Supplementary Online Content

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This supplementary material has been provided by the authors to give readers additional information about their work.

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eAppendix 1. eMicrobiology Methods

Surveillance cultures

Twice weekly (Monday/Thursday) rectum and respiratory surveillance samples were collected from study patients and inoculated on ESBL-selective agar (chromID, bioMérieux). Species determination and antibiotic susceptibility testing were performed according to local protocols, with a minimum common set of antibiotics tested for GNB (Supplementary eTable 2). Five centers used automated antibiotic susceptibility testing (AST), five relied on disk diffusion and three used both methods (Supplementary eTable 1). Colistin testing was performed at least once per species per tractus (rectum or respiratory) per patient, by automatic testing in five centers and by E-testing (bioMérieux) in eleven centers; three centers used both approaches.

Point prevalence cultures

Rectal swabs and respiratory tract secretions were obtained for point prevalence surveys once monthly, from all patients present in the unit on that day, and inoculated on ESBL- (both samples), VRE- (rectal swabs) and MRSA-selective agar (respiratory tract samples, all chromID, bioMérieux). Growing isolates were processed using the laboratory procedures described above. Once every three months, these samples were also inoculated on plain MacConkey agar, from which a maximum of three morphologically distinct colonies were selected for species determination (MALDI-TOF, Bruker) and automated susceptibility testing (BD Phoenix, BD Diagnostic Systems) at the University Medical Center Utrecht to determine the prevalence of colistin susceptibility among GNB isolated from non-selective media (Supplementary eTable 11).

Clinical cultures

Results from clinical respiratory and blood samples were collected for all study patients. These samples were processed according to local microbiological procedures.

eAppendix 2. Sample Size Calculation

The sample size calculation in the study protocol states that "to demonstrate a 10% relative difference in 28-day mortality for each intervention, 10,800 patients were required (using a baseline 28-day mortality of 27.5%, alpha 0.05, beta 0.80), including a margin of 600 patients per study arm to include cluster-effects and differences in baseline characteristics. The required sample size to detect a 50% reduction of ICU-acquired BSI due to MDRGNB for each intervention (from a baseline incidence of 1.4%) was 6,700."

In fact, a power of 80% instead of a beta of 0.80 was used. .

It was discovered – after study completion - that an error had occurred in the calculation of variance between study groups in the formula which led to lower patient numbers than required. This mistake was partly attenuated by the higher mortality rate observed in the baseline period.

The ICC was determined with the final data with the R-package ICCbin (rm method) and yielded a value of 0.001.

The funding for this study was obtained from a call text that specifically asked for evaluation of interventions in ICU that could reduce the incidence of ICU-acquired BSI. We, therefore, designed our study with this primary endpoint, but decided to power the study on mortality, as this is the more clinically relevant endpoint.

eFigure. Proportion of Surveillance Samples Positive for Antibiotic-Resistant Gram-Negative Bacteria on Day 1-14 of ICU Admission (ESBL-Selective Media)



Positivity of rectal samples

No. of rectum samples															
baseline period	275	558	465	438	378	396	345	299	279	232	238	196	212	180	159
СНХ	239	458	444	444	366	361	322	299	281	226	225	179	185	174	163
SOD	178	476	474	469	368	381	373	341	246	246	234	197	185	186	148
SDD	236	437	446	430	382	304	359	321	246	227	229	202	157	178	169

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eFigure (continued). Proportion of Surveillance Samples Positive for Antibiotic-Resistant Gram-Negative Bacteria on Day 1-14 of ICU Admission (ESBL-Selective Media)



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ICU	Coun -try	no. of beds	Type of ICU (medical /surgical	CHX oral care in standa	CHX oral Standard care in (study pe standa		Randomization sequence (order of study periods)		No. of patients included (% of screened)	prevalence of colonization with 3GC-R	susce testii surve cul	eptibility ng GNB eillance Itures	
			/mixed)	rd care	Start date	Duration (months) ^(d)	2	3	4		Enterobacter iaceae in rectum (standard care)	disk diffu- sion	automate d
1 ^(a)	BE	36	mixed	0.12%	01-Dec- 2013	6	CHX (5.6)	SOD (6)	SDD (6)	900 (14.9%)	19.0%	х	
2 ^(a)	BE	24	mixed	0.20%	01-Jan- 2014	6	CHX (4.6)	SDD (6)	SOD (6)	317 (12.8%)	13.6%		phoenix
3	SL	12	medical	none	06-Jan- 2014	6	SOD (6)	SDD (6)	CHX (6)	168 (22.0%)	7.5%	х	
4	BE	42	mixed	0.20%	01-Mar- 2014	6	SDD (7)	CHX (6)	SOD(6)	1231 (19.4%)	7.2%	х	
5	SP	30	mixed	0.12%	14-Apr- 2014	6	SOD (8.5)	CHX (6)	SDD (6)	1043 (32.7%)	30.0%	х	sensititre
6	ІТ	12	mixed	0.20%	15-May- 2014	6	SDD (6)	SOD (3.5+2. 5)	CHX (6)	598 (76.4%)	23.2%	х	vitek / phoenix
7	PT	10	mixed	0.12%	01-Jun- 2014	14.5	SDD (6)	CHX (6)	SOD (6)	639 ^(b)	5.7%		vitek/mi- croscan
8	BE	15	mixed	0.20%	01-Jul- 2014	6	SOD (6)	SDD (6)	CHX (6)	337 (20.9%)	15.2%	х	vitek
9	BE	42	mixed	0.20%	01-Sep- 2014	6	CHX (6)	SOD (5)	SDD (6)	1297 (28.0%)	18.8%		vitek
10	SP	24	medical	0.12%	01-Nov- 2014	8	SDD (6)	SOD (6)	CHX (6)	375 (18.3%)	35.2%	х	
11	SP	8	mixed	0.12%	03-Nov- 2014	9	CHX (6)	SDD (5)	SOD (6)	237 (23.1%)	11.3%		phoenix
12	UK	22	mixed	none	11-Jan- 2015	8	SOD (6)	CHX (6)	SDD (6)	1109 (44.1%)	8.9%	х	
13	PT	9	mixed	0.20%	19-Jan- 2015	7	SDD (6)	SOD (6)	CHX (6)	414 (53.1%)	0.0% ^(c)		micro- scan

eTable 1. Characteristics of Participating Centers

eTable 1. Characteristics of Participating Centers

Abbreviations: 3GC-R, resistance to third generation cephalosporins (Cefotaxime or Ceftriaxone or Ceftazidime); BE, Belgium, CHX, chlorhexidine mouthwash; GNB, Gram-negative bacteria; IT, Italy; PT, Portugal; SL, Slovenia; SP, Spain; UK, United Kingdom; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination.

a) Oral mucosal lesions occurred during the CHX study period with CHX 2% concentration, which was replaced by CHX 1% for the remainder of the study.

b) Only excluded patients with point prevalence data were registered as part of the screened population.

c) 31/31 rectal samples were negative for 3GC-R; in the respiratory tract the prevalence of 3GC-R Enterobacteriaceae was 2/31 (6.5%).

d) In some ICUs, the baseline period was extended pending authorization to conduct the trial.

eTable 2. Micro-organisms in Positive Blood Cultures That are Not Included in the Study Definition of ICU-Acquired Bloodstream Infection

Study period Number of patients	Baseline n=2251		CHX n=2108		SOD n=2224		SDD n=2082	
	No. of episodes ^a	Column%						
Other ICU-acquired positive blood cultures ^b	106		136		142		148	
Coagulase negative staphylococcus spp.	97	91.5%	117	86.0%	127	89.4%	135	91.2%
Clostridium spp.	0	0.0%	1	0.7%	1	0.7%	1	0.7%
Non-pneumococcal streptococcus spp.	5	4.7%	13	9.6%	8	5.6%	8	5.4%
Micrococcus spp.	0	0.0%	0	0.0%	2	1.4%	1	0.7%
Other	4	3.8%	5	3.7%	4	2.8%	3	2.0%

Abbreviations: BSI, Bloodstream infection; CHX, chlorhexidine mouthwash; ICU, intensive care unit; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination; spp., species.

a) First occurrence of unique species on day 2 of ICU admission onwards.b) Excluding species considered contaminants (Bacillus, Atopobium, Corynebacterium and Propionibacterium species).

eTable 3. Antibiotic Susceptibility Testing and Definition of Highly Resistant Microorganism

Minimum required antibiotic susceptibility testing list	HRMO criteria ^a						
Multidrug resistant Gram-negative bacteria (primary outcome)							
Enterobacteriaceae							
Imipenem or Meropenem	А						
Colistin (for species other than naturally resistant)*	A*						
Cefotaxime or Ceftriaxone or Ceftazidime	A and B****						
Ciprofloxacin**	В						
Gentamicin	В						
Amikacin	В						
Piperacillin or Pip/tazobactam	В						
Trimethoprim-Sulfamethoxazole	В						
Glucose non-fermenting Gram-negative bacteria							
Imipenem or Meropenem or Doripenem	A***						
Colistin	А						
Ceftazidime	В						
Ciprofloxacin**	В						
Gentamicin	В						
Amikacin	В						
Piperacillin or Pip/tazobactam or Ticarcillin	В						
MDRGNB regardless of susceptibility testing result							
Stenotrophomonas spp.							
Achromobacter spp.							
Burkholderia spp.							
Gram-positive HRMO							
Methicillin-resistant S. aureus (MRSA)							
Vancomycin resistant <i>E.faecium/E.faecalis</i> (VRE)							

Abbreviations: HRMO highly resistant micro-organism; spp. Species.

- a) Modified from: A.P. Magiorakos et al., Multidrug-resistant, extensively drug-resistant and pandrug-resistant bacteria: an international expert proposal for interim standard definitions for acquired resistance, Clinical Microbiology and Infection, April 2011.
- A Intermediate/resistant result for an antibacterial agent from any one of the indicated groups of this category is sufficient to define the micro-organism as HRMO.
- B Intermediate/resistant result for antibacterial agents from at least three of the indicated groups in this category is required to define the micro-organism as HRMO.
- * Except intrinsically resistant Proteus spp., Providencia spp., Morganella spp., Serratia spp. and H. alvei; these species are defined as a HRMO when they meet resistance criteria for the other antibiotics listed.
- ** Result from ofloxacin or levofloxacin was used as alternative if ciprofloxacin was not available.
- *** For *Pseudomonas aeruginosa*, resistance to one of the other antibiotic groups was necessary to be reported as carbapenem resistant HRMO.
- **** Resistance to third-generation cephalosporins directly qualified for HRMO and was counted as one of three antibiotic classes for criterium B.

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	baseline	СНХ	SOD	SDD
	8106	7898	8407	8522
ICU admission in prior 30 days	356 (4.4%)	442 (5.6%)	441 (5.2%)	493 (5.8%)
Age (mean, SD)	61.3 (17.7)	60.7 (18.6)	61.3 (18.1)	60.9 (18.6)
Sex (male)	4996 (61.6%)	4928 (62.4%)	5238 (62.3%)	5264 (61.8%)
APACHE II (mean, SD) - 5 hospitals*	16.1 (7.9)	15.4 (7.6)	16.3 (7.9)	16.6 (8.0)
SAPS II (mean, SD) - 8 hospitals**	42.4 (18.6)	43.3 (18.9)	43.8 (18.9)	43.8 (18.7)
ICU-LOS (median, IQR)	4 [2 – 7]	4 [2 – 7]	4 [3-7]	4 [2-7]

<u>Abbreviations</u>: CHX, chlorhexidine mouthwash; ICU, intensive care unit; LOS, length of stay (days); SD, standard deviation; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination.

* 137 missing values.
** 2306 missing values (within these selected hospitals).

eTable 5. All Baseline Characteristics	(Study Population)
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	baseline	СНХ	SOD	SDD
	n = 2251	n = 2108	n = 2224	n = 2082
ICU admission in prior 30 days with inclusion	45 (2.0%)	41 (1.9%)	39 (1.8%)	44 (2.1%)
Age (mean, SD)	62.0 (15.6)	61.4 (15.7)	61.6 (15.7)	62.8 (15.5)
Male gender (%)	1420 (63.1%)	1358 (64.4%)	1439 (64.7%)	1344 (64.6%)
APACHE II (mean, SD) - 5 hospitals	20.3 (8.6)	19.8 (8.2)	20.5 (9.3)	21.8 (8.7)
SAPS II (mean, SD) - 8 hospitals	53.0 (18.0)	54.8 (17.9)	54.4 (17.5)	55.0 (18.0)
Type of ICU-admission				
medical	1464 (65.3%)	1323 (63.0%)	1442 (64.9%)	1385 (66.6%)
trauma with surgery	138 (6.2%)	142 (6.8%)	156 (7.0%)	115 (5.5%)
trauma, no surgery	113 (5.0%)	88 (4.2%)	104 (4.7%)	88 (4.2%)
surgical, scheduled	198 (8.8%)	173 (8.2%)	173 (7.8%)	178 (8.6%)
surgical, unscheduled	328 (14.6%)	374 (17.8%)	346 (15.6%)	314 (15.1%)
surgical, unspecified	10 (0.4%)	8 (0.4%)	3 (0.1%)	2 (0.1%)
Location before ICU admission				
Same hospital	1020 (45.3%)	1032 (49.0%)	1025 (46.1%)	1035 (49.7%)
Another hospital or long term care facility	400 (17.8%)	312 (14.8%)	316 (14.2%)	301 (14.5%)
Home (directly or via emergency room)	831 (36.9%)	764 (36.2%)	883 (39.7%)	746 (35.8%)
If admitted from a hospital or long-term care facility, prior location				
Operating room	379 (27.1%)	409 (30.9%)	371 (28.4%)	360 (27.8%)
Other ICU	229 (16.4%)	221 (16.7%)	221 (16.9%)	176 (13.6%)

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	baseline	СНХ	SOD	SDD	
continued	n = 2251	n = 2108	n = 2224	n = 2082	
If admitted from a hospital or long-term care facility, prior location (continued)					
Acute care ward	726 (52.0%)	656 (49.5%)	642 (49.1%)	710 (54.8%)	
Rehabilitation or long term care facility	63 (4.5%)	38 (2.9%)	74 (5.7%)	49 (3.8%)	
Antibiotic at the time of ICU admission	943 (41.9%)	832 (39.5%)	992 (44.6%)	744 (35.8%)	
Acute illness on ICU-admission	1999 (88.9%)	1888 (89.6%)	2007 (90.2%)	1932 (92.8%)	
Sites of organ failure					
Respiratory illness	1023 (45.5%)	990 (47.0%)	998 (44.9%)	985 (47.3%)	
Cardiovascular illness	828 (36.8%)	811 (38.5%)	835 (37.5%)	792 (38.0%)	
Neurologic illness	686 (30.5%)	674 (32.0%)	615 (27.7%)	603 (29.0%)	
Renal illness	232 (10.3%)	203 (9.6%)	220 (9.9%)	201 (9.7%)	
Hepatic illness	148 (6.6%)	122 (5.8%)	124 (5.6%)	104 (5.0%)	
Metabolic illness	128 (5.7%)	146 (6.9%)	145 (6.5%)	106 (5.1%)	
Hematologic illness	135 (6.0%)	111 (5.3%)	155 (7.0%)	109 (5.2%)	
Other illness	210 (9.3%)	253 (12.0%)	345 (15.5%)	344 (16.5%)	
Charlson comorbidity Index (mean, SD)	2.15 (2.42)	2.38 (2.49)	2.35 (2.42)	2.42 (2.56)	
Charlson comorbidity Index (categories)					
0	738 (32.8%)	631 (29.9%)	653 (29.4%)	626 (30.1%)	
1-2	759 (33.7%)	674 (32.0%)	718 (32.3%)	654 (31.4%)	
3-4	399 (17.7%)	398 (18.9%)	461 (20.7%)	410 (19.7%)	
>4	355 (15.8%)	405 (19.2%)	392 (17.6%)	392 (18.8%)	

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	baseline	СНХ	SOD	SDD
continued	n = 2251	n = 2108	n = 2224	n = 2082
Comorbidities				
Any malignancy (incl. leukemia and lymphoma)	319 (14.2%)	296 (14.0%)	329 (14.8%)	289 (13.9%)
Metastatic solid tumor	108 (4.8%)	129 (6.1%)	104 (4.7%)	143 (6.9%)
Hematologic cancer	71 (3.2%)	66 (3.1%)	80 (3.6%)	61 (2.9%)
Immunodepression (incl. steroids / immunesupression)	190 (8.4%)	223 (10.6%)	193 (8.7%)	206 (9.9%)
Periferal vascular disease	260 (11.6%)	254 (12.0%)	262 (11.8%)	325 (15.6%)
Myocardial infarction	232 (10.3%)	230 (10.9%)	219 (9.8%)	209 (10.0%)
Cerebrovascular disease	225 (10.0%)	200 (9.5%)	177 (8.0%)	199 (9.6%)
Congestive heart failure	340 (15.1%)	425 (20.2%)	469 (21.1%)	443 (21.3%)
Pulmonary disease / chronic respiratory failure	384 (17.1%)	398 (18.9%)	400 (18.0%)	395 (19.0%)
DM without chronic complication	269 (12.0%)	298 (14.1%)	251 (11.3%)	280 (13.4%)
DM with chronic complication	187 (8.3%)	147 (7.0%)	180 (8.1%)	148 (7.1%)
Chronic renal failure	257 (11.4%)	278 (13.2%)	337 (15.2%)	303 (14.6%)
Peptic ulcer disease	139 (6.2%)	168 (8.0%)	189 (8.5%)	145 (7.0%)
Dementia	43 (1.9%)	52 (2.5%)	54 (2.4%)	55 (2.6%)
Hemiplegia / paraplegia	50 (2.2%)	47 (2.2%)	56 (2.5%)	53 (2.5%)
Moderate / Severe liver disease	132 (5.9%)	124 (5.9%)	119 (5.4%)	88 (4.2%)
Mild liver disease	56 (2.5%)	87 (4.1%)	58 (2.6%)	67 (3.2%)
Rheumatologic disease	63 (2.8%)	102 (4.8%)	91 (4.1%)	76 (3.7%)
AIDS/HIV infection	20 (0.9%)	17 (0.8%)	29 (1.3%)	21 (1.0%)

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<u>Abbreviations</u>: AIDS/HIV, acquired immunodeficiency syndrome/human immunodeficiency virus; CHX, chlorhexidine mouthwash; DM, diabetes mellitus; ICU, intensive care unit; incl., including; SD, standard deviation; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination.

eTable 6. Compliance Measures

	СНХ	SOD	SDD
Protocol compliance (monthly unit-wide point prevalence screening)			
Patients admitted	1667	1693	1588
Ventilated (% of admitted) - representing % eligible	722 (43.1%)	708 (41.8%)	640 (40.3%)
Included	693	682	604
Patient received 4 doses of study intervention in past 24h (% of included)	641 (92.5%)	630 (92.4%)	569 (94.2%)
Treatment compliance (amongst included patients)	n = 2108	n = 2224	n = 2082
Treatment compliance (% of included)			
Intervention interrupted for more than > 24h (4 gifts or more) before extubation	54 (2.6%)	82 (3.7%)	100 (4.8%)
Type of medication interrupted			
СНХ	54 (2.6%)	2 (0.1%)	0 (0.0%)
SOD	0 (0.0%)	75 (3.4%)	0 (0.0%)
SDD, paste only	0 (0.0%)	0 (0.0%)	13 (0.6%)
SDD, suspension only	0 (0.0%)	1 (0.0%)	54 (2.6%)
SDD, both paste and suspension	0 (0.0%)	2 (0.1%)	33 (1.6%)
Duration of interruption			
1-3 days	28 (1.3%)	47 (2.1%)	58 (2.8%)
4-7 days	17 (0.8%)	25 (1.1%)	30 (1.4%)
8-14 days	1 (0.0%)	2 (0.1%)	7 (0.3%)
15 days or more	1 (0.0%)	4 (0.2%)	2 (0.1%)
Reason for interruption			
Allergy	0 (0.0%)	1 (0.0%)	0 (0.0%)
Intolerance	15 (0.7%)	2 (0.1%)	1 (0.0%)
Colonization with HRMO, only sensitive to colistin	0 (0.0%)	1 (0.0%)	0 (0.0%)
Colonization with HRMO, that is resistant to carbapenems <i>and</i> tobramycin (or if tobramycin was not tested, to gentamicin)	0 (0.0%)	0 (0.0%)	0 (0.0%)
Gastrointestinal contra-indication	2 (0.1%)	0 (0.0%)	33 (1.6%)
Doctor's decision	4 (0.2%)	17 (0.8%)	8 (0.4%)
Patient's (or proxy) decision	10 (0.5%)	4 (0.2%)	10 (0.5%)

<u>Abbreviations</u>: CHX, chlorhexidine mouthwash; HRMO, highly resistant micro-organism; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination.

Center	Baseline period	СНХ	SOD	SDD
ICU1	45.3%	59.1%	60.3%	54.7%
ICU2	66.0%	74.8%	66.2%	66.3%
ICU3	76.2%	80.7%	71.9%	78.8%
ICU4	67.0%	71.2%	74.1%	76.1%
ICU5	61.4%	70.3%	75.0%	73.7%
ICU6	74.1%	73.3%	64.2%	75.2%
ICU7	64.9%	86.5%	93.6%	65.0%
ICU8	66.3%	70.4%	70.3%	74.6%
ICU9	61.1%	70.3%	75.5%	64.3%
ICU10	58.4%	72.7%	70.6%	65.0%
ICU11	61.7%	67.3%	74.0%	76.6%
ICU12	32.6%	61.7%	48.7%	78.0%
ICU13	86.5%	86.4%	91.1%	87.7%
Overall	64.1%	72.4%	72.5%	72.2%

eTable 7. Average Hand Hygiene Compliance per Study Period

<u>Abbreviations</u>: CHX, chlorhexidine mouthwash; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination.

	b	aseline		СНХ		SOD	:	SDD		Total
Centre	n	incidence n (%)	n	incidence n (%)	n	incidence n (%)	n	incidence n (%)	n	incidence n (%)
ICU1	212	4 (1.9%)	214	7 (3.3%)	245	2 (0.8%)	2 (229)	0.9%	900	15 (1.7%)
ICU2	77	1 (1.3%)	59	0 (0.0%)	101	3 (3.0%)	2 (80)	2.5%	317	6 (1.9%)
ICU3	45	0 (0.0%)	46	0 (0.0%)	36	0 (0.0%)	0 (41)	0.0%	168	0 (0.0%)
ICU4	266	0 (0.0%)	285	0 (0.0%)	334	0 (0.0%)	0 (346)	0.0%	1231	0 (0.0%)
ICU5	169	4 (2.4%)	277	4 (1.4%)	349	3 (0.9%)	2 (248)	0.8%	1043	13 (1.3%)
ICU6	122	9 (7.4%)	177	7 (4.0%)	155	8 (5.2%)	3 (144)	2.1%	598	27 (4.5%)
ICU7	297	0 (0.0%)	109	1 (0.9%)	104	1 (1.0%)	1 (129)	0.8%	639	3 (0.5%)
ICU8	85	2 (2.4%)	92	2 (2.2%)	85	0 (0.0%)	0 (75)	0.0%	337	4 (1.2%)
ICU9	333	9 (2.7%)	338	11 (3.3%)	309	8 (2.6%)	8 (317)	2.5%	1297	36 (2.8%)
ICU10	113	8 (7.1%)	85	1 (1.2%)	85	7 (8.2%)	6 (92)	6.5%	375	22 (5.9%)
ICU11	63	1 (1.6%)	50	2 (4.0%)	70	0 (0.0%)	1 (54)	1.9%	237	4 (1.7%)
ICU12	352	6 (1.7%)	272	2 (0.7%)	248	0 (0.0%)	1 (237)	0.4%	1109	9 (0.8%)
ICU13	117	3 (2.6%)	104	1 (1.0%)	103	1 (1.0%)	0 (90)	0.0%	414	5 (1.2%)
Total	2251	47 (2.1%)	2108	38 (1.8%)	2224	33 (1.5%)	2082	26 (1.2%)	8665	144 (1.7%)

eTable 8. Incidence of ICU-Acquired Bloodstream Infection With Multidrug Resistant Gram-Negative Bacteria (Primary Outcome) per ICU ^a

<u>Abbreviations</u>: CHX, chlorhexidine; n, number of inclusions; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination.

a) Incidence refers to the number of patients with an ICU-acquired bloodstream infection with multidrug resistant Gram-negative bacteria (n=144); there were 154 unique episodes

eTable 9. Systemic Antibiotics Used^a

Study period	Baseline	СНХ	SOD	SDD			
% Patient days included	54%	56%	54%	52%			
Antibiotic type (DDD per 1,000 patient days)							
Penicillin + beta-lactamase inhibitor	402	355	371	422			
Cephalosporins	176	160	180	188			
Carbapenems	179	177	172	186			
Fluoroquinolones	104	99	85	94			
Aminoglycosides	30	26	24	23			
Colistin	24	25	38	36			
Tetracyclins	9	16	25	15			
Macrolides	53	54	58	68			
Lincosamides	25	22	23	23			
Others	56	66	54	60			
Total	1,058	1,000	1,030	1,115			

<u>Abbreviations</u>: CHX, chlorhexidine mouthwash; DDD, defined daily dose; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination.

a) Secondary exploratory analysis

eTable 10. Prevalence of Unit-Wide Carriage of Antibiotic Resistant Microorganisms in Rectum and Respiratory Tract (Complete Results)

	baseline	СНХ		SOD		SDD	
Rectum	prev.	prev.	aRR ª	prev.	aRR ª	prev.	aRR ª
HRMO Enterobacteriaceae	16.1%	21.7%	1.07 (0.99-1.16)	19.7%	1.04 (0.96-1.13)	13.9%	1.05 (0.95-1.16)
third-generation cephalosporin resistance	15.8%	21.5%	1.07 (0.99-1.16)	19.2%	1.04 (0.96-1.13)	13.7%	1.07 (0.97-1.18)
carbapenem resistance	3.2%	3.1%	0.68 (0.54-0.86)	2.9%	0.85 (0.71-1.03)	2.6%	0.80 (0.64-1.01)
resistance to ≥3 antibiotics (or classes)	10.8%	15.5%	1.07 (0.97-1.19)	14.2%	1.06 (0.96-1.17)	10.0%	1.10 (0.97-1.24)
colistin resistance ^b	0.5%	1.6%	0.81 (0.54-1.21)	1.8%	0.97 (0.65-1.45)	1.3%	0.96 (0.60-1.54)
HRMO Glucose non-fermenting GNB	3.2%	3.2%	0.77 (0.62-0.95)	3.3%	0.93 (0.76-1.14)	2.3%	0.81 (0.63-1.04)
carbapenem resistance ^c	2.9%	2.9%	0.75 (0.60-0.93)	2.7%	0.95 (0.76-1.18)	1.8%	0.80 (0.60-1.06)
colistin resistance	0.0%	0.1%		0.1%		0.3%	
resistance to ≥3 antibiotics (or classes)	2.4%	2.0%	0.66 (0.50-0.89)	2.7%	0.82 (0.65-1.05)	1.7%	0.71 (0.52-0.98)
MDRGNB, regardless of antibiotic susceptibility	1.0%	1.5%	0.80 (0.50-1.27)	1.1%	0.80 (0.49-1.30)	1.6%	1.01 (0.64-1.58)
Any MDRGNB (aggregate)	19.3%	25.3%	1.03 (0.96-1.11)	23.0%	1.03 (0.96-1.11)	17.1%	1.04 (0.96-1.14)
VRE	2.2%	1.5%	0.96 (0.74-1.24)	1.8%	0.94 (0.73-1.21)	4.2%	1.03 (0.84-1.27)
Enterobacteriaceae or GNF-GNB with gentamicin resistance ^d	8.3%	9.0%	0.95 (0.84-1.08)	10.4%	0.99 (0.89-1.10)	7.2%	1.00 (0.87-1.15)

eTable 10. Prevalence of Unit-Wide Carriage of Antibiotic Resistant Microorganisms in Rectum and Respiratory Tract (Complete Results) (continued)

	baseline	СНХ		SOD		SDD	
Respiratory tract	prev.	prev.	aRR ª	prev.	aRR ª	prev.	aRR ª
HRMO Enterobacteriaceae	6.6%	7.6%	0.94 (0.81-1.09)	4.2%	0.93 (0.80-1.09)	4.7%	0.94 (0.78-1.13)
third-generation cephalosporin resistance	6.4%	7.4%	0.95 (0.82-1.10)	4.2%	0.93 (0.80-1.09)	4.5%	0.94 (0.78-1.13)
carbapenem resistance	1.4%	1.1%	0.71 (0.47-1.07)	0.9%	0.68 (0.48-0.94)	0.5%	0.59 (0.37-0.97)
resistance to ≥3 antibiotics (or classes)	4.0%	5.2%	1.02 (0.84-1.23)	3.3%	0.92 (0.76-1.12)	3.5%	1.04 (0.83-1.31)
colistin resistance ^b	0.1%	0.8%	0.57 (0.29-1.14)	0.9%	0.66 (0.36-1.21)	0.3%	0.61 (0.30-1.22)
HRMO Glucose non-fermenting GNB	3.4%	2.9%	0.80 (0.64-1.00)	3.8%	0.84 (0.70-1.00)	2.7%	0.75 (0.58-0.96)
carbapenem resistance ^c	3.1%	2.8%	0.80 (0.63-1.00)	3.4%	0.83 (0.69-1.00)	2.4%	0.80 (0.62-1.04)
colistin resistance	0.3%	0.0%		0.1%		0.2%	
MDR	2.5%	1.7%	0.75 (0.55-1.01)	2.6%	0.83 (0.67-1.03)	2.2%	0.76 (0.57-1.01)
MDRGNB, regardless of antibiotic susceptibility	3.8%	5.2%	1.16 (0.94-1.44)	3.2%	0.97 (0.77-1.22)	3.6%	1.04 (0.83-1.31)
Any MDRGNB (aggregate)	12.9%	15.2%	0.98 (0.88-1.08)	10.3%	0.93 (0.84-1.04)	10.2%	0.94 (0.83-1.06)
MRSA	1.7%	1.1%	0.95 (0.66-1.36)	1.3%	0.77 (0.59-1.00)	1.7%	0.73 (0.54-0.97)
Enterobacteriaceae or GNF-GNB with gentamicin resistance ^d	4.5%	4.3%	0.84 (0.69-1.02)	4.4%	0.86 (0.72-1.02)	4.0%	0.85 (0.68-1.06)

<u>Abbreviations</u>: aRR; adjusted relative risk per month; CHX, chlorhexidine; GNB, Gram-negative bacteria; HRMO, highly resistant micro-organism; MDRGNB, multidrug resistant Gramnegative bacteria (resistance to 3 or more antibiotics/classes of antibiotics [(Supplementary eTable 2]);MRSA, methicillin resistant *S. aureus*; SDD, selective digestive decontamination; SOD, selective oropharyngeal decontamination; VRE, vancomycin-resistant *E. faecium/E. faecalis*.

a) all models were corrected for underlying time trends per center.

b) excluding Enterobacteriaceae with intrinsinc colistin resistance (Proteus spp., Morganella spp., Serratia spp., Providencia spp., Hafnia alvei).

c) for Pseudomonas spp., resistance to at least one other antibiotic was required to be reported as carbapenem-R.

d) these were not part of the definition of HRMO.

	Red	ctum	Respiratory tract	
	EB	GNF-GNB	ЕВ	GNF-GNB
Number of unique micro-organisms	1,153	467	559	399
Antibiotics for which AST was requested				
colistin	96.9%	93.8%	97.3%	92.0%
meropenem or imipenem	98.0%	n/a	98.0%	n/a
meropenem or imipenem or doripenem	n/a	97.2%	n/a	94.0%
piperacillin or piperacillin/tazobactam	97.9%	n/a	98.7%	n/a
piperacillin or piperacillin/tazobactam or ticarcillin	n/a	95.9%	n/a	89.2%
cefotaxim or ceftriaxone or ceftazidime	98.0%	n/a	98.6%	n/a
ceftazidime	n/a	95.9%	n/a	89.0%
gentamicin	97.2%	97.0%	98.4%	93.5%
amikacin	98.0%	97.4%	98.7%	93.7%
fluoroquinolone	98.2%	97.2%	98.7%	94.0%
sulfamethoxazole/trimethoprim	97.7%	n/a	98.2%	n/a
no. of AST results missing	204	119	69	218
no. of AST results expected	9183	3736	4362	3192
Completeness of AST	97.8%	96.8%	98.4%	93.2%
Completeness of AST per tractus	97	.5%	96.2	%

eTable 11. Compliance With Antibiotic Susceptibility Testing in Point Prevalence Samples

Abbreviations: AST, antibiotic susceptibility testing; EB Enterobacteriaceae; GNF-GNB Glucose non-fermenting Gram-negative bacteria (including Pseudomonas spp.); no., number.

eTable 12. Prevalence of Colistin-Resistant Gram-Negative Bacteria in 3 Monthly Point Prevalence Surveys (10/13 Centers)

	baseline	СНХ	SOD	SDD			
No. of point prevalence surveys	19	19	18	22			
Number of point prevalence samples							
Rectum	360	409	354	402			
Respiratory tract	360	396	337	377			
Colonization with GNB							
Rectum	256 (71.1%)	289 (70.7%)	253 (71.5%)	245 (60.9%)			
Respiratory tract	112 (31.1%)	146 (36.9%)	85 (25.2%)	105 (27.9%)			
Colonization with colistin-R GNB (excl. intrins	ically colistin resistar	nt species ^a)					
Rectum	4 (1.1%)	3 (0.7%)	8 (2.0%)	8 (2.3%)			
Respiratory tract	1 (0.2%)	3 (0.8%)	3 (0.9%)	0 (0.0%)			
Colonization with <i>intrinsically</i> colistin-R GNB ^a							
Rectum	38 (10.6%)	43 (10.5%)	24 (6.8%)	36 (9.0%)			
Respiratory tract	20 (5.6%)	23 (5.8%)	13 (3.9%)	23 (6.1%)			

Abbreviations: CHX, chlorhexidine mouthwash; GNB, Gram-negative bacteria; no., number; SDD, selective digestive tract decontamination; SOD, selective oropharyngeal decontamination.

<u>Methods</u>: The use of the colistin E-test in eleven hospitals (see supplementary microbiological methods) might have underestimated the prevalence of colistin resistance during monthly point prevalence surveys. Therefore, once every three months, rectum and respiratory samples obtained during point prevalence surveys were inoculated on plain MacConkey agar from which a maximum of three morphologically distinct colonies were selected in the local laboratory. These isolates were shipped to the University Medical Center of Utrecht for species determination (MALDI-TOF, Bruker) and automated susceptibility testing (BD Phoenix, BD) to determine the prevalence of colistin susceptibility among GNB isolated from non-selective media.

^a intrinsically colistin resistant species included Proteus spp., Morganella spp., Serratia spp., Providencia spp., Hafnia alvei.

eTable 13. Results of Sensitivity Analyses

A) Post-hoc sensitivity analysis on the effects of SDD on mortality ^a

	Results from adjusted analysis ^b
	SDD vs. baseline
ICU mortality (aHR, 95% CI)	0.93 (0.79-1.08)
Hospital mortality (aHR, 95% CI)	0.93 (0.79-1.10)
28-day mortality (aOR, 95% CI)	1.01 (0.78-1.29)

Abbreviations: aHR, adjusted hazard ratio; aOR, adjusted odds ratio; CI, confidence interval; ICU, intensive care unit; SDD, selective digestive decontamination.

a) In this analysis all SDD-treated patients with ICU-acquired BSI caused by a pathogen susceptible to third-generation cephalosporins during the first four days and/or with ICUacquired BSI with any pathogen in the absence of mechanical ventilation, were considered alive at all mortality endpoints.

b) all models accounted for clustering using a fixed effect on ICU and a random effect on study period (13 ICUs x 4 study periods) and were adjusted for age, gender, the Charlson comorbidity index, APACHE II or SAPS II score, admission type, antibiotic use on ICU admission, location before ICU admission (in both propensity score and final models) and mean hand hygiene compliance per study period (only in final models).

B) A prior sensitivity analysis: Effects of interventions on mortality, excluding patients with ICU-LOS <3 days.

	Results from adjusted analysis ^a					
	CHX vs. baseline	SOD vs. baseline	SDD vs. baseline			
ICU mortality (aHR, 95% CI)	1.09 (0.96-1.24)	1.04 (0.92-1.19)	0.99 (0.84-1.15)			
Hospital mortality (aHR, 95% CI)	0.99 (0.86-1.15)	1.02 (0.89-1.17)	0.98 (0.84-1.15)			
28-day mortality (aOR, 95% CI)	1.08 (0.87-1.35)	1.06 (0.86-1.32)	1.04 (0.81-1.34)			

Abbreviations: aHR, adjusted hazard ratio; aOR, adjusted odds ratio; CHX, chlorhexidine; CI, confidence interval; HR, hazard ratio; ICU, intensive care unit; OR, odds ratio; SDD, selective digestive decontamination; SOD, selective oropharyngeal decontamination.

a) all models accounted for clustering using a fixed effect on ICU and a random effect on study period (13 ICUs x 4 study periods) and were adjusted for age, gender, the Charlson comorbidity index, APACHE II or SAPS II score, admission type, antibiotic use on ICU admission, location before ICU admission (in both propensity score and final models) and mean hand hygiene compliance per study period (only in final models).

eTable 14. Head-to-Head Comparisons Between Study Interventions; Primary and Secondary Outcomes

	SDD vs. CHX	SOD vs. CHX	SDD vs. SOD
Primary outcome ^(a)			
ICU-acquired BSI with MDRGNB (aHR, 95% CI) ^(b)	0.62 (0.39-0.98)	0.79 (0.53-1.17)	0.79 (0.52-1.19)
Secondary outcomes ^(a)			
ICU-acquired BSI with HRMO (aHR, 95% CI)	0.72 (0.43-1.19)	0.78 (0.55-1.10)	0.92 (0.57-1.48)
ICU-acquired BSI with any pathogen (aHR, 95% CI)	0.73 (0.57-0.93)	0.87 (0.75-1.00)	0.84 (0.67-1.05)
Mortality in ICU (aHR, 95% CI)	0.92 (0.82-1.02)	0.97 (0.90-1.05)	0.94 (0.84-1.06)
Mortality in hospital (aHR, 95% CI)	0.99 (0.90-1.08)	1.03 (0.95-1.12)	0.96 (0.87-1.06)
Mortality at 28th day from ICU admission (aOR, 95% CI)	0.96 (0.82-1.14)	0.98 (0.86-1.12)	0.98 (0.83-1.16)

<u>Abbreviations</u>: aHR, adjusted hazard ratio; aOR adjusted odds ratio; BSI, bloodstream infection; CHX, chlorhexidine; CI, confidence interval; MDRGNB, multidrug resistant Gramnegative bacteria; SDD, selective digestive decontamination; SOD, selective oropharyngeal decontamination.

a) all models accounted for clustering using a fixed effect on ICU and a random effect on study period (13 ICUs x 4 study periods) and were adjusted for age, gender, the Charlson comorbidity index, APACHE II or SAPS II score, admission type, antibiotic use on ICU admission, location before ICU admission (in both propensity score and final models) and mean hand hygiene compliance per study period (only in final models).

b) multidrug resistant Gram-negative bacteria are defined in Supplementary eTable 2.

Study period (comparisons with baseline period)	СНХ	SOD	SDD
Competing events	CSHR (95% CI)	CSHR (95% CI)	CSHR (95% CI)
ICU-acquired BSI with multidrug resistant Gr	am-negative bacteria	l	
ICU discharge	0.97 (0.87-1.07)	1.00 (0.90-1.12)	0.99 (0.88-1.11)
Death in ICU	1.02 (0.90-1.15)	0.99 (0.87-1.12)	0.94 (0.80-1.09)
ICU-acquired BSI with highly-resistant micro-	-organisms		
ICU discharge	0.97 (0.87-1.07)	1.00 (0.90-1.12)	0.99 (0.88-1.11)
Death in ICU	1.01 (0.89-1.15)	0.98 (0.86-1.12)	0.92 (0.79-1.08)
ICU-acquired BSI with any pathogen			
ICU discharge	0.96 (0.87-1.06)	1.01 (0.91-1.12)	0.98 (0.87-1.09)
Death in ICU	1.00 (0.88-1.14)	0.97 (0.85-1.11)	0.93 (0.79-1.09)
Mortality in ICU			
ICU discharge	0.97 (0.88-1.08)	1.01 (0.91-1.12)	1.00 (0.90-1.11)
Mortality in hospital			
Hospital discharge	1.03 (0.93-1.14)	1.10 (1.02-1.18)	1.03 (0.96-1.11)

eTable 15. Results for Competing End Points for Primary and Secondary Outcomes

Abbreviations: CSHR: cause specific hazard ratio; CHX, chlorhexidine; CI: confidence interval; ICU: intensive care unit; SDD, selective digestive decontamination; SOD, selective oropharyngeal decontamination.