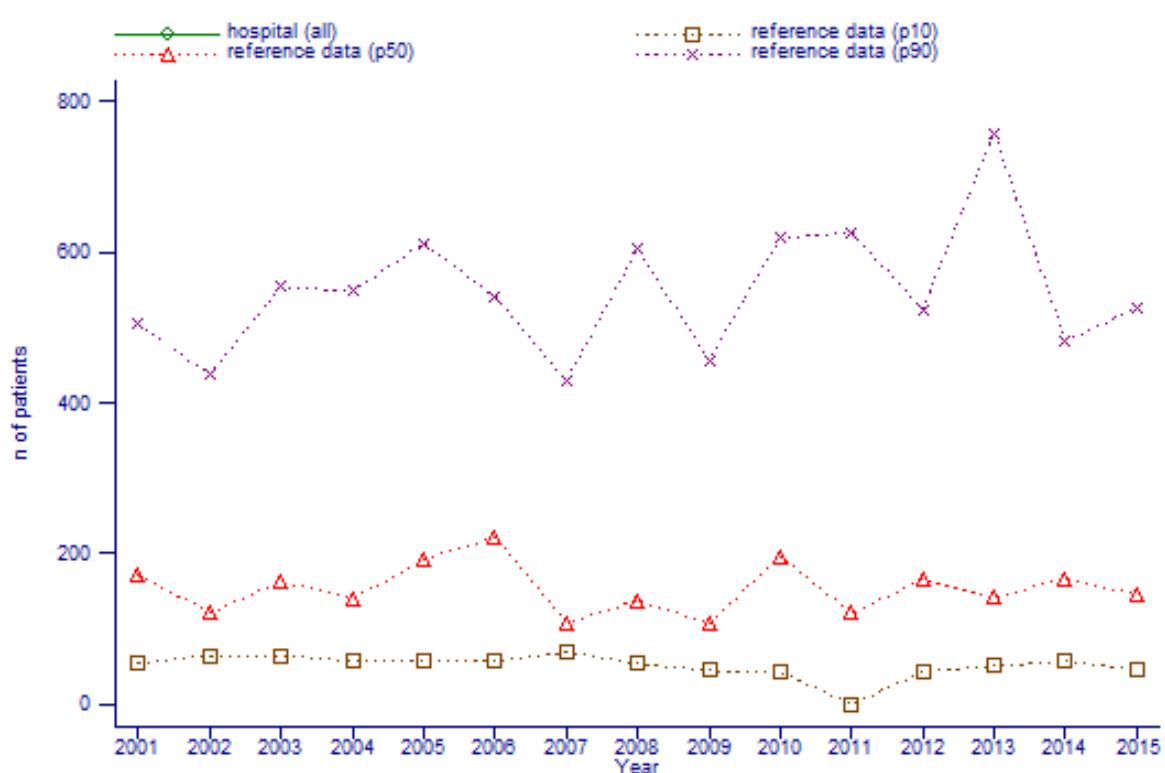


Figure 13: Evolution: Number of patients

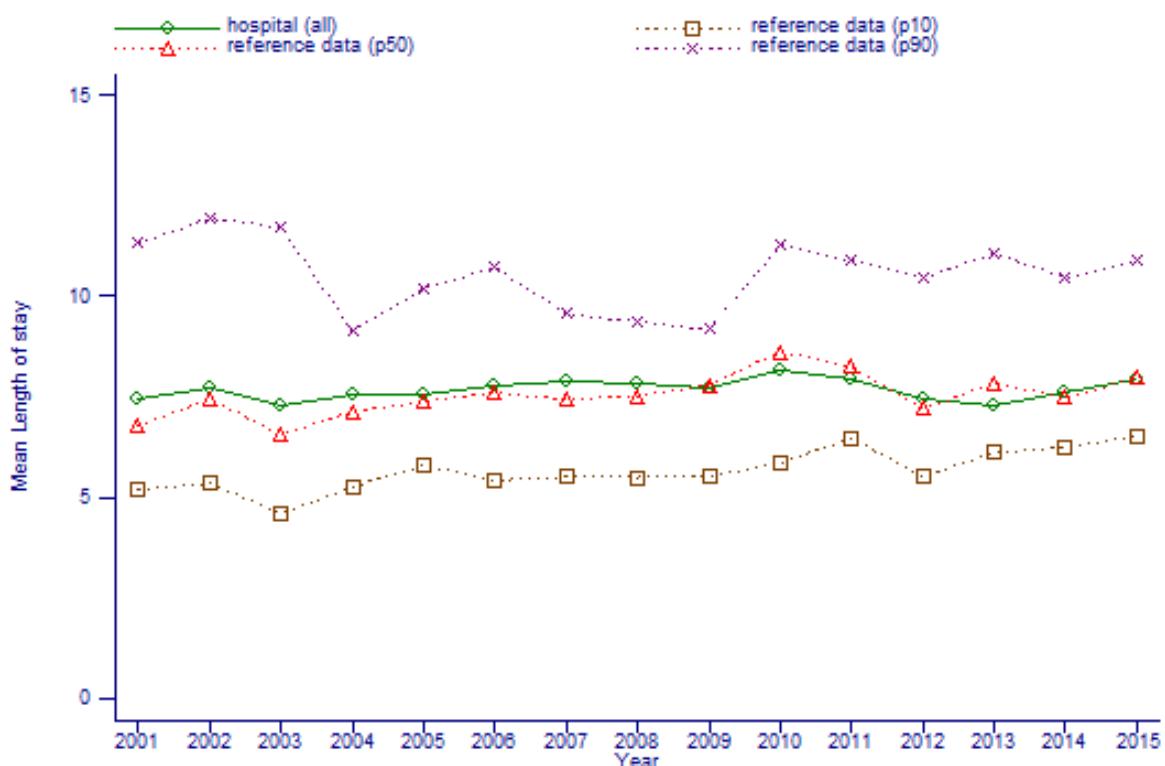


Figure 14: Evolution: Mean Length of stay

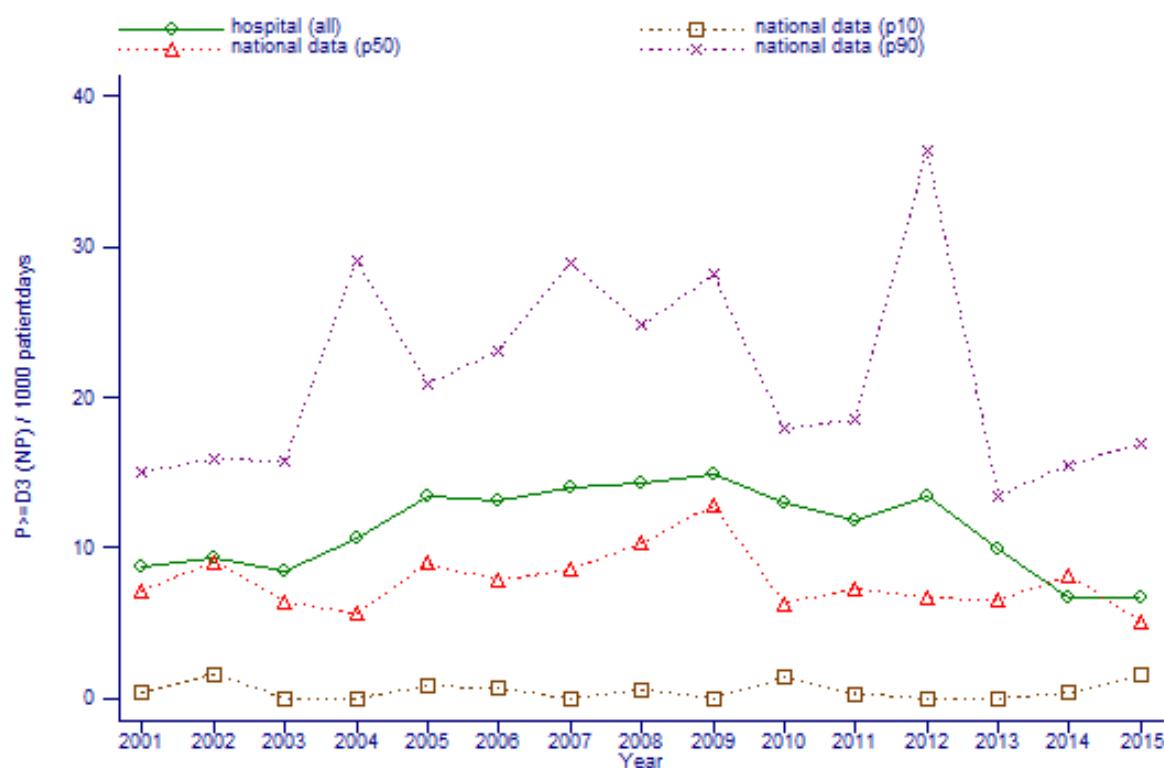


Figure 15: Evolution:  $P \geq D3 (NP) / 1000$  patientdays

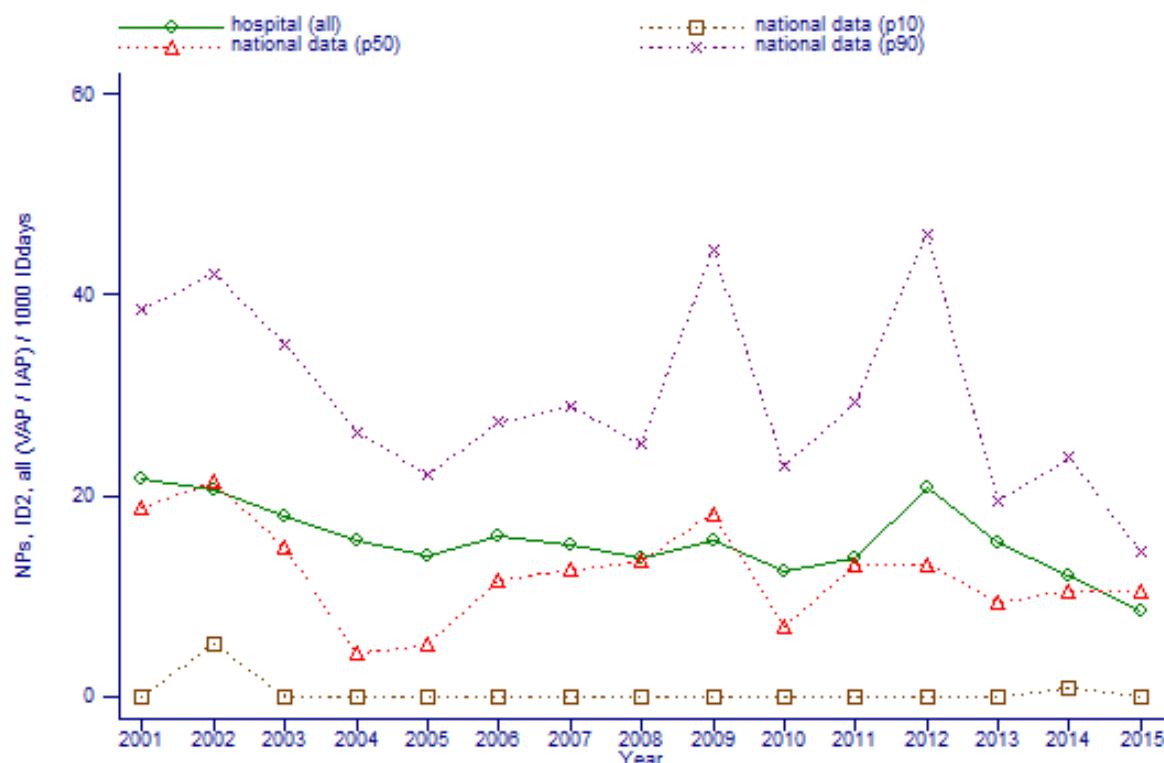


Figure 16: Evolution: NPs, ID2, all / 1000 IDdays ; NPs, with intubation in 2 days before onset of NP, all episodes / 1000 intubation days

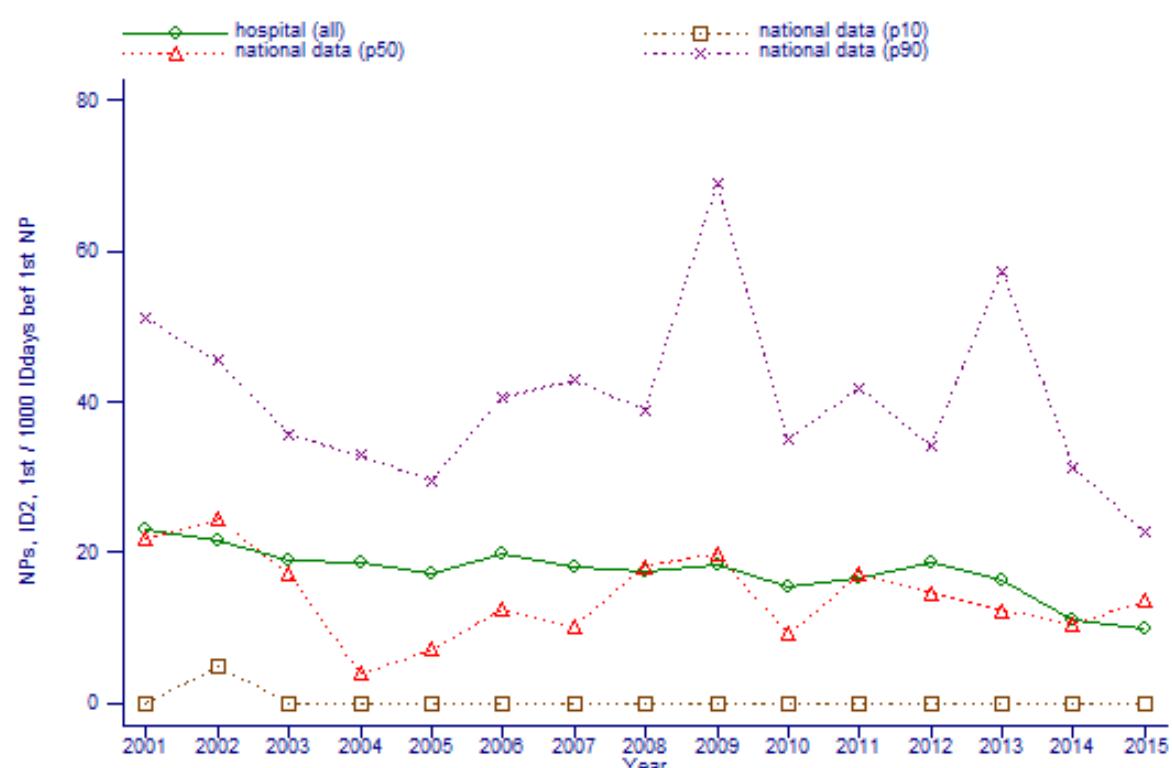


Figure 17: Evolution: NPs, ID2, 1st / 1000 IDdays bef 1st NP ; NPs, with intubation in 2 days before onset of NP, 1st episode only / 1000 intubation days

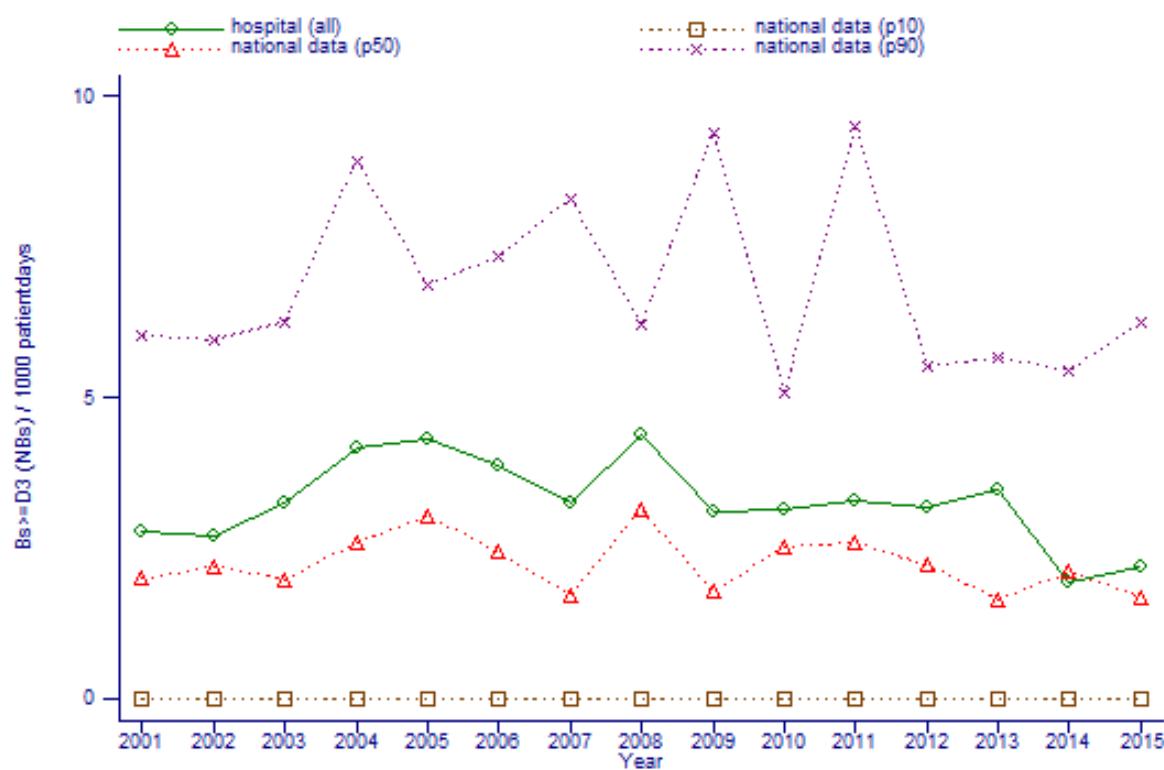


Figure 18: Evolution:  $BS \geq D3 (NBs) / 1000 \text{ patientdays}$

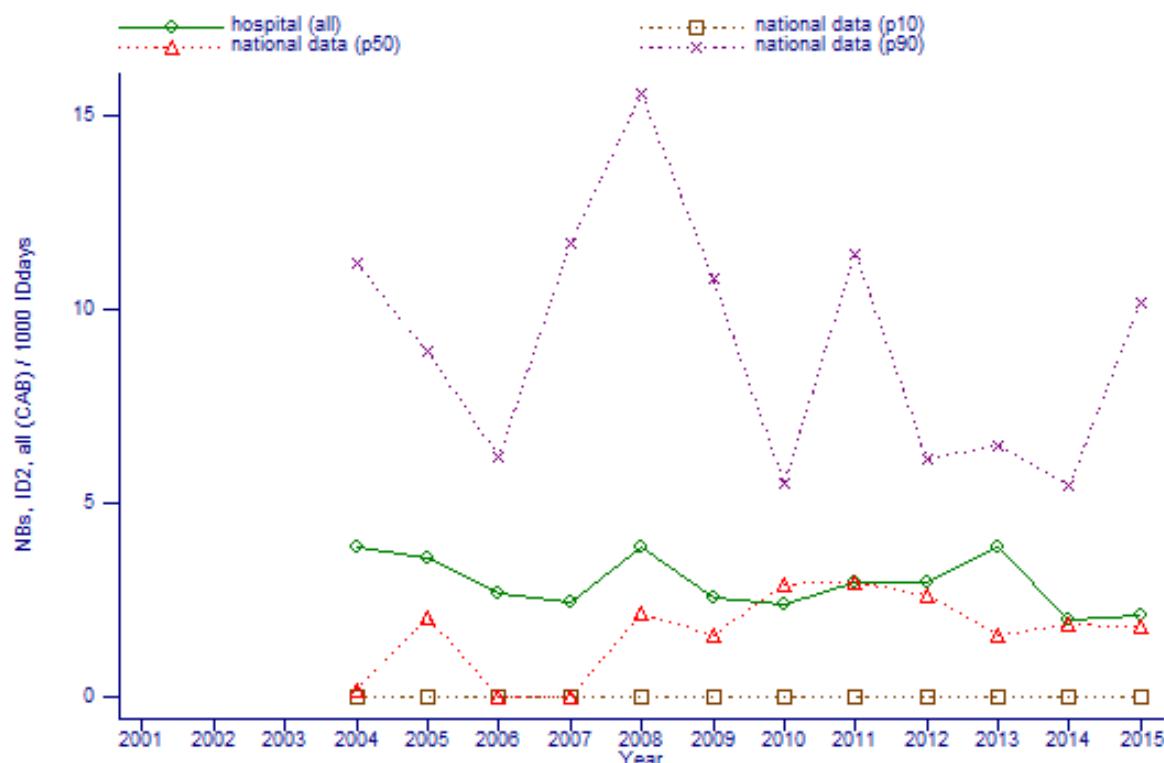


Figure 19: Evolution:  $NBs, ID2, \text{all} / 1000 \text{ IDdays}$ ;  $NBs, \text{with catheter use in 2 days preceding onset of infection, all episodes} / 1000 \text{ catheterdays}$

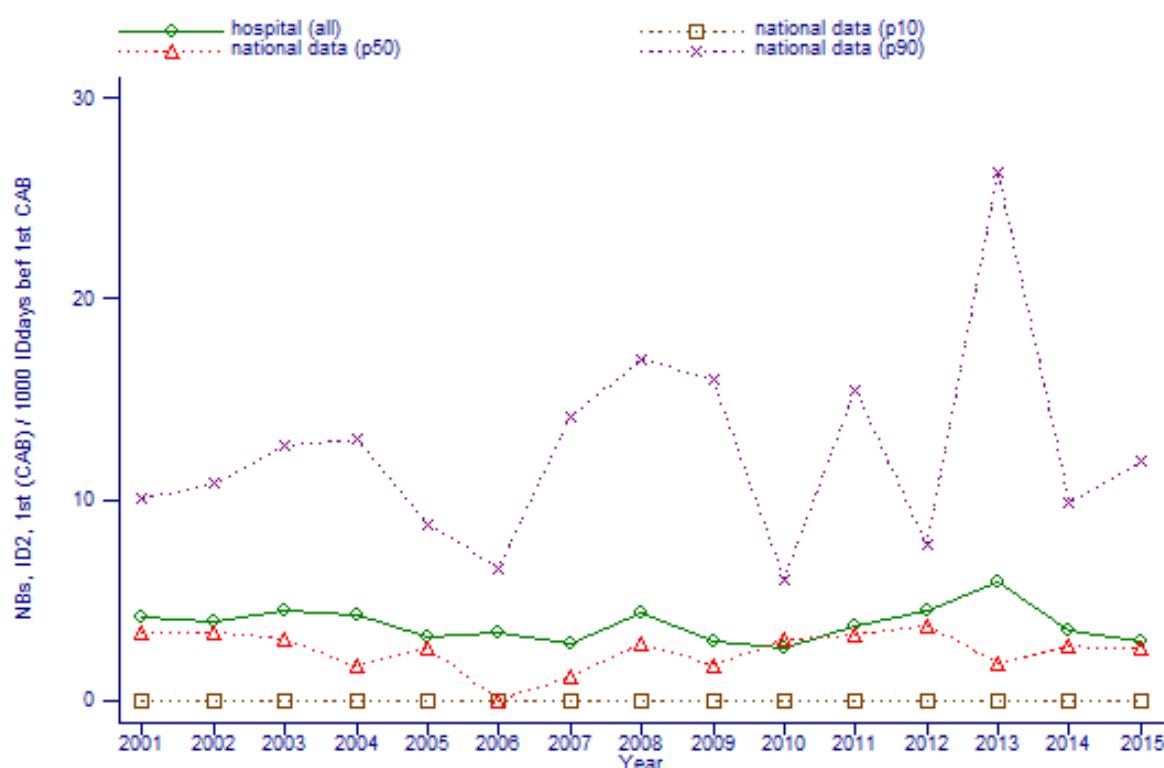


Figure 20: Evolution: NBs, ID2, 1st / 1000 IDdays bef 1st CAB; NBs, with catheter use in 2days preceding onset of infection, 1st episodes only / 1000 catheterdays

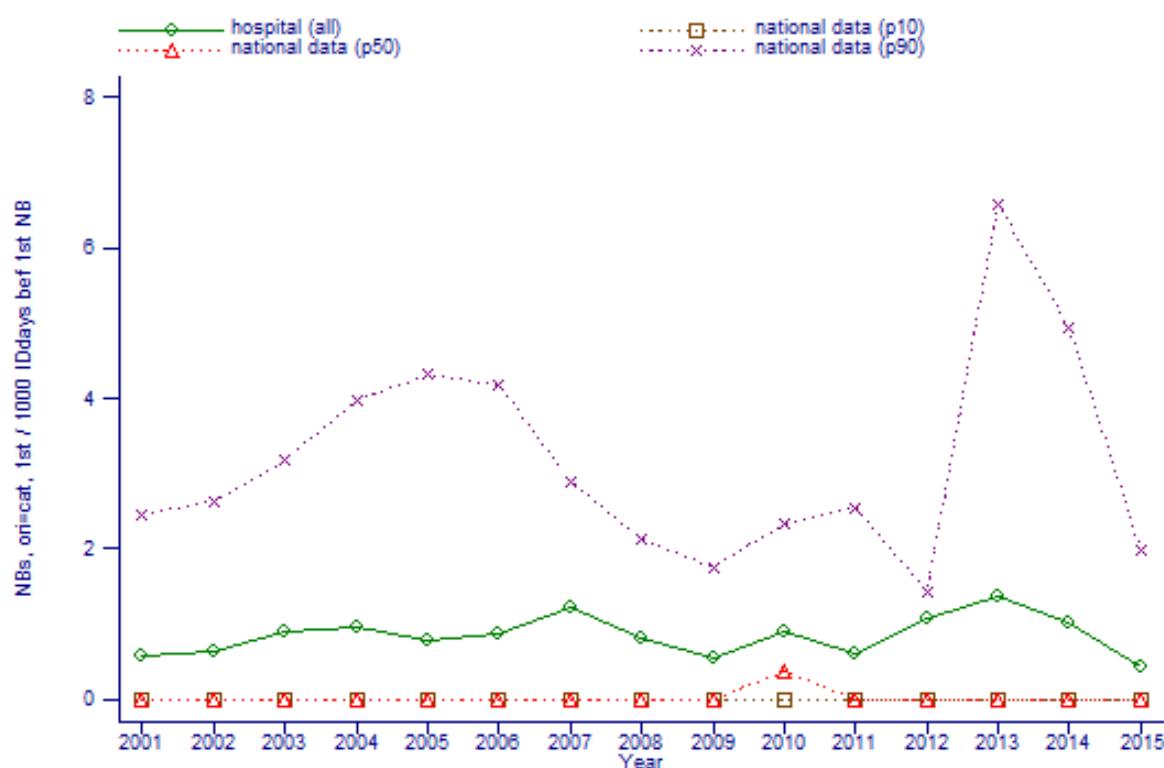


Figure 21: Evolution: NBs, ori=cat, 1st / 1000 IDdays bef 1st NB ; definite catheter associated NBs, 1st episodes only / 1000 catheterdays

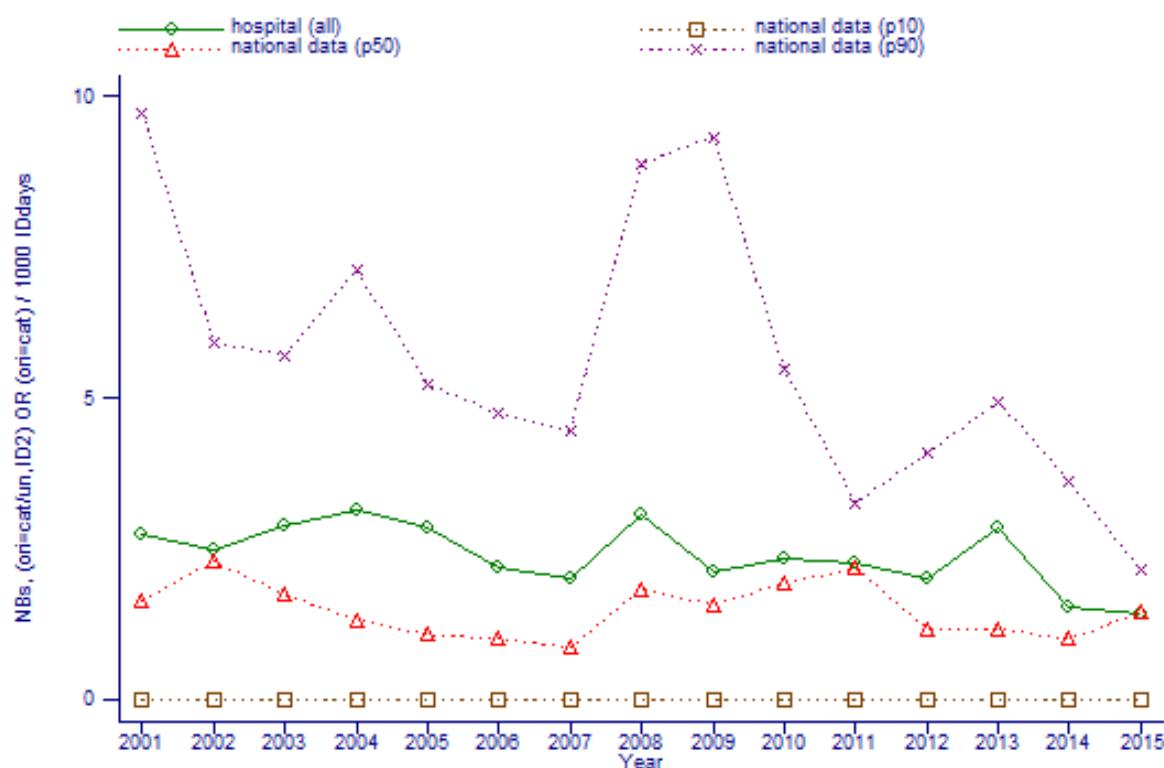


Figure 22: Evolution: NBs, (ori=cat/un, ID2) OR (ori=cat) / 1000 IDdays ; catheter associated NBs / 1000 catheterdays (CDC)

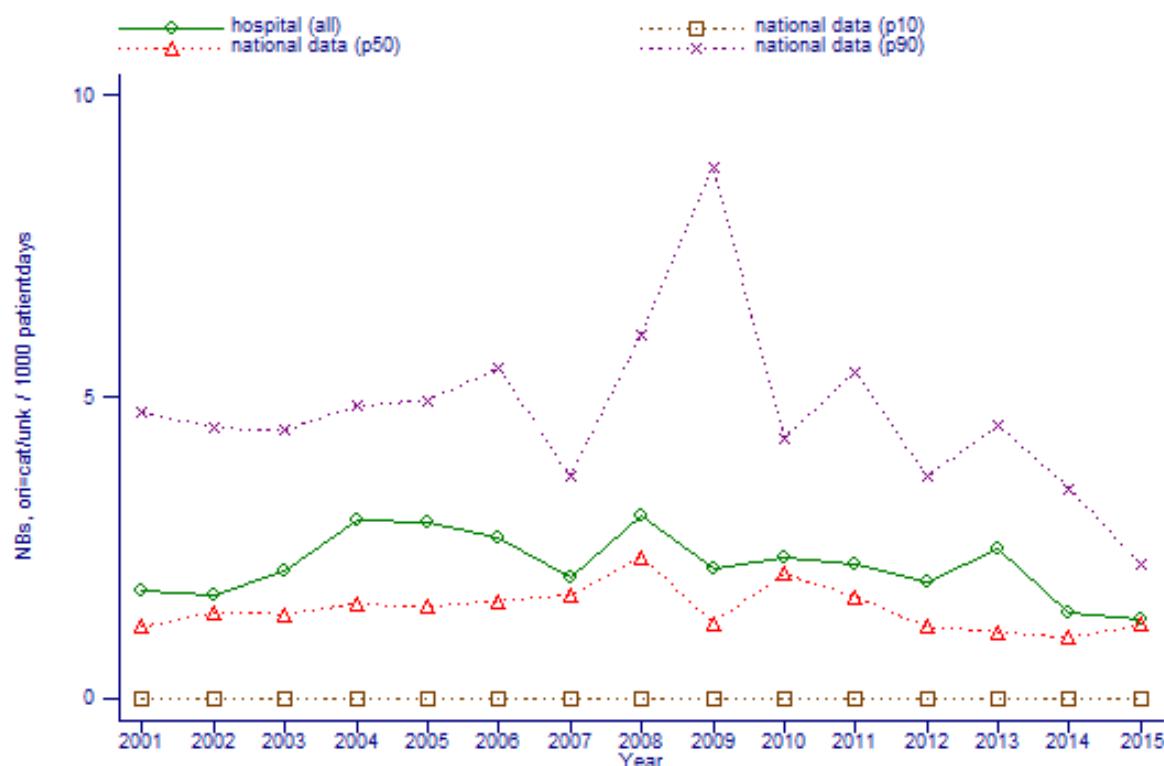


Figure 23: Evolution: NBs, ori=cat/unk / 1000 patientdays ; primary NBs / 1000 patientdays

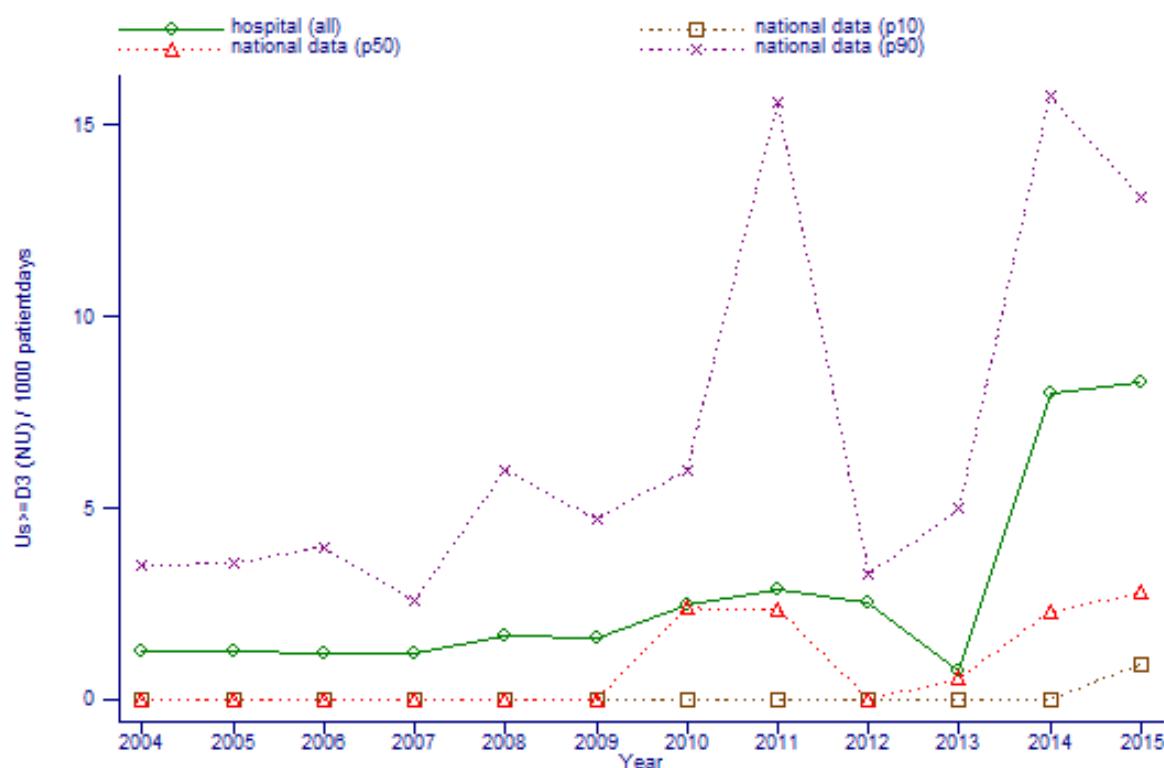


Figure 24: Evolution:  $Us \geq D3 (NU) / 1000$  patientdays ;  $NUs / 1000$  patientdays

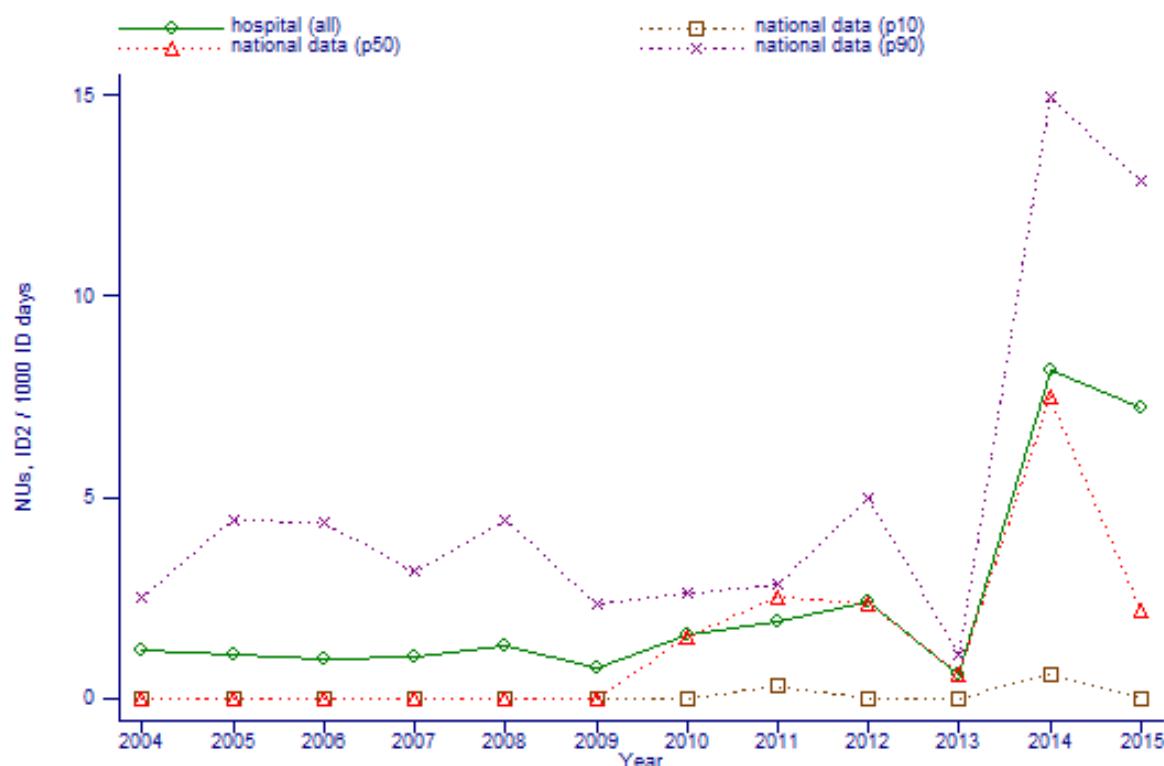


Figure 25: Evolution:  $NUs, ID2 / 1000$  ID days ; catheter associated  $NUs / 1000$  urinary catheterdays