

Identification of the clinical most relevant bacteria for blood and blood components

Result of expert consultation meeting organized by the European Centre for Disease Prevention and Control (ECDC)¹

Literature review¹



Step 1

List of confirmed and probable transmissions of bacterial infections from published reviews and original research in the English language

40 bacteria



Responsible for:

213
transmissions

46
fatalities

1. Prioritizing of bacterial infections transmitted through substances of human origin in Europe, D. Domanovic et al, TRANSFUSION 2017;57;1311–1317

40 bacteria



Step 2

Selecting most frequently transmitted bacteria for ranking exercise

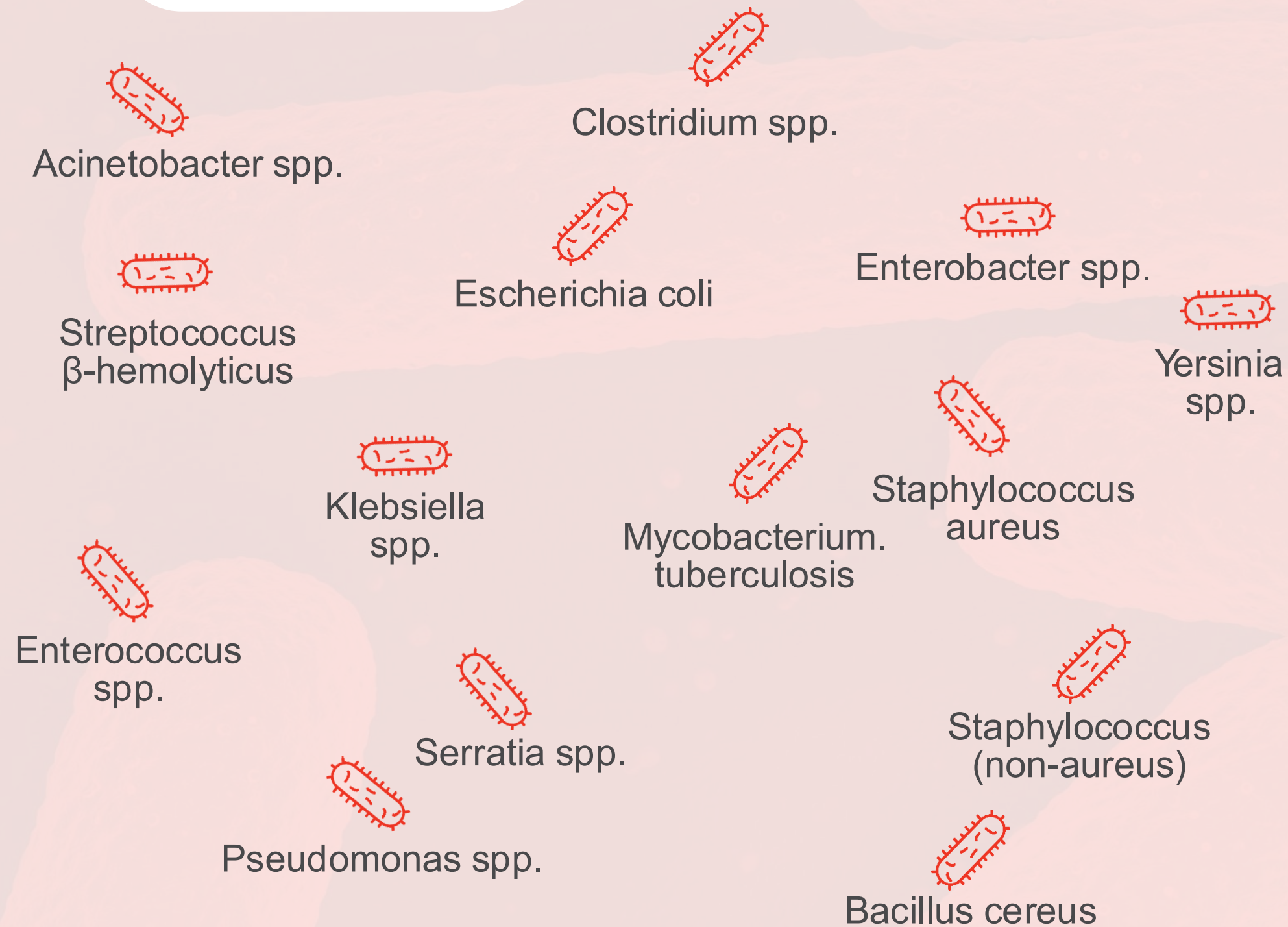




Step 2

Selecting most frequently transmitted bacteria for ranking exercise

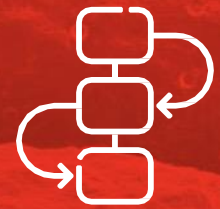
14 bacteria



Responsible for:

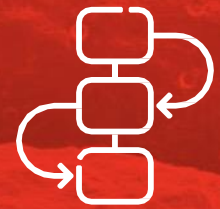
84%
transmissions

93%
fatalities



Step 3 Prioritization of clinically most relevant bacteria

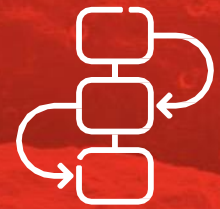
RISK OF TRANSMISSION: reflects an overall prevalence of bacterial transmissions through substances of human origin (SoHO) in the EU adjusted for a given bacteria by a frequency of reported cases of transmission or contamination and survival during processing and storage.



Step 3 Prioritization of clinically most relevant bacteria

RISK OF TRANSMISSION: reflects an overall prevalence of bacterial transmissions through substances of human origin (SoHO) in the EU adjusted for a given bacteria by a frequency of reported cases of transmission or contamination and survival during processing and storage.

SEVERITY OF DISEASE: reflects virulence, toxins production by multiplying during a storage, and survival of a given bacteria after transmission. Health consequences, expressed in disability-adjusted life-year (DALYs) per case, are assessed according to the loss of life, permanent disability, and seriousness of illness.



Step 3 Prioritization of clinically most relevant bacteria

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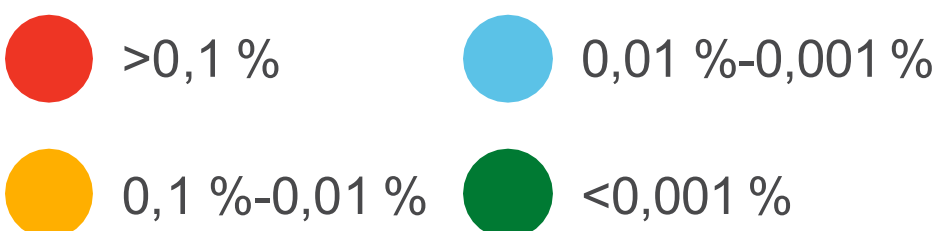
ANTIMICROBIAL RESISTANCE: reflects a probability that a given bacteria is resistant to antimicrobials that is assessed from reported prevalence in the EU, resistance to single or multiple antimicrobials, and trend in the past 4 years developing resistance.



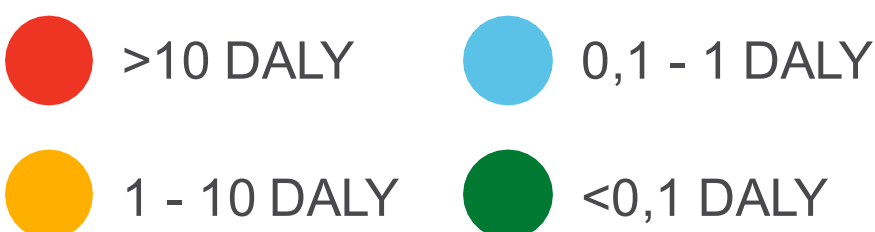
Step 3 Prioritization of clinically most relevant bacteria

Scoring Criteria

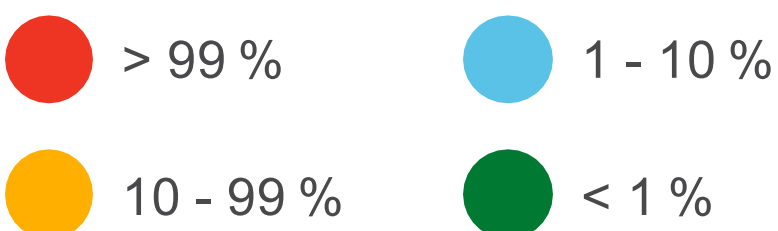
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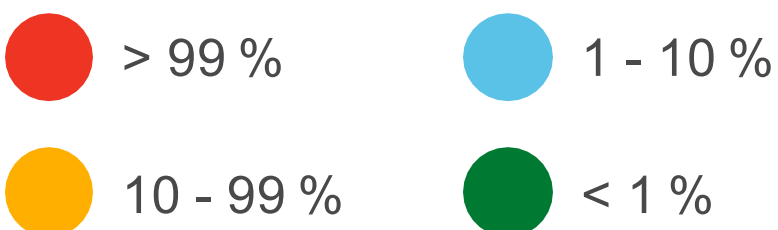
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ANTIMICROBIAL RESISTANCE: reflects a probability that a given bacteria is resistant to antimicrobials that is assessed from reported prevalence in the EU, resistance to single or multiple antimicrobials, and trend in the past 4 years developing resistance.



THREAT EVOLUTION: reflects a probability expressed in the percentage that the magnitude of the threat posed by a given pathogen doubles in the next 5 years.



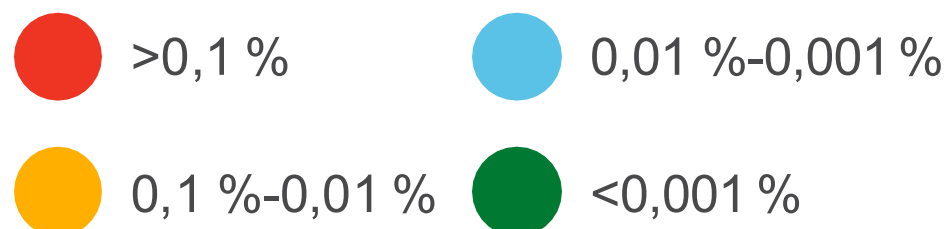
1. Prioritizing of bacterial infections transmitted through substances of human origin in Europe, D. Domanovic et al, TRANSFUSION 2017;57;1311–1317



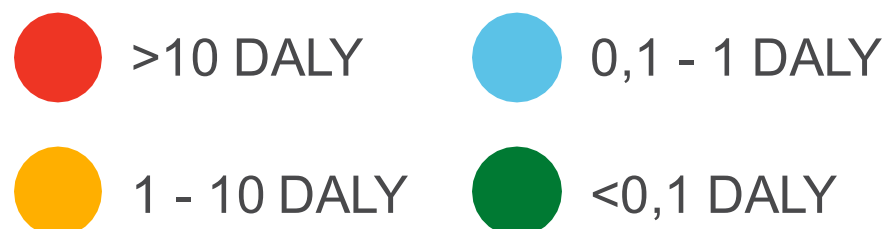
Step 3 Prioritization of clinically most relevant bacteria

Scoring Criteria

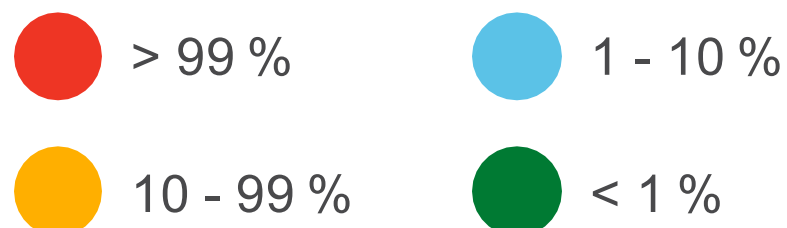
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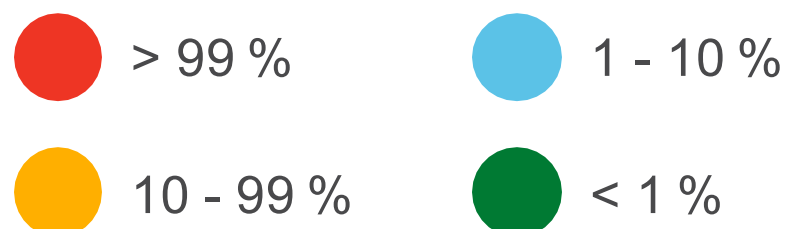
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Weight score

90%

Weight score

10%

1. Prioritizing of bacterial infections transmitted through substances of human origin in Europe, D. Domanovic et al, TRANSFUSION 2017;57;1311–1317

Final result¹



Step 4 Final result ranking exercise

Priority	Ranking	Pathogen	Probability of transmission	Severity of disease	Antimicrobial resistance	Threat evolution
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1. Prioritizing of bacterial infections transmitted through substances of human origin in Europe, D. Domanovic et al, TRANSFUSION 2017;57;1311–1317

 Very high

 Moderately high

 Moderately low

● Very low

Final result¹



Step 4 Final result ranking exercise

Priority	Ranking	Pathogen	Probability of transmission	Severity of disease	Antimicrobial resistance	Threat evolution
Tier 1	1	<i>Staphylococcus aureus</i>				
	2	<i>Streptococcus b-hemolyticus</i>				

1. Prioritizing of bacterial infections transmitted through substances of human origin in Europe, D. Domanovic et al, TRANSFUSION 2017;57;1311–1317



Final result¹



Step 4 Final result ranking exercise

Priority	Ranking	Pathogen	Probability of transmission	Severity of disease	Antimicrobial resistance	Threat evolution
Tier 1	1	<i>Staphylococcus aureus</i>	●	●	●	●
	2	<i>Streptococcus b-hemolyticus</i>	●	●	●	●
Tier 2	3	<i>Klebsiella spp.</i>	●	●	●	●
	4	<i>Escherichia coli</i>	●	●	●	●
	5	<i>Pseudomonas spp.</i>	●	●	●	●
	6	<i>Enterobacter spp.</i>	●	●	●	●
	7	<i>Yersinia enterocolitica</i>	●	●	●	●

1. Prioritizing of bacterial infections transmitted through substances of human origin in Europe, D. Domanovic et al, TRANSFUSION 2017;57;1311–1317



Very high



Moderately high



Moderately low



Very low

Final result¹



Step 4 Final result ranking exercise

Priority	Ranking	Pathogen	Probability of transmission	Severity of disease	Antimicrobial resistance	Threat evolution
Tier 1	1	<i>Staphylococcus aureus</i>	●	●	●	●
	2	<i>Streptococcus b-hemolyticus</i>	●	●	●	●
Tier 2	3	<i>Klebsiella spp.</i>	●	●	●	●
	4	<i>Escherichia coli</i>	●	●	●	●
	5	<i>Pseudomonas spp.</i>	●	●	●	●
	6	<i>Enterobacter spp.</i>	●	●	●	●
	7	<i>Yersinia enterocolitica</i>	●	●	●	●
Tier 3	8	<i>Acinetobacter spp.</i>	●	●	●	●
	9	<i>Staphylococcus (non-aureus)</i>	●	●	●	●
	10	<i>Serratia spp.</i>	●	●	●	●
	11	<i>Clostridium spp.</i>	●	●	●	●

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Final result¹



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Priority	Ranking	Pathogen	Probability of transmission	Severity of disease	Antimicrobial resistance	Threat evolution
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	2	<i>Streptococcus b-hemolyticus</i>	●	●	●	●
Tier 2	3	<i>Klebsiella spp.</i>	●	●	●	●
	4	<i>Escherichia coli</i>	●	●	●	●
	5	<i>Pseudomonas spp.</i>	●	●	●	●
	6	<i>Enterobacter spp.</i>	●	●	●	●
	7	<i>Yersinia enterocolitica</i>	●	●	●	●
Tier 3	8	<i>Acinetobacter spp.</i>	●	●	●	●
	9	<i>Staphylococcus (non-aureus)</i>	●	●	●	●
	10	<i>Serratia spp.</i>	●	●	●	●
	11	<i>Clostridium spp.</i>	●	●	●	●
Tier 4	12	<i>Enterococcus spp.</i>	●	●	●	●
	13	<i>M. tuberculosis</i>	●	●	●	●
	14	<i>Bacillus spp.</i>	●	●	●	●

1. Prioritizing of bacterial infections transmitted through substances of human origin in Europe, D. Domanovic et al, TRANSFUSION 2017;57;1311–1317



Very high



Moderately high



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Final result¹



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Tier 1	1	<i>Staphylococcus aureus</i>	●	●	●	●
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	5	<i>Pseudomonas spp.</i>	●	●	●	●
	6	<i>Enterobacter spp.</i>	●	●	●	●
	7	<i>Yersinia enterocolitica</i>	●	●	●	●
Tier 3	8	<i>Acinetobacter spp.</i>	●	●	●	●
	9	<i>Staphylococcus (non-aureus)</i>	●	●	●	●
	10	<i>Serratia spp.</i>	●	●	●	●
	11	<i>Clostridium spp.</i>	●	●	●	●
Tier 4	12	<i>Enterococcus spp.</i>	●	●	●	●
	13	<i>M. tuberculosis</i>	●	●	●	●
	14	<i>Bacillus spp.</i>	●	●	●	●



“Bacteria ranked in tiers 1 and 2 represent a substantial threat to the safety of blood and blood components.”¹

1. Prioritizing of bacterial infections transmitted through substances of human origin in Europe, D. Domanovic et al, TRANSFUSION 2017;57;1311–1317

INTERCEPT™ Blood System provides robust pathogen inactivation for the clinically most relevant bacteria

ECDC

Published pathogen inactivation data²

Priority	Ranking	Organism	System	PI Data	N° studies ³	Extend of inactivation (log reduction)									
						1	2	3	4	5	6	7	8	9	10
Tier 1	1	<i>Staphylococcus aureus</i>													
	2	<i>Streptococcus pyogenes</i>													
Tier 2	3	<i>Klebsiella pneumoniae</i>													
	4	<i>Escherichia Coli</i>													
	5	<i>Pseudomonas aeruginosa</i>													
	6	<i>Enterobacter cloacae</i>													
	7	<i>Yersinia enterocolitica</i>													
Total number of peer reviewed publications									> 4 log						

1. Mirasol® Pathogen Reduction Technology (PRT) System
2. Peer reviewed publications only, no abstracts
3. References peer reviewed publication, see slide 22

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Priority	Ranking	Organism	System	PI Data	N° studies ³	Extend of inactivation (log reduction)									
						1	2	3	4	5	6	7	8	9	10
Tier 1	1	<i>Staphylococcus aureus</i>	INTERCEPT	✓	4					>5	6,1 6,6				>10
	2	<i>Streptococcus pyogenes</i>	INTERCEPT	✓	2					>5	>6,8				
Tier 2	3	<i>Klebsiella pneumoniae</i>	INTERCEPT	✓	2					>5,6	>6				
	4	<i>Escherichia Coli</i>	INTERCEPT	✓	3						>6,2 >6,4			>9	
	5	<i>Pseudomonas aeruginosa</i>	INTERCEPT	✓	1				4,5						
	6	<i>Enterobacter cloacae</i>	INTERCEPT	✓	1						5,9				
	7	<i>Yersinia enterocolitica</i>	INTERCEPT	✓	2					>5	>5,9				
Total number of peer reviewed publications			INTERCEPT		15				> 4 log						

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	4	<i>Escherichia Coli</i>	INTERCEPT	✓	3						>6,2 >6,4			>9	
	5	<i>Pseudomonas aeruginosa</i>	INTERCEPT	✓	1				4,5						
	6	<i>Enterobacter cloacae</i>	INTERCEPT	✓	1						5,9				
			MIRASOL ¹	✗	0				No peer reviewed data published						
	7	<i>Yersinia enterocolitica</i>	INTERCEPT	✓	2					>5	>5,9				
Total number of peer reviewed publications			INTERCEPT		15				> 4 log						

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Tier 2	3	<i>Klebsiella pneumoniae</i>	INTERCEPT	✓	2					>5,6	>6				
			MIRASOL ¹	✓	2		2,7	2,8							
	4	<i>Escherichia Coli</i>	INTERCEPT	✓	3						>6,2 >6,4			>9	
	5	<i>Pseudomonas aeruginosa</i>	INTERCEPT	✓	1				4,5						
	6	<i>Enterobacter cloacae</i>	INTERCEPT	✓	1						5,9				
			MIRASOL ¹	x	0				No peer reviewed data published						
	7	<i>Yersinia enterocolitica</i>	INTERCEPT	✓	2					>5	>5,9				
			MIRASOL ¹	✓	1			3,3							
Total number of peer reviewed publications			INTERCEPT		15				> 4 log						

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Tier 1	1	Staphylococcus aureus	INTERCEPT	✓	4					>5	6,1	6,6			>10
			MIRASOL ¹	✓	2				4,26	4,8					
	2	Streptococcus pyogenes	INTERCEPT	✓	2					>5		>6,8			
			MIRASOL ¹	✓	2		2,6			>6					
Tier 2	3	Klebsiella pneumoniae	INTERCEPT	✓	2						>5,6	>6			
			MIRASOL ¹	✓	2		2,7	2,8							
	4	Escherichia Coli	INTERCEPT	✓	3						>6,2	>6,4		>9	
			MIRASOL ¹	✓	4				>4,4	5,4	>5,45	>6			
	5	Pseudomonas aeruginosa	INTERCEPT	✓	1				4,5						
			MIRASOL ¹	✓	1					4,7					
	6	Enterobacter cloacae	INTERCEPT	✓	1							5,9			
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	7	Yersinia enterocolitica	INTERCEPT	✓	2					>5	>5,9				
			MIRASOL ¹	✓	1			3,3							
Total number of peer reviewed publications			INTERCEPT		15	> 4 log									
			MIRASOL ¹		12										

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			MIRASOL ¹	✓	2			2,6			>6					
Tier 2	3	Klebsiella pneumoniae	INTERCEPT	✓	2						>5,6	>6				
			MIRASOL ¹	✓	2		2,7	2,8								
	4	Escherichia Coli	INTERCEPT	✓	3						>6,2	>6,4		>9		
			MIRASOL ¹	✓	4				>4,4	5,4	>5,45	>6				
	5	Pseudomonas aeruginosa	INTERCEPT	✓	1				4,5							
			MIRASOL ¹	✓	1					4,7						
	6	Enterobacter cloacae	INTERCEPT	✓	1							5,9				
			MIRASOL ¹	x	0				No peer reviewed data published							
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			MIRASOL ¹	✓	1			3,3								
Total number of peer reviewed publications			INTERCEPT		15	> 4 log										
			MIRASOL ¹		12											

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Mirasol: For ~43 % of the clinically most relevant bacteria the extent of inactivation is “< 4 log” or “no peer reviewed data” have been published

The background of the entire image is a microscopic view of numerous red, rod-shaped bacteria. The bacteria are densely packed in some areas and more sparse in others, creating a textured, organic appearance. They vary in orientation, with some appearing horizontally and others at angles. The lighting gives them a three-dimensional look, with highlights and shadows on their surfaces.

Would you like to know more?

**Ask access to the
INACTIVATION APP**

contact@cerus.com

Pathogen inactivation data - References peer reviewed publication

Priority	Ranking	Organism	INTERCEPT		Mirasol ¹	
			Reference	Extent of inactivation (log reduction)	Reference	Extent of inactivation (log reduction)
Tier 1	1	<i>Staphylococcus aureus</i>	Schmidt M et al., Transfusion, 2015	>5	Kwon SY et al., Vox Sanguinis, 2014	4,26
			Nath Makroo et al., Asian Journal of Transfusion Science, 2018	>6,1	Keil SD et al., J. Vis. Exp., 2015	4,8
			Irsch J et al., Transfusion Medicine and Hemotherapy, 2011	6,6		
			Kwon SY et al., Vox Sanguinis, 2014	>10		
	2	<i>Streptococcus pyogenes</i>	Schmidt M et al., Transfusion, 2015	>5	Keil SD et al., J. Vis. Exp., 2015	2,6
			Irsch J et al., Transfusion Medicine and Hemotherapy, 2011	>6,8	Bello-López et al., Transfusion Clinique et Biologique, 2018	>6
Tier 2	3	<i>Klebsiella pneumoniae</i>	Irsch J et al., Transfusion Medicine and Hemotherapy, 2011	>5,6	Bello-López et al., Transfusion Clinique et Biologique, 2018	2,7
			Schmidt M et al., Transfusion, 2015	>6	Keil SD et al., J. Vis. Exp., 2015	2,8
	4	<i>Escherichia Coli</i>	Nath Makroo et al., Asian Journal of Transfusion Science, 2018	>6,2	Ruane HP et al., Transfusion, 2004	>4,4
			Irsch J et al., Transfusion Medicine and Hemotherapy, 2011	>6,4	Keil SD et al., J. Vis. Exp., 2015	>5,4
			Kwon SY et al., Vox Sanguinis, 2014	>9	Kwon SY et al., Vox Sanguinis, 2014	5,45
					Bello-López et al., Transfusion Clinique et Biologique, 2018	>6
	5	<i>Pseudomonas aeruginosa</i>	Irsch J et al., Transfusion Medicine and Hemotherapy, 2011	4,5	Keil SD et al., J. Vis. Exp., 2015	4,7
	6	<i>Enterobacter cloacae</i>	Irsch J et al., Transfusion Medicine and Hemotherapy, 2011	5,9	No peer reviewed data published	
	7	<i>Yersinia enterocolitica</i>	Schmidt M et al., Transfusion, 2015	>5	Keil SD et al., J. Vis. Exp., 2015	3,3
			Irsch J et al., Transfusion Medicine and Hemotherapy, 2011	>5,9		

