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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



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) 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



Table 1: Performance of the two plating methods		
	Conventional plating	Spotting
Number of time-kill per week	8-12	48-96
ean 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



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**.

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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).

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

Agreement Interval — Regression — 95 % Prediction Interva

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

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4) Francq et al. 10.1002/sim.8709 2) Francq and Govaerts, <u>10.1002/sim.6872</u>



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