

Table S1: Characteristics of NH/NP isolates collected by the CNR-Strep between 2007 and 2013.

Isolate*	Patient	Invasive / non-invasive	EOD / LOD <sup>a</sup>	Source <sup>b,c</sup>	Syndrome <sup>d</sup>	CSP	ST <sup>d</sup>
20130013	neonate	Invasive	EOD	Blood	Bacteremia	Ib	ND
209800983	neonate	Invasive	EOD	Blood	Bacteremia	III	ND
20131298	neonate	Invasive	EOD	Blood	Bacteremia	VI	ND
CCH1330*	neonate	Invasive	EOD	Blood	Bacteremia	III	17
CCH1352*	neonate	Invasive	EOD	CSF <sup>b</sup>	Meningitis	III	17
CCH1322*	neonate	Invasive	LOD	Blood	Bacteremia	Ia	23
CCH1319*	neonate	Invasive	LOD	Blood	Bacteremia	III	17
CCH1320*	neonate	Invasive	LOD	Blood	Bacteremia	III	17
CCH1336*	neonate	Invasive	LOD	Blood	Bacteremia	III	17
CCH1337*	neonate	Invasive	LOD	Blood	Bacteremia	III	27
20130207	neonate	Invasive	LOD	Blood	Bacteremia	III	ND
CCH1340*	neonate	Invasive	LOD	Blood	Bacteremia	IV	459
CCH1318*	neonate	Invasive	LOD	CSF <sup>b</sup>	Meningitis	III	17
CCH1341*	neonate	Invasive	LOD	CSF <sup>b</sup>	Meningitis	III	17
CCH1333*	neonate	Invasive	LOD	CSF <sup>b</sup>	Meningitis	III	17
206800210	neonate	Invasive	LOD	Blood	Bacteremia	III	ND
20130673	neonate	Non-invasive	-	Gastric fluid	Colonization	IV	ND
211800629	neonate	Non-invasive	-	Gastric fluid	Colonization	III	ND
210800492	neonate	Non-invasive	-	Gastric fluid	Colonization	Ia	ND
210800493	neonate	Non-invasive	-	Gastric fluid	Colonization	Ia	ND
20130069	neonate	Non-invasive	-	Stools	Colonization	Ia	ND
CCH1344*	adult	Invasive	-	Blood	Bacteremia	Ib	8
20130718	adult	Invasive	-	Blood	Bacteremia	Ib	ND
20130856	adult	Invasive	-	Blood	Bacteremia	Ib	ND
CCH1351*	adult	Invasive	-	Blood	Bacteremia	II	2
CCH1345*	adult	Invasive	-	Blood	Bacteremia	II	10
CCH1329*	adult	Invasive	-	Blood	Bacteremia	II	12
CCH1343*	adult	Invasive	-	Blood	Bacteremia	II	22
20121061	adult	Invasive	-	Blood	Bacteremia	II	ND
20130897	adult	Invasive	-	Blood	Bacteremia	II	ND
CCH1347*	adult	Invasive	-	Blood	Bacteremia	III	17
CCH1323*	adult	Invasive	-	Blood	Bacteremia	III	27
209800984	adult	Invasive	-	Blood	Bacteremia	III	ND
20121067	adult	Invasive	-	Blood	Bacteremia	III	ND
20121576	adult	Invasive	-	Blood	Bacteremia	III	ND
CCH1348*	adult	Invasive	-	Blood	Bacteremia	IV	459
20131002	adult	Invasive	-	Blood	Bacteremia	IV	ND
CCH1350*	adult	Invasive	-	Blood	Bacteremia	V	1
CCH1328*	adult	Invasive	-	Blood	Bacteremia	V	1
209800638	adult	Invasive	-	Blood	Bacteremia	V	ND
20121276	adult	Invasive	-	Blood	Bacteremia	V	ND
20121410	adult	Invasive	-	Blood	Bacteremia	V	ND
20121597	adult	Invasive	-	Blood	Bacteremia	V	ND

CCH1334*	adult	Invasive	-	Blood	Erysipelas	III	17
CCH1325*	adult	Invasive	-	Blood	Endocarditis	III	1
CCH1324*	adult	Invasive	-	Blood	Endocarditis	V	1
20121165	adult	Invasive	-	Blood	Bone/joint infection	III	ND
207801032	adult	Invasive	-	Bone biopsy	Bone/joint infection	II	ND
CCH1349*	adult	Invasive	-	Bone biopsy	Bone/joint infection	Ib	1
20131427	adult	Invasive	-	Bone biopsy	Bone/joint infection	II	ND
CCH1331*	adult	Invasive	-	Joint fluid	Bone/joint infection	V	1
CCH1338*	adult	Invasive	-	CSF <sup>b</sup>	Meningitis	III	366
CCH1342*	adult	Non-invasive	-	Sperm	Colonization	III	27
20130156	adult	Non-invasive	-	Urine	UTI <sup>c</sup>	III	ND
CCH1327*	adult	Non-invasive	-	Abscess	Abces	II	28
CCH1326*	adult	Non-invasive	-	Skin	Ulcer	III	17
CCH1339*	adult	Non-invasive	-	Vagina	Colonization	Ia	136
CCH1321*	adult	Non-invasive	-	Vagina	Colonization	III	106
CCH1084*	adult	Non-invasive	-	Vagina	Colonization	III	1
20130765	adult	Non-invasive	-	Vagina	Colonization	III	ND
20131184	adult	Non-invasive	-	Vagina	Colonization	III	ND
CCH1346*	adult	Non-invasive	-	Vagina	Colonization	IV	1
20130064	adult	Non-invasive	-	Vagina	Colonization	IV	ND

\*: Sequenced isolates

<sup>a</sup>: EOD/LOD: Early Onset Disease; Late Onset Disease

<sup>b</sup>: Cerebrospinal Fluid

<sup>c</sup>: Urinary Track Infection

<sup>d</sup> : ND Not Determined

Table S2: Bacterial strains used in this study.

<b>Strain</b>	<b>Relevant properties</b>	<b>Source</b>
<b><i>E. coli</i></b>		
TG1	F', traD36, lacIQ, proA+B+, lacZΔ, supE, hsdΔ5	
HB101 / pRK24	Strains for <i>E. coli</i> to GBS conjugation	[1]
NEM3535	HB101 / prK24 + pTCV- <i>erm</i>	This study
NEM3536	HB101 / prK24 + pTCVΩP <sub>cyl+</sub> - <i>abx1</i>	This study
<b><i>S. agalactiae</i></b>		
NEM316	Clinical isolate, Serotype III ST-23	[2]
BM110	Clinical isolate, Serotype III ST-17	[3, 4]
A909	Clinical isolate, Serotype Ia	[5]
2603V/R	Clinical isolate, Serotype V	[6]
515	Clinical isolate, Serotype Ia	[5]
H36B	Clinical isolate, Serotype Ib	[5]
CCH1627	NEM316Δ <i>cylIX</i>	This study
CCH1628	NEM316Δ <i>cylID</i>	This study
CCH1605	NEM316Δ <i>cylG</i>	This study
NEM4190	NEM316Δ <i>cylAB</i>	This study
NEM4192	NEM316Δ <i>cylII</i>	This study
CCH10	NEM316Δ <i>cylE</i>	[7]
CCH1675	NEM316Δ <i>cylJ</i>	[7]
CCH1676	NEM316Δ <i>cylK</i>	[7]
CCH1353	CCH1084ΔIS:: <i>cyl</i>	This study
CCH1620	NEM316+IS:: <i>cyl</i>	This study
NEM3566	CCH1328+pTCV- <i>erm</i>	This study
NEM3567	CCH1328+pTCVΩP <sub>cyl+</sub> - <i>abx1</i>	This study
NEM3568	CCH1331+pTCV- <i>erm</i>	This study
NEM3569	CCH1331+pTCVΩP <sub>cyl+</sub> - <i>abx1</i>	This study
NEM3570	CCH1350+pTCV- <i>erm</i>	This study
NEM3571	CCH1350+pTCVΩP <sub>cyl+</sub> - <i>abx1</i>	This study
NEM3564	CCH1324+pTCV- <i>erm</i>	This study
NEM3565	CCH1324+pTCVΩP <sub>cyl+</sub> - <i>abx1</i>	This study
NEM3244	2603V/R+pTCV- <i>erm</i>	[8]
NEM3249	2603V/R+pTCVΩP <sub>cyl+</sub> - <i>abx1</i>	[8]

Table S3: Plasmids used in this study.

<b>Plasmids</b>	<b>Markers</b>	<b>Relevant properties</b>	<b>Source</b>
pGEM-T easy	Amp	T7 and SP6 promoters	Promega
pG+host5	Ery	ColE1 replicon, thermosensitive derivative of pGK12	[9]
pTCV- <i>erm</i>	Ery, Km	Mob <sup>+</sup> (IncP); <i>oriR</i> pACYC184; <i>oriR</i> pAM_1	[10]
pTCVΩP <sub>cyl+</sub> - <i>abx1</i>	Ery, Km	<i>abx1</i> complementing vector, promoter P <sub>abx1</sub>	[8]
pG+host5ΩΔ <i>cylX</i>	Ery	<i>cylX</i> in-frame deletion vector	This study
pG+host5ΩΔ <i>cylD</i>	Ery	<i>cylD</i> in-frame deletion vector	This study
pG+host5ΩΔ <i>cylG</i>	Ery	<i>cylG</i> in-frame deletion vector	This study
pG+host5ΩΔ <i>cylAB</i>	Ery	<i>cylAB</i> in-frame deletion vector	This study
pG+host5ΩΔ <i>cylII</i>	Ery	<i>cylII</i> in-frame deletion vector	This study
pG+host5ΩΔIS	Ery	IS1381 in-frame deletion vector	This study
pG+host5ΩIS	Ery	Truncated IS1381 in-frame insertion vector	This study

Amp: Ampicillin; Ery: Erythromycin; Km: Kanamycin.

Table S4 : Primers used in this study.

<b>Primer</b>	<b>Sequence (5'-3')*</b>	<b>Gene target</b>
For plasmids constructions		
cylX-5b	aaaGGTACCAggcaaggccattaatttttgagt	$\Delta cylX$
cylX-3b	aaaGGATCCcgcatcaaagaagaacgggtgactt	$\Delta cylX$
cylX-int1	ggcacgcgggtgctgccgcctactcaaaatattagaacg	$\Delta cylX$
cylX-int2	gcggcagcacccggcgtccctattaatggaaaggattgaa	$\Delta cylX$
cylAB-5b	atctGAATTCatagtgaggatatagtgaccaagc	$\Delta cylAB$
cylAB-3b	acatGGATCCtgtttatcatactttgaagcca	$\Delta cylAB$
cylAB-int1	aatgagggaatgacataaaatcaacttaattccatttctgataccaac	$\Delta cylAB$
cylAB-int2	gttggtatcagaaaatggaaattaatggattatgtcattgcctcattc	$\Delta cylAB$
cylD-5b	aaaGAATTCTactcatgtttgtgaattcaac	$\Delta cylD$
cylD-3b	aaaGGATCCcatatctgttcaatgaatc	$\Delta cylD$
cylD-int1	ggcacgcgggtgctgccgcctgaacattaccctgtcctcc	$\Delta cylD$
cylD-int2	gcggcagcacccggcgtccgcaggggaggaaacagttatga	$\Delta cylD$
cylG-5b	aaaGGTACCGtctgaagagcttggcttttg	$\Delta cylG$
cylG-3b	aaaGGATCCcaatcatcaaaacacccggaaac	$\Delta cylG$
cylG-int1	ggcacgcgggtgctgccgcaccaccagtaactattgctac	$\Delta cylG$
cylG-int2	gcggcagcacccggcgtccgtgattgtatggaggaaatgatt	$\Delta cylG$
cylI-5b	acatGAATTCatctgctagattggtttctgg	$\Delta cylI$
cylI-3b	acatGGATCCtgttccacgttttaacatttgcgtt	$\Delta cylI$
cylI-int1	aataatagcaaaagagataccaccttattccactaacaatatacgctc	$\Delta cylI$
cylI-int2	gagcgatatgttagtggataggaggtggatctttgtctattattg	$\Delta cylI$
cyl628-EcoR1	gatatGAATTGaaagatgattgtgaaagggt	$\Delta IS$
cyl628-Pst1	gacaaCTGCAGataagtggcacgacgtt	$\Delta IS$
cyl628-int1	ggatccaaaggaaaggtacccaccaacaaggcgctaaatcg	cyl::IS
cyl628-int2	ggtaccccttggatcccatacgaaatcgacgcaaacgg	cyl::IS
For sequence analysis		
covR_5'	ggttgggttagaagcggtttatcag	covR
covR_3'	ctttccccatggatttaccaatag	covR
covS_5'	gaaagatcgatttatagggtaaaac	covS
covS_3'	caaagcgacataagtaatctc	covS
covS2_5'	tcaagaagacgaaattggagaattg	covS
abx1-fwd	agtttcgcactgaccccttccac	abx1
abx1-rev	tggcgataacttggctatttg	abx1
stk1-F	gctacagtactaacacaag	stk1
stk1-R	ctaggcatggcttaccatagc	stk1
stk1-F2	gttaggtatgtttcattt	stk1
stk1-F3	ctccaaacaagagttcactc	stk1
cylX-5kon	cccgctgcttacttctacaag	cylX
cyl628-5kon	gtgttggaaataatgtccctatag	cylX
cyl1F	caacacttaagtcatgacccatt	cylX
cylD-3kon	gcatcaatagatgttagcttaatc	cylD
cyl628-3kon	gctacttccataactgtttcc	cylD
cyl1.1F	gttcatcgctatgttagc	cylD

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cyl1.1R	ccatctgatgcaaattcttcgc	<i>cylG</i>
cyl2F	gccctagaactgtcaataggga	<i>acpC</i>
cyl1R	ggaactataggacgttgtgg	<i>cylZ</i>
cyl2.2F	cttggacctaattggagctggta	<i>cylA</i>
cyl2.3R	ttagctatcgaaagtgggtc	<i>cylA</i>
cyl2.2R	gcaaagatgcaagagaacatcgctc	<i>cylB</i>
<i>cylEF</i>	gcgtaatgctgtatggagaaaggct	<i>cylB</i>
cyl2R	aggcctaccgtctggaaattga	<i>cylE</i>
<i>cylE.1F</i>	gtcgtatggacaggcaatcatt	<i>cylE</i>
<i>cylE.1R</i>	cctccgatgtatgtgtgaaatcc	<i>cylE</i>
cyl4F	ggaactcaccttatgctacaatgt	<i>cylE</i>
<i>cylER</i>	ggaatcacaaggcttaagg	<i>cylF</i>
cyl4.1F	agcagagtatgggatgaacttga	<i>cylF</i>
cyl4R	gcaggtaatgtttctagagggtga	<i>cylI</i>
cyl4.1R	gctcctagtgtatgtgaagcc	<i>cylI</i>
cyl5F	gcggcaatagaggaacagactgt	<i>cylI</i>
cyl5.1F	tagtggagcgcttgatggta	<i>cylI</i>
cyl5R	gctccatcgataaaactgtttct	<i>cylJ</i>
cyl5.1R	gacggtaacctgtcaatacagc	<i>cylJ</i>
cyl6F	actgacaagctgtggtgaggctat	<i>cylJ</i>
<i>cylKbis</i>	cacatgtgcgagtttagttatctgg	<i>cylK</i>
cyl6R	tgcgtatcttcggcgtgatatg	<i>gbs0657</i>
cyl7F	gatggcttatgctggactaacctc	<i>gbs0657</i>
cyl7R	aaccagacttcctcaggctcact	<i>gbs0658</i>

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\* Restriction sites are in capital letters.

## References cited in Supplementary Tables

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