

Clinical prediction rule for bacteremia among maintenance hemodialysis patients in outpatient settings

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Background

In general population, clinical prediction rules to identify bacteremia were established.

•However, these clinical prediction rules are difficult to use in hemodialysis (HD) patients, since they includes items related to renal function and electrolyte concentration which are affected of the timing of the dialysis. (Eliakim. 2015, Ratzinger. 2014, van Werkhoven. 2014, Takeshima. 2016)

 Further, etiology and pathogen of bacteremia in maintenance HD patients are different from the general population. (Khayr.2003, Nielsen. 2014, Chan.2012, Girndt.2015, Vandecasteele.2009) **Objectives**

We aimed to establish a clinical prediction rule (CPR) for bacteremia among HD patients in outpatients settings. Methods

Study Design: A multi-centre retrospective cohort study

Setting: "Japanese investigators with innovative network about Kidney Disease (JOINT-KD)

six tertiary care institutions and one secondary care institution [August 2011 ~ July 2013]

Participants: Inclusion criteria

(1) HD patients who had any set of blood cultures drawn for suspected infection within 24 hr from first arrival at Hp. (2) age>=18

Discussion

A simple clinical prediction rule specific for HD patients will be helpful for decision making about admission and early ABx administration.

Since, "Fistula First" recent awareness campaign for initiation of AVF as vascular access(VA) in HD induction has been promoted by National Kidney Foundation, in the future, it is expected that AVF becomes mainstream in the world.

Our prediction rule developed in Japanese HD cohort, most of VA are AVF, could be a suitable model for the future hemodialysis patients in the world.

Bacteremia in HD patients, because they have more simple etiology compared as the general population, were considered to be predictable by simple CPR comprised of small items.

Exclusion criteria

(1) low frequency of hemodialysis (<1 time per week), (2) combination of peritoneal dialysis, (3) less than two weeks from the introduction of HD, (4) hospitalized patients referred from another hospital **Outcome:** Bacteremia "Identification of bacteria in blood culture specimen

w/o fulfilling the definition of bacterial contamination"

Candidate predictors: All candidate predictors were selected thorough literature review Statistical analysis:

Derivation set "complete dataset for candidate predictors"

- Description
- Conversion of predictors to binary variables
- Development of CPR :
- (1)Select predictors for bacteremia

CPR1; from predictors among general population using step-wise regression (select items with P<0.05)

CPR2; from predictors among general population & HD patients using stepwise regression

CPR3; from predictors among general population & HD patients with clinical expertise

2 Multivariate logistic regression analysis

(3)Scoring (β -coefficient based)

- <u>Calibration</u>: Hosmer Lemeshow test
- Assessment of test performance
- Validation set "bootstrap method (200 interaction)"
- Internal validation : bootstrap method (200 interaction) Discrimination - C-statistics (95%CI)

CPR1 is considered to be clinically easy to use even with excellent predictive ability.

Limitations

- The reasons to draw blood cultures will be undetected.
- Cases with undetectable bacteremia (blood culture negative) could exist.
- Participants will be restricted to tertiary centers.
- Because of the complete dataset analysis, subjects with relatively mild clinical presentation w/o detailed history taking or laboratory test could had been excluded
- Because it is retrospective cohort study, there is uncertainty of the data extracted from the medical records.
- Relatively small number of samples were enrolled.
- External validation was not performed because of small sample size.

Conclusion

We established simple clinical prediction rule for bacteremia among maintenance hemodialysis patients in outpatients settings

Results Figure 1. Study flow

Table 3. Scoring and calibration

Table 5. Results of internal validation

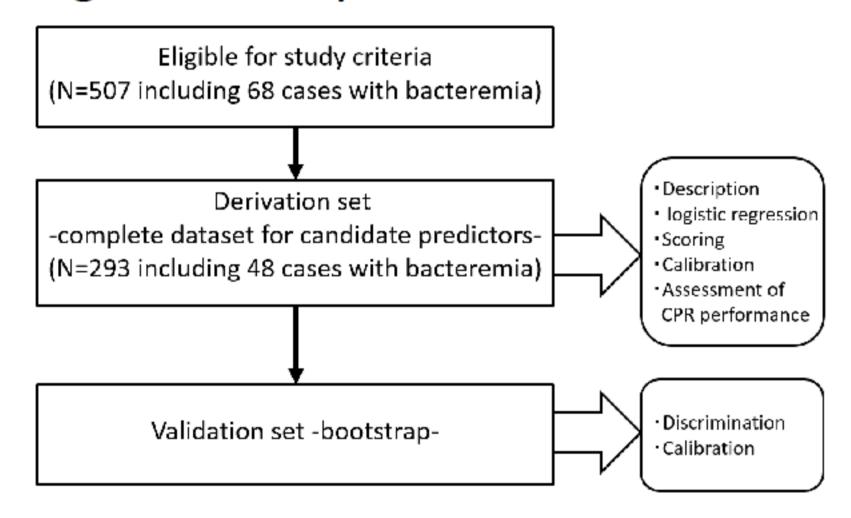


Table 1. Baseline characteristics

				CPR3	Body ter	nperature	≧ 38.3 °C	0.99 0.22,	1.75 0	.01	1		Stud	y facili	
	median (quartile or %)		median (quartile or %)		Pulse rat	te ≧125 /r	min	1.17 0.08,	1.93 0	.04	1		Jua	yracin	
Sex		Medications			CRP ≥ 1	0 mg/dL		1.35 0.67	, 2.0 <0	.01	1		1. Departm	ent of Health	
Male	195 (66.6)	Steroid use	33 (11.3)		No prior	Abx withir	n 1w	1.23 0.13,	2.44 0	.03	1 P=0	0.4	Health in	the Graduate	
Геmale	98 (33.5)	Immunosuppressant use	5 (1.7)	Kyoto, Japan											
Age*	74 (66, 81)	Antibiotics use within 1w*	48 (16.4)	CPR1: ag	e≧65, GCS	5≦14, BT ≧	≧38.3, SBP<90, F	PR≧125, SpO2<	90,Focal a	bdominal	sign, chill, nausea	а,		r Innovative	
Vital signs		Symptoms			-	-	P>360, Na<130,		0, no prio	r antibioti	cs within 1w			ce (CIRC2LE),	
Body temperature*	37.2 (36.6, 38.1)	Chill*	13 (4.4)		were entered into a stephnoe logistie regression models.								3. Division of Nephrology		
Systolic blood pressure*	136 (114, 160)	Nausea*	28 (9.6)	WBC \geq 15000 Pltz15 X 10 ⁴ ALP > 360 Naz130 Albz3 0 CRP \geq 10 no prior aptibiotics within 1w											
Pulse rate*	84 (74, 100)	Focal abdominal sign*	29 (9.9)		AVF, past history of bacteremia, DM were entered into a stepwise logistic regression models. CPR3: age≧65, BT ≧38.3, SBP<90, PR≧125, SpO2<90,chill, nausea, WBC≧15000, CRP≧10, AVF, DM were entered into logistic regression models.							Showa University Fujig 4. Division of Nephrology			
SpO2*	97 (95, 98)	Causes of CKD											University, Kawasaki,		
(FiO2)	0.21 (0.21, 0.21)	Diabetic nephropathy	123 (42.0)	AVF, DIVI											
GCS<15*	45 (15.4)	Hypertensive nephrosclerosis	61 (20.8)												
Vintage of HD (month)	61 (23,112)	Chronic glomerulonephritis	45 (15.4)	Table	Table 4. Assessment of test performance										
Vascular access**		Others and unknown	64 (21.8)				.			10.1.					
AV fistula	245 (83.6)	Laboratory findings		CPR	Cutoff	Total	Bacteremia	Sensitivity (95%Cl)	-	cificity 5%CI)	LR+ (95%Cl)	LR- (95%Cl)	PPV (95%Cl)	NPV (95%Cl)	
AV graft	28 (9.6)	White blood cell (/µL)*	8400 (5900, 11300)		≧1	278	48	100		6.1		0	17.3	100	
Superficial artery	16 (5.5)	Platelet count (/µL)*	14.9 (10.5, 20)		<u></u>	270	40	(92.6, 100)		5, 9.9)	1.1 (1, 1.1)	0	(13, 22.2)	(78.2, 100)	
Permanent Catheter	4 (1.4)	Albumine (mg/dL)*	3.3 (2.9, 3.6)		≧2	162	43	89.6	-	51.4	1.8	0.2	26.5	96.2	
Past history of bacteremia**	31 (10.6)	ALP (IU/L)	271 (212, 382)			101	10	(77.3, 96.5)		, 57.8)	(1.6, 2.2)	(0.1, 0.5)	(19.9, 34)	(91.3, 98.7)	
Comorbidities		Urea nitrogen (mg/dL)	39 (26, 56)	CPR1	≧3	54	22	45.8	8	86.9	3.5	0.6	40.7	89.1	
Diabetes mellitus**	131(44.7)	Sodium (mEq/L)*	137 (135, 139)					(31.4, 60.8)	(82.)	1, 90.9)	(2.3, 5.5)	(0.5, 0.8)	(27.6, 55)	(84.5, 92.8)	
Malignancy	33 (11.3)	C-reacted protein (mg/dl)*	6.1 (1.8, 12.8)		≧4	9	5	10.4	9	98.4	6.4	0.9	55.6	84.9	
		Bacteremia	48 (16.3)					(3.5, 22.7)	(95.9	9, 99.6)	(1.8, 22.9)	(0.8 , 1)	(21.2, 86.3)	(80.2, 88.8)	
* Candidate prec	dictors for bact	teremia among general	population		≧5	0									
** Candidate predictors for bacteremia, specific for HD patients					≧1	278	48	100 (92.6, 100)		6.1 5, 9.9)	1.1 (1, 1.1)	0	17.3 (13, 22.2)	100 (78.2, 100)	
Table 2. Pat	hogens o	of bacteremia			≥2	173	45	93.8 (82.8, 98.7)		17.8 4, 54 . 2)	1.8 (1.6, 2.1)	0.1 (0, 0.4)	26 (19.6, 33.2)	97.5 (92.9, 99.5)	
Bacteria		N Bacteria	N	CPR2	≧3	58	23	47.9 (33.3, 62.8)		35.7 7, 89.8)	3.4 (2.2, 5.1)	0.6 (0.5, 0.8)	39.7 (27, 53.4)	89.4 (84.7, 93)	
Staphylococcus aureus		¹⁹ Enterococcus faecalis	1		≧4	7	4	8.3	ç	8.8	6.8	0.9	57.1	84.6	
Methicillin-sensitive Staphy	lococcus aureus	12 Pseudomonas aerugin	osa 1					(2.3, 20)	(96.)	5, 99.7)	(1.6, 29.4)	(0.9, 1)	(18.4, 90.1)	(79.9, 88.6)	
Methicillin-resistant Staphy	lococcus aureus	⁷ Streptococcus salivariu	s 1		≧5	0									
Klebsiella pneumoniae		⁹ Streptococcus pneumo	niae 1		≧1	271	48	100 (02 6 100)		9.0	1.1	0	17.7	100	
Escherichia coli		7 Streptococcus mutans	1		N -1			(92.6,100)	-	, 13.3)	(1.1, 1.1)		(13.4, 22.8)	(84.6, 100)	
Coagulase-negative staphyle	ococcus species	5 Parabacteroides distas	onis 1	CPR3	≧2	127	37	77.1 (62.7, 88)		53.3 9, 69 . 3)	2.1 (1.7, 2.6)	0.4 (0.2, 0.6)	29.1 (21.4, 37.9)	93.4 (88.5, 96.6)	
Clostridium perfringens		2 Helicobacter cinaedi	1	CFR3	≧3	21	8	16.7 (7.5, 30.2)		94.7 1, 9 7.1)	3.1 (1.4, 7.2)	0.9 (0.8, 1)	38.1 (18.1, 61.6)	85.3 (80.5, 89.3)	
Bacteroides		 Anaerobic gram-negat 2 	ive bacilli ¹		≧4	3	3	6.3	:	100	-	0.9	100	84.5	
Enterococcus faecium		_						(1.3, 17.2)	(90)	5, 100)		(0.9 , 1)	(29.2, 100)	(79.8, 88.5)	

CPR	Selected variables	B-coefficient	95% CI	p-value	Score	Hosmer-Lemeshow χ2test
CPR1	Body temperature \geq 38.3 °C	1.12	0.34, 1.91	<0.01	1	
	Pulse rate \geq 125 /min	1.12	0.01, 2.22	0.04	1	
	CRP ≧10 mg/dL	1.31	0.60, 2.01	<0.01	1	
	ALP > 360 IU/L	1.05	0.35, 1.74	<0.01	1	
	no prior Abx within 1w	1.30	0.15, 2.45	0.03	1	P=0.57
CPR2	Body temperature \geq 38.3 °C	1.40	0.63, 2.17	<0.01	1	
	CRP ≧10 mg/dL	1.35	0.64, 2.06	<0.01	1	
	ALP > 360 IU/L	1.08	0.38, 1.78	<0.01	1	
	No prior Abx within 1w	1.44	0.27, 2.61	<0.01	1	
	Past history of bacteremia	1.22	0.29, 2.14	0.01	1	P=0.11
CPR3	Body temperature \geq 38.3 °C	0.99	0.22, 1.75	0.01	1	
	Pulse rate \geq 125 /min	1.17	0.08, 1.93	0.04	1	
	CRP ≧10 mg/dL	1.35	0.67, 2.0	<0.01	1	
	No prior Abx within 1w	1.23	0.13, 2.44	0.03	1	P=0.4

Vascular access**		Others and unknown	64 (21.8)										
AV fistula	245 (83.6)	Laboratory findings	τ, γ	CPR	Cutoff	Total	Bacteremia	Sensitivity (95%Cl)	Specificity (95%CI)	LR+ (95%Cl)	LR- (95%CI)	PPV (95%Cl)	NPV (95%Cl)
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Superficial artery	16 (5.5)	Platelet count (/µL)*	14.9 (10.5, 20)					(92.6, 100)	(3.5, 9.9)	(1, 1.1)		(13, 22.2)	(78.2, 100)
Permanent Catheter	4 (1.4)	Albumine (mg/dL)*	3.3 (2.9, 3.6)		≧2	162	43	89.6	51.4	1.8	0.2	26.5	96.2
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Comorbidities Urea nitrogen (mg/dL) 39 (26, 56)		CPR1	≧3	54	22	45.8	86.9	3.5	0.6	40.7	89.1		
Diabetes mellitus**	131(44.7)	Sodium (mEq/L)*	137 (135, 139)					(31.4, 60.8)	(82.1, 90.9)	(2.3, 5.5)	(0.5, 0.8)	(27.6, 55)	(84.5, 92.8)
Malignancy	33 (11.3)	C-reacted protein (mg/dl)*	6.1 (1.8, 12.8)		≧4	9	5	10.4	98.4	6.4	0.9	55.6	84.9
	Bacteremia 48 (16.3)						(3.5, 22.7)	(95.9, 99.6)	(1.8, 22.9)	(0.8 , 1)	(21.2, 86.3)	(80.2, 88.8)	
* Candidate predi	ctors for bac	teremia among general	population		≧5	0							
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Table 2. Path	ogens o	of bacteremia			≥2	173	45	93.8	47.8	1.8	0.1	26	97.5
								(82.8 <i>,</i> 98.7)	(41.4, 54.2)	(1.6, 2.1)	(0, 0.4)	(19.6, 33.2)	(92.9 <i>,</i> 99.5)
Bacteria		N Bacter	ia N	CPR2	≧3	58	23	47.9	85.7	3.4	0.6	39.7	89.4
		N Bacter	1	CPR2		58 7	23	47.9 (33.3, 62.8)	85.7 (80.7, 89.8)	3.4 (2.2, 5.1)	0.6 (0.5, 0.8)	39.7 (27, 53.4)	89.4 (84.7, 93)
Bacteria Staphylococcus aureus Methicillin-sensitive <i>Staphyle</i>	ococcus aureus	10	1a 1	CPR2	≧3 ≧4	58 7	23 4	47.9	85.7	3.4	0.6	39.7	89.4
Staphylococcus aureus		¹⁹ Enterococcus faecalis	1 nosa 1	CPR2		58 7 0	23 4	47.9 (33.3, 62.8) 8.3	85.7 (80.7, 89.8) 98.8	3.4 (2.2, 5.1) 6.8	0.6 (0.5, 0.8) 0.9	39.7 (27, 53.4) 57.1	89.4 (84.7, 93) 84.6
Staphylococcus aureus Methicillin-sensitive <i>Staphylo</i> Methicillin-resistant <i>Staphylo</i>		 ¹⁹ Enterococcus faecalis 12 Pseudomonas aerugi 7 	1 nosa 1 ius 1	CPR2	≧4	7	23 4 48	47.9 (33.3, 62.8) 8.3 (2.3, 20) 100	85.7 (80.7, 89.8) 98.8 (96.5, 99.7) 9.0	3.4 (2.2, 5.1) 6.8 (1.6, 29.4) 1.1	0.6 (0.5, 0.8) 0.9	39.7 (27, 53.4) 57.1 (18.4, 90.1) 17.7	89.4 (84.7, 93) 84.6 (79.9, 88.6) 100
Staphylococcus aureus Methicillin-sensitive <i>Staphyle</i> Methicillin-resistant <i>Staphyle</i> Klebsiella pneumoniae		 ¹⁹ Enterococcus faecalis 12 Pseudomonas aerugi 7 Streptococcus salivar 	nosa 1 ius 1 oniae 1	CPR2	≧4 ≧5 ≧1	7 0 271	4	47.9 (33.3, 62.8) 8.3 (2.3, 20) 100 (92.6,100)	85.7 (80.7, 89.8) 98.8 (96.5, 99.7) 9.0 (5.7, 13.3)	3.4 (2.2, 5.1) 6.8 (1.6, 29.4) 1.1 (1.1, 1.1)	0.6 (0.5, 0.8) 0.9 (0.9, 1)	39.7 (27, 53.4) 57.1 (18.4, 90.1) 17.7 (13.4, 22.8)	89.4 (84.7, 93) 84.6 (79.9, 88.6) 100 (84.6, 100)
Staphylococcus aureus Methicillin-sensitive <i>Staphylo</i> Methicillin-resistant <i>Staphylo</i> Klebsiella pneumoniae Escherichia coli	ococcus aureus	 ¹⁹ Enterococcus faecalis 12 Pseudomonas aerugi 7 Streptococcus salivar 9 Streptococcus pneum 	1 nosa 1 ius 1 oniae 1 s 1		≧4 ≧5	7 0	4	47.9 (33.3, 62.8) 8.3 (2.3, 20) 100	85.7 (80.7, 89.8) 98.8 (96.5, 99.7) 9.0	3.4 (2.2, 5.1) 6.8 (1.6, 29.4) 1.1	0.6 (0.5, 0.8) 0.9 (0.9, 1)	39.7 (27, 53.4) 57.1 (18.4, 90.1) 17.7	89.4 (84.7, 93) 84.6 (79.9, 88.6) 100
<i>Staphylococcus aureus</i> Methicillin-sensitive <i>Staphylo</i>	ococcus aureus	 ¹⁹ Enterococcus faecalis 12 Pseudomonas aerugi 7 Streptococcus salivar 9 Streptococcus pneum 7 Streptococcus mutan 	1 nosa 1 ius 1 oniae 1 s 1	CPR2	≧4 ≧5 ≧1	7 0 271	4	47.9 (33.3, 62.8) 8.3 (2.3, 20) 100 (92.6,100) 77.1 (62.7, 88) 16.7	85.7 (80.7, 89.8) 98.8 (96.5, 99.7) 9.0 (5.7, 13.3) 63.3 (56.9, 69.3) 94.7	3.4 (2.2, 5.1) 6.8 (1.6, 29.4) 1.1 (1.1, 1.1) 2.1 (1.7, 2.6) 3.1	0.6 (0.5, 0.8) 0.9 (0.9, 1) 0 0.4 (0.2, 0.6) 0.9	39.7 (27, 53.4) 57.1 (18.4, 90.1) 17.7 (13.4, 22.8) 29.1 (21.4, 37.9) 38.1	89.4 (84.7, 93) 84.6 (79.9, 88.6) 100 (84.6, 100) 93.4 (88.5, 96.6) 85.3
Staphylococcus aureus Methicillin-sensitive Staphylo Methicillin-resistant Staphylo Klebsiella pneumoniae Escherichia coli Coagulase-negative staphyloo	ococcus aureus	 ¹⁹ Enterococcus faecalis 12 Pseudomonas aerugi 7 Streptococcus salivari 9 Streptococcus pneum 7 Streptococcus mutani 5 Parabacteroides dista 	1 nosa 1 ius 1 oniae 1 s 1 isonis 1 1		≧4 ≧5 ≧1 ≧2	7 0 271 127	4 48 37	47.9 (33.3, 62.8) 8.3 (2.3, 20) 100 (92.6,100) 77.1 (62.7, 88)	85.7 (80.7, 89.8) 98.8 (96.5, 99.7) 9.0 (5.7, 13.3) 63.3 (56.9, 69.3)	3.4 (2.2, 5.1) 6.8 (1.6, 29.4) 1.1 (1.1, 1.1) 2.1 (1.7, 2.6)	0.6 (0.5, 0.8) 0.9 (0.9, 1) 0 0.4 (0.2, 0.6)	39.7 (27, 53.4) 57.1 (18.4, 90.1) 17.7 (13.4, 22.8) 29.1 (21.4, 37.9)	89.4 (84.7, 93) 84.6 (79.9, 88.6) 100 (84.6, 100) 93.4 (88.5, 96.6)

CPR	C-statistics	95%CI			
CPR1	0.77	0.70, 0.83			
CPR2	0.77	0.68, 0.83			
CPR3	0.73	0.64, 0.79			

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