

Supplementary Material

Application of Next-Generation Sequencing to Enterobacter Hormaechei Subspecies Analysis during a Neonatal Intensive Care Unit Outbreak

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Table S1	NEONATES			MOTHERS
	2MRGN-NeoPaed	3 or 4MRGN, MRSA, VRE	SERMA	2, 3 or 4MRGN, VRE, MRSA with negative neonate.
Patient room	Parents must wear DG.	24 h-RI: Parents do not need to wear DG; consequent HD required. No 24 h-RI: Parents must wear DG.	Parents must wear DG. No 24 h-RI allowed.	Parents must wear disposable gowns. No 24 h-RI allowed.
Breastfeeding	yes	yes (certain limitations apply - see Table S2, 4MRGN)	yes (certain limitations apply – see Table S2, 4MRGN)	With 2MRGN-NeoPaed + VRE: yes. With 3MRGN + 4MRGN, MRSA: no.
Kangaroo care	yes	yes (certain limitations apply - see Table S2, 4MRGN)	yes - however not in case of a breakout (see suppl. Table S2, 4MRGN)	With VRE, 2MRGN-NeoPaed + 3MRGN: yes. With 4MRGN + MRSA: no.
When parents leave the patient room	DG, Dispose gown, HD.	24 h-Rooming-in: Parents must put on disposable gowns. No 24h-Rooming in: Dispose gown, hand disinfection.	Dispose Gown, HD. During a breakout: Replace disposable gown by new one.	Dispose Gown, HD. Minimize potential contact points outside the patient room. 3MRGN +4MRGN+MRSA: Replace DG by new one.

Table S1. NICU hygiene measures regarding handling of neonates and mothers colonized with multiresistant bacteria (MRB) and *Serratia marcescens* (SERMA). Breastfeeding of mothers with evidence of MRB/child (still) negative: Since a routine examination of every portion of breast milk was not feasible with reasonable effort in everyday clinical practice, breast milk was preliminary discarded in this scenario. An exemption was made for 2MRGN and VRE. If necessary, breastfeeding was stopped depending on the mother's wishes. Definition of an outbreak scenario: two or more nosocomial infections in which an epidemiologic connection is likely or suspected (§6 Abs. 3, German Infection Protection Act, IfSG). Legend: DG= disposable gown (i.e., longsleeve gown covering upper and lower body), RI= rooming-in, HD= hand disinfection, Legend: MRGN= multidrug-resistant gram-negative bacteria, MRSA= Methicillin-resistant staphylococcus aureus, VRE= Vancomycin-resistant Enterococcus.

Table S2	Neonate with positive bacterial screening				SERMA	
	Multiresistant bacteria					
	2MRGN- NeoPaed	3MRGN, MRSA, VRE	4MRGN			
Breastfeeding	yes	yes	yes, but only after prior education about potential infection risk and sufficient maternal compliance, e.g. hand disinfection		yes - regulation for 4MRGN applies	
Kangaroo care	yes	yes	yes, but only after prior education about potential infection risk and sufficient maternal compliance, e.g., hand disinfection		individual detection: yes outbreak scenario: no	

Table S2. Breastfeeding and Kangaroo care in neonates with positive bacterial screening for multiresistant bacteria (MRB) and *Serratia marcescens* (SERMA). Legend: MRGN= multidrug-resistant gram-negative bacteria, MRSA= Methicillin-resistant staphylococcus aureus, VRE= Vancomycin-resistant Enterococcus. Disposable gown= longsleeve gown covering upper and lower body.

Isolate	Genetic element	Plamid size	Plasmid contig number(s)	MGE (contig number)	AMR genes (contig number)
VA33829	genome				
	p1	2495 bp	1 (circular)		oqxA9B9 (18), fosA (22), blaCT-40 (22)
	p2	4665 bp	2 (circular)		
VA33831	genome				
	p1	2495 bp	1 (circular)		oqxA9B9 (2), fosA (33), blaCT-40 (33)
	p2	3713 + 709 bp	6, 27		
VA33836	genome				
	p1	2346 bp	6		oqxA9B9 (2), fosA (13), blaCT-40 (13)
	p2	4665 bp	1 (circular)		
VA33843	genome			ISKpn34 (43)	
	p1	2495 bp	1 (circular)		oqxA10B15 (11), fosA (29), blaCT-5 (16)
	p2	4665 bp	2 (circular)		
VA34552	genome			ISBras1 (45), ISKpn38 (66)	
	p1	2463 bp	21		oqxA9B9 (13), fosA (16), blaCT-40 (18)
	p2	3635 bp	36		
VA34560	genome				oqxA9B9 (12), fosA (24), blaCT-40 (17)
VA35386	genome				oqxA9B9 (12), fosA (41), blaCT-40 (19)
	p1	595 + 710 + 812 + 1073 bp	17, 16, 4, 21		
VA36175	genome				oqxA10B15 (35), fosA (29), blaCT-5 (17)
	p1	2495 bp	1 (circular)		
	p2	4665 bp	2 (circular)		
VA41244	genome			IS904 (92), IS5708 (97)	oqxA10B5 (52), fosA (60), blaCT-67 (100)
	p1	2494 bp	1 (circular)		
	p2	4307 bp	2 (circular)		
	p3	8814 bp	79, 152		
	p4	14103 bp	46, 157	IS903 (46)	
	p5	21597 bp	86, 99		
VA42547	genome			ISEcl1 (66), ISKpn26 (91), ISKpn34 (135), ISEcl1 (135)	
	p1	2495 bp	66, 72, 91, 107, 135, 146, 153	IS903B (78)	oqxA9B9 (8), blaCT-55 (111)
	p2	5405 bp	1 (circular)		
			2, 56	ISKpn26 (56)	

Table S3. Overview of predicted plasmids. Legend: Plasmid size= either predicted total plasmid size or sum of individual contig sizes likely resembling one plasmid, Contigs= contigs forming the plasmid and whether the assembly is circular. Legend: MGE= mobile genetic elements, AMR= antimicrobial resistance, blaACT gene – coding for a C-type beta-lactamase; fosA gene – coding for a glutathione-S-transferase that inactivates fosfomycin; oqxA gene – coding for a resistance-nodulation-cell division (RND) efflux pump conferring resistance to fluoroquinolone.