Online Resource 5: Screening results differentiated by different methods to treat missing values

1. Method

Studies on the predictive ability of screening tools use a variety of methods to handle EMS' documentation deficiencies. To allow comparability between studies, we report screening results using different methods of handling missing values for screening relevant variables such as temperature or heart rate:

- <u>Method #1:</u> All missing values were considered as normal/healthy values
- <u>Method #2:</u> Eligible cases had to have at least two screening-relevant variables documented in *all* screening tools alike
- <u>Method #3:</u> Eligible cases had to have at least two variables filled which are necessary for the *respective* screening tool
- <u>Method #4:</u> Missing values were imputed using multiple imputation (see Appendix 3) and main paper

All methods enabled EMS cases to reach a score threshold that either equals positive or negative screening results.

The sepsis guidelines mention qSOFA, MEWS, SIRS and NEWS2 for adults [1], whereas the pediatric guidelines do not recommend any specific tool [2]. Thus, the following results base on persons age \geq 18 only.

2. Results

The following results base on the linked EMS + health claims data set (dataset #3) that allowed analyses of the screening tools' ability to predict an inpatient sepsis. Independent of how missing values were treated, qSOFA had the highest specificity and positive predictive value, while NEWS2 had the highest sensitivity, negative predictive value and area under the ROC curve (AUROC) of all tools (see Tab. 1).

Tab. 1: Screening results with qSOFA, MEWS,	SIRS and NEWS2 di	lifferentiated by metho	od for treating mis	ssing values (v	arying sample sizes, l	based on
linked dataset #3; patient age ≥ 18 years)		-	-			

	Method #1				Method #2				Method	I #3			Method #4			
	Cases with missing values treated as "healthy" values (n=4,979)				Cases with at least two screening-relevant variables per tool filled in <i>all</i> screening tools (n=2,061)				Cases v screening for the <i>r</i>	with at leang-releva	ast two nt variabl e screenir	es filled ng tool	Cases with imputed data (n=4,503)			
	qSOFA	MEWS	SIRS	NEWS2	qSOFA	MEWS	SIRS	NEWS2	OSb 4 (n= 3,873)	B S (n= 4,235)	≌ິສ ທ (n= 2,101)	B (n= 4,258)	qSOFA	MEWS	SIRS	NEWS2
% of positive screening results per respective tool	2.3% [1.9; 2.7]	6.6% [5.9; 7.3]	3.7% [3.1; 4.2]	15.5% [14.5; 16.5]	4.0% [3.1; 4.8]	11.6% [10.2; 13.0]	8.6% [7.4; 9.8]	22.0% [20.2; 23.8]	3.0% [2.4; 3.5]	7.8% [7.0; 8.6]	8.7% [7.5; 9.9]	18.1% [17.0; 19.3]	3.7% [3.2; 4.3]	12.5% [11.6; 13.5]	6.1% [5.4; 6.8]	19.4% [18.3; 20.6]
Sensitivity	18.4%	37.9%	21.8%	63.2%	26.0%	56.0%	38.0%	84.0%	23.2%	44.0%	36.5%	73.3%	23.1%	48.7%	28.2%	73.1%
(Se; %)	[17.3; 19.5]	[36.6; 39.3]	[20.7; 23.0]	[61.9; 64.6]	[24.1; 27.9]	[53.9; 58.1]	[35.9; 40.1]	[82.4; 85.6]	[21.9; 24.5]	[42.5; 45.5]	[34.5; 38.6]	[72.0; 74.7]	[21.8; 24.3]	[47.3; 50.2]	[26.9; 29.5]	[71.8; 74.4]
Specificity	98.0%	93.9%	96.7%	85.4%	96.6%	89.5%	92.1%	79.6%	97.4%	92.9%	92.0%	82.9%	96.6%	88.1%	94.3%	81.6%
(Sp; %)	[97.6; 98.4]	[93.3; 94.6]	[96.2; 97.2]	[84.4; 86.3]	[95.8; 97.4]	[88.2; 90.8]	[90.9; 93.3]	[77.8; 81.3]	[96.9; 97.9]	[92.1; 93.7]	[90.9; 93.2]	[81.8; 84.0]	[96.1; 97.1]	[87.2; 89.1]	[93.6; 95.0]	[80.4; 82.7]
Positive predictive value	13.9%	10.0%	10.4%	7.1%	15.9%	11.7%	10.7%	9.3%	13.9%	10.0%	10.4%	7.1%	10.7%	6.7%	8.0%	6.5%
Negative predictive value	98.5%	98.8%	98.6%	99.2%	98.1%	98.8%	98.4%	99.5%	98.6%	98.9%	98.3%	99.4%	98.6%	99.0%	98.7%	99.4%
Area under the ROC curve	0.582	0.659	0.593	0.743	0.613	0.728	0.650	0.818	0.603	0.684	0.643	0.781	0.598	0.684	0.613	0.773

(AUROC, Scores dichotomou s)	[0.514 ; 0.650]	[0.591 ; 0.728]	[0.525 ; 0.660]	[0.683 ; 0.803]	[0.522 ; 0.704]	[0.643 ; 0.812]	[0.561 ; 0.740]	[0.758 ; 0.877]	[0.526 ; 0.680]	[0.612 ; 0.757]	[0.555 ; 0.731]	[0.723 ; 0.840]	[0.526 ; 0.670]	[0.615 ; 0.753]	[0.541 ; 0.684]	[0.716 ; 0.831]
Positive Likelihood Ratio (LR+)	9.1	6.3	6.6	4.3	7.6	5.3	4.8	4.1	8.9	6.2	4.6	4.3	6.8	4.1	5.0	4.0
Negative Likelihood Ratio (LR-)	0.8	0.7	0.8	0.4	0.8	0.5	0.7	0.2	0.8	0.6	0.7	0.3	0.8	0.6	0.8	0.3
% of positive cases in <i>all</i> four screenings tools	0.5% n <i>all</i> ngs (Se: 5.7%; Sp: 99.6%; PPV: 20.8%; NPV: 98.3%, AUROC: 0.527)				1.2% (Se: 10.0%; Sp: 99.1%; PPV: 20.8%; NPV: 97.8%; AUROC: 0.545)			Not app sample	licable du sizes per	ue to diffe tool	erent	0.8% (Se: 7.7%; Sp: 99.3%; PPV: 16.7%; NPV: 98.4%, AUROC: 0.535)				
% of positive cases in <i>any</i> of the four screenings tools	17.4% (Se: 65 6.6%; N 0.745)	.5%; Sp: / IPV: 99.3	83.5%; P %, AURC	PV: DC:	26.2% [Se: 88.0%; Sp: 75.4%, PPV: 8.2%; NPV: 99.6%; AUROC: 0.817)				Not app sample	licable du sizes per	ue to diffe · tool	erent	24.2% (Se: 76.9%; Sp: 76.7%; PPV: 5.5%; NPV: 99.5%, AUROC: 0.768)			

References

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