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Spotting method as a high throughput alternative to the conventional spread plating method for bacterial time-kill experiments.

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Introduction

In vitro time-kill experiments are commonly performed to obtain information about the dynamics of a drug antimicrobial activity. It is however labor intensive due to the conventional spread plating method used to count bacteria. In this work we compare a spotting method with the conventional spread plating method.

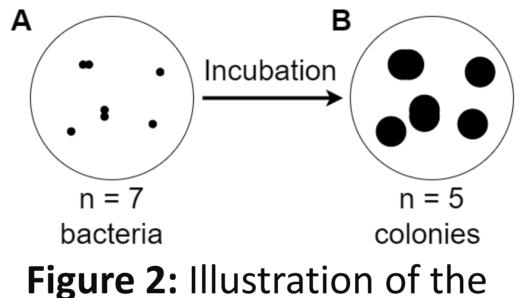
Materials and Methods Conventional 4 strains: plating (n=3) 2 Pseudomonas 25 µL per third of aeruginosa plate Escherischia coli 1 Klebsiella pneumoniae For each strain For each For each For each and antibotic concentration dilution sample X 6 0*MIC 4*MIC 3 samples at 6 4 antibiotics: Spotting (n=3) times between 0 Antibiotic Ceftazidime 1 μL per spot and 30h concentrations Ciprofloxacin (n=6)Colistin Serial dilutions (n=8) Tobramycin Figure 1: Experimental protocol.

Correction of colony counts to account for overcrowding

- Colony forming unit counts were converted to bacterial densities after correcting for overcrowding (Figure 2) by the « most probable number » method from Martini et al.(1)
- Under this method, for a given measurement, the probability of overcrowding is higher when the number of counted colonies (**Figure 3**) is close to the maximal number of countable colonies.

Validation of the spotting method

 To validate the spotting method against the conventional plating method we applied correlated bivariate least square regression developed by Francq and Govaerts (2) on log₁₀ transformed bacterial densities after exclusion of measurements below 10³ CFU/mL (limit of detection of the spotting method)



overcrowding problem.

A. Number of bacteria at time of plating.

B. Number of colonies after

incubation

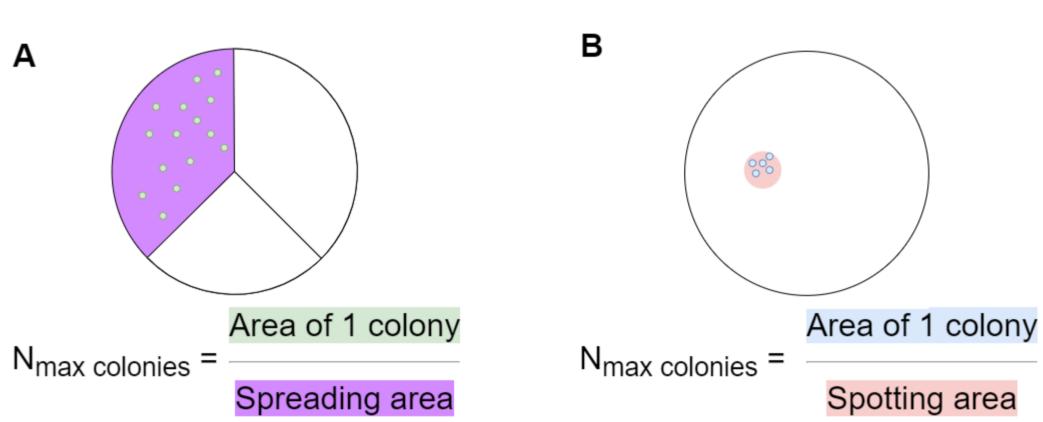


Figure 3: Illustration of how the maximal number of countable colonies was determined.

A. Conventional plating B. Spotting

Mathematical methods and software

• Optimization of the likelihood function of the « most probable number » (1) method was performed using the R function optim (3). Correlated bivariate least square regression was performed using the BivRegBLS package (4). Data wrangling and graphing was performed with the tidyverse package (5)

Results

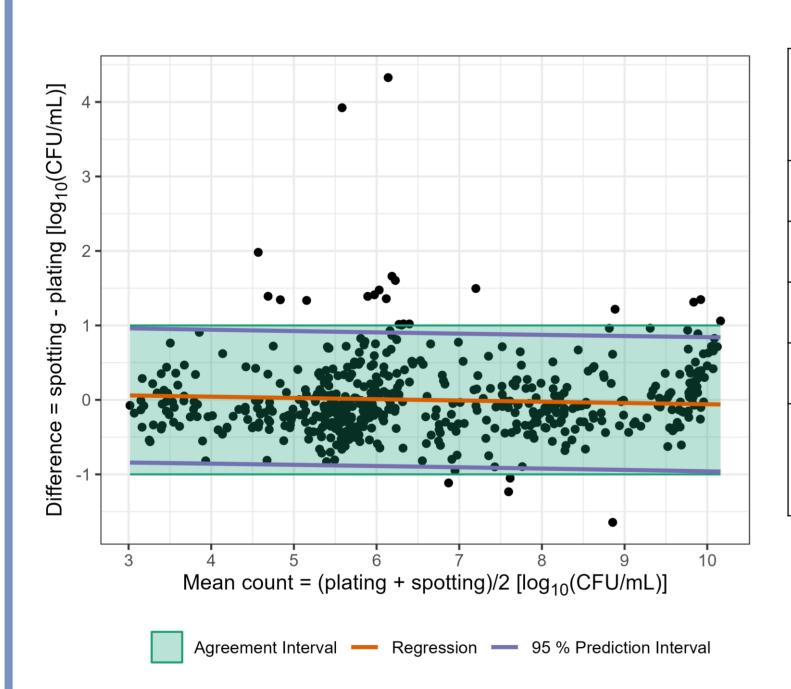


Figure 4: Bland-Altman plot (n=551).

Table 1: Performance of the two plating methods

	Conventional plating	Spotting
Number of time-kill per week	8-12	48-96
Mean maximal number of colonies per plating	675	7.84
Limit of detection (CFU/mL)	4*10 ¹	10 ³
Average bias (log ₁₀ CFU/mL)		6.6 * 10-4
Repeatability standard deviation (log ₁₀ CFU/mL)	0.105	0.445

CST, PAE32, plating

CST, PAE32, spot

CST, PAE32, spot

CST, PAE32, spot

Time (h)

CST, PAE32, spot

Above A Below

XMIC

Figure 5: Typical time-kill results. A. Conventional plating B. Spotting

- As seen in **Table 1** spotting method was accurate but exhibited lower precision than the conventional plating method.
- The accuracy was confirmed over the whole range of measurements as evidenced by the red regression line on Figure 4.
- The two measurement methods were considered in agreement when the difference was between -1 and +1 \log_{10} CFU/mL (**Figure 4** green area). In our experiment they agreed since the 95% prediction interval of the differences (**Figure 4** purple lines) lied inside the agreement interval (**Figure 4** green area).
- This agreement is qualitatively shown by **Figure 5** where typical time-kill profiles obtained by plating (**Figure 5A**) and by spotting (**Figure 5B**) are comparable.

Conclusion

The spotting method yields bacterial counting results comparable with the conventional plating method while being significantly higher throughput.

References

- 1) Martini et al. <u>10.1101/2023.05.18.541301</u> 3) R Core Team (2023) <u>www.R-project.org</u> 5) Wickham et al. <u>10.21105/joss.01686</u>
- 2) Francq and Govaerts, <u>10.1002/sim.6872</u> 4) Francq et al. <u>10.1002/sim.8709</u>



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