TABLE S1 Statistical significance when comparing the results from the different detection methods

Method (n=no. of	No.	No.	Significant difference Fisher's exact test (p-value)				
labs)	VME/	ME/	VME and ME		VME		
	total R	total S	versus disk diffusion	versus agar screen	versus disk diffusion	versus agar screen	
EUCAST disk diffusion (n=28)	53/756	2/84	-	-	-	-	
CLSI agar screen (n=18)	32/486	3/54	NS	-	NS	-	
VITEK 2 (n=5)	18/135	0/15	SD (p=0.0263)	SD (p=0.0360)	SD (p=0.0231)	SD (p=0.0186)	

VME; classified as S when containing *van* genotype, ME; classified as R when containing no *van* genotype, NS; no significant difference, SD; significant difference.

Table S2 Number of very major (VME) and major errors (ME) calculated for EUCAST disk diffusion and CLSI agar screen divided into both experience with method and type of agar used for $n \ge 5$

Method	Media (no. of labs)	VME in % (IE/E)	No. VME/ total no. R IE, E	ME in % (IE/E)	No. ME/ total no. S IE, E
Disk diffusion	Oxoid MH (4IE/12E) BBL MH II (9IE/1E)	2.9 (7.4/1.9) 14 (14/11)	8/108, 6/324 34/243, 3/27	0 6.7 (7.4/0)	2/27, 0/3
Agar screen	BHI Difco (7E/1IE) BHI Oxoid (5E)	4.2 (4/4) 11	8/189, 1/27 15/135	0 0	

VME; classified as S when containing *van* genotype, ME; classified as R when containing no *van* genotype, E; experienced, IE; inexperienced, NS; no significant difference, SD; significant difference.

Table S3 Statistical significance when comparing results by both experience with method and type of agar used for $n \ge 5$

Method	Media (no. of	No.	No.	Significant difference Fisher's exact test (p-value)			
	labs)	VME/	ME/	VME and ME		VME versus IE	
		total	total	versus Oxoid MH	versus BBL MH II	versus Oxoid MH	versus BBL MH II
		no. R	no. S	IE	IE	IE	IE
EUCAST	BBL MH II (9IE)	34/243	2/27	NS	-	NS	-
disk	BBL MH II (1E)	3/27	0/3	-	NS	-	NS
diffusion	Oxoid MH (4IE)	8/108	0/12	-	-	-	-
	Oxoid MH (12E)	6/324	0/36	SD (p=0.0093)	-	SD (p=0.0092)	-
				versus Difco BHI E		versus Difco BHI E	
CLSI agar	BHI Difco (7E)	8/189	0/21	-		-	
screen	BHI Oxoid (5E)	15/135	0/15	SD (p=0.0270)		SD (p=0.0264)	

VME; classified as S when containing van genotype, ME; classified as R when containing no van genotype, E; experienced, IE; inexperienced,

NS; no significant difference, SD; significant difference.