Quantifying the bias of the viable cell enumeration process and its impact on microbial inactivation



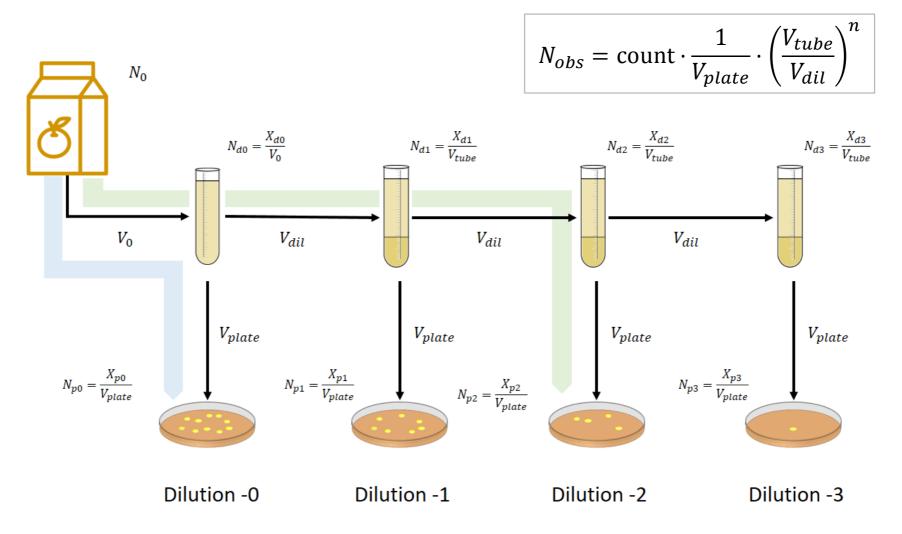
International Conference on Uncertainty in Risk Assessment

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Serial dilution/plate count







Sources of uncertainty



- (Sub)sampling error.
- Log-transformation.
- Errors in volumes (calibration errors).
- Counting errors.
- Clustering of bacterial cells.

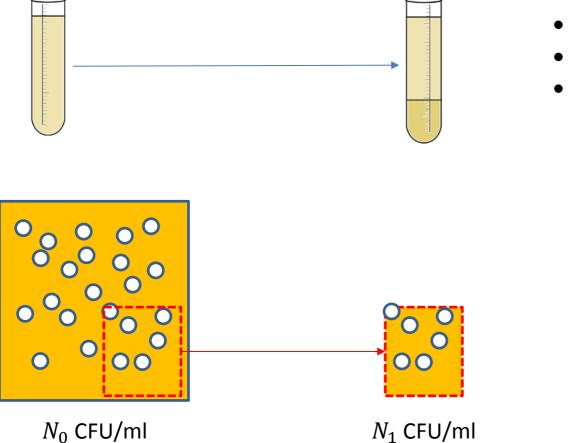




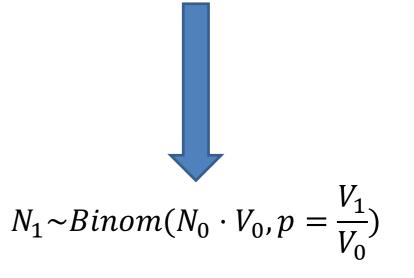
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Modelling of one (sub)sampling





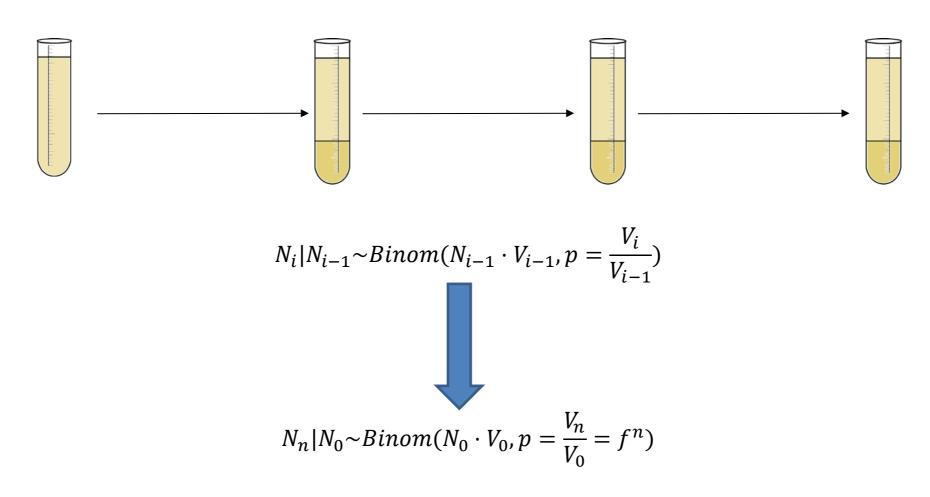
- Finite volumes.
- Finite number of particles.
- Uncorrelated positions (no clumps; perfect mixing).





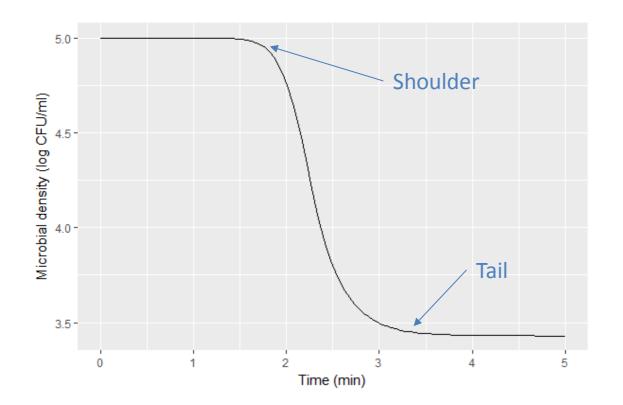
Modelling of serial dilutions







Case study: microbial inactivation



Survivor curve typically non-lineal

Shoulder: Se debe de mantener el tratamiento para inactivar

Tail: More resistant fraction/acclimation.

Very relevant for food safety

How are non-linearities affected by experimental error?





Simulation of inactivation experiments

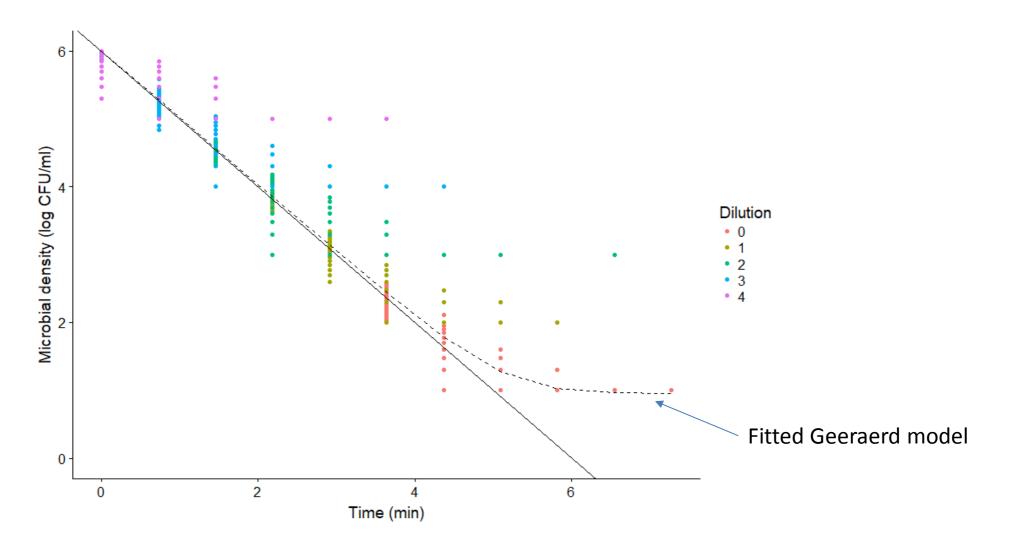


- 1. Log-linear survivor curve $\log N(t) = \log N_0 \frac{1}{D}t$
- 2. 11 sampling points. Dilutions 0 to -5.
- 3. Plate count according to Binomial distribution.
- 4. Plates with >300 colonies omitted.
- 5. 600 experiments simulated.



Uncertainty can cause bias







Experimental design affects bias



	0.1 mL en placa	0.5 mL en placa	1 mL en placa
0 dilutions	0.87 log CFU/ml	0.17 log CFU/ml	-0.17 log CFU/ml
1 dilution	1.86 log CFU/ml	1.17 log CFU/ml	0.84 log CFU/ml
2 dilutions	2.89 log CFU/ml	2.17 log CFU/ml	1.88 log CFU/ml
3 dilutions	3.94 log CFU/ml	3.22 log CFU/ml	2.92 log CFU/ml
4 dilutions	5.00 log CFU/ml	4.25 log CFU/ml	3.93 log CFU/ml



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Conclusions/outlook



- Uncertainty related to experimental design.
- Uncertainty can cause bias.
- Reporting of volumes plated/dilutions in scientific publications.
- Do not mix different dilutions/volumes plated.
- Detection limit (Chik et al. 2018, *Learning Something From Nothing: The Critical Importance of Rethinking Microbial Non-detects*. Frontiers in Microbiology).





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Tail or artefact? Illustration of the impact that uncertainty of the serial dilution and cell enumeration methods has on microbial inactivation

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Danke schön! ¡Muchas gracias!



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