ELECTRONIC SUPPLEMENTARY INFORMATION FOR THE PAPER ENTITLED:

Fecal coliform population dynamics associated with the thermophilic stabilization of treated sewage sludge

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Table S1. UniFrac P-vales based on pairwise phylogenic population difference comparisons. Values are corrected for the number of pairwise comparisons between each set of populations. P values > 0.1, indicate that no significant phylogenetic differences exist between the compared populations.

Unifrac P-values for pairwise difference comparisons

.	50°C Pre	50°C Post	55°C Pre	55°C Post	60°C Pre	60°C Post	Patho-genic	Non-path.
50°C Pre		1	1	1	1	1	1	1
50°C Post			1	0.28	0.28	1	0.28	0.28
55°C Pre				1	1	1	1	1
55°C Post					1	1	1	1
60°C Pre						0.28	1	1
60°C Post							1	0.84
Pathogenic								1
Non-path.								

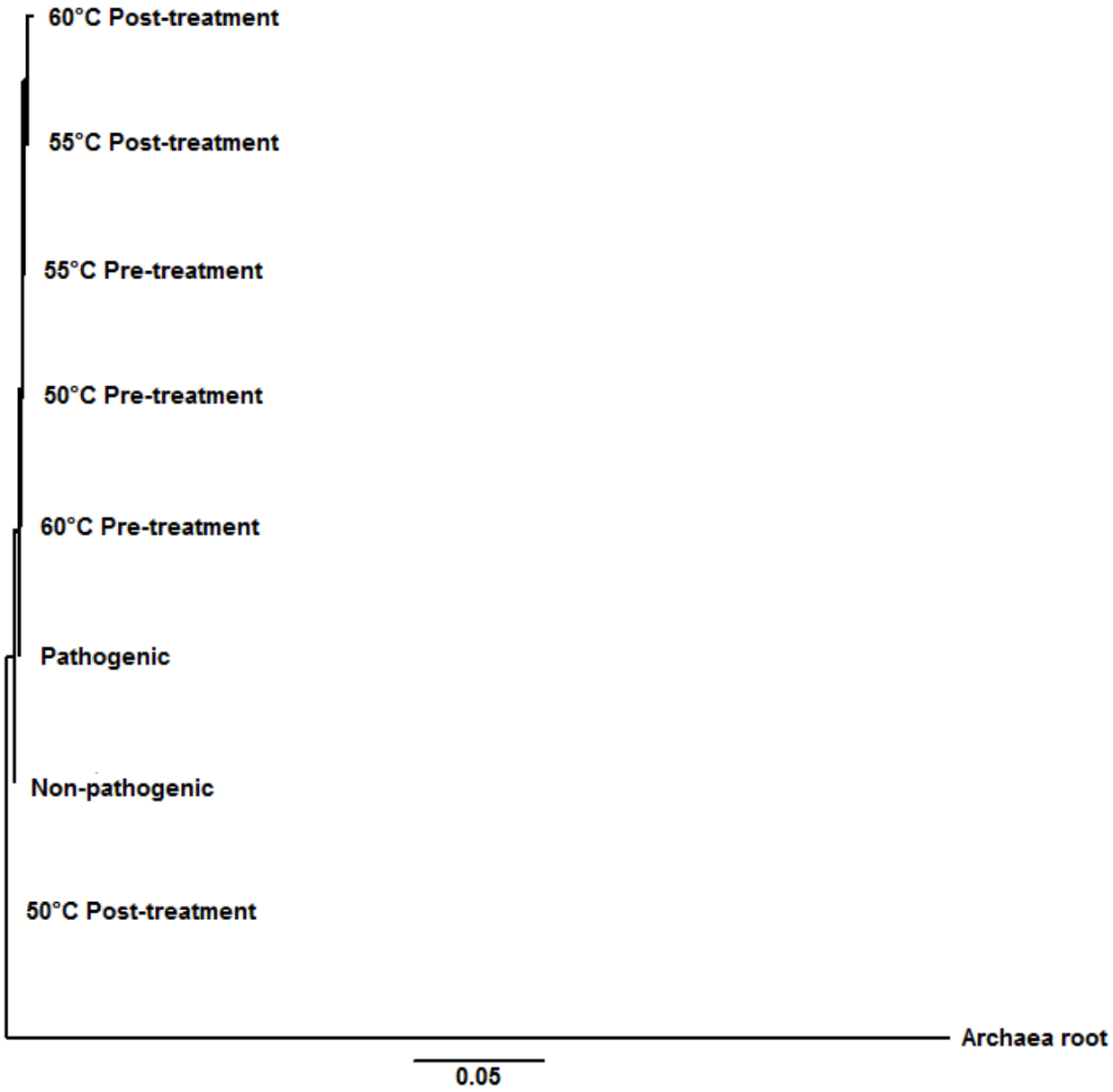

Figure S1. Phylogenic tree of consensus sequences for *E. coli* isolates sampled pre and post thermophilic batch treatment at each temperature compared to consensus sequences for pathogenic and non-pathogenic *E. coli* strains. The scale of 0.05 corresponds to a distance representing 5 deviations in 100 base pairs. The tree is rooted with the Archaebacteria *Methanobacterium congolense*.

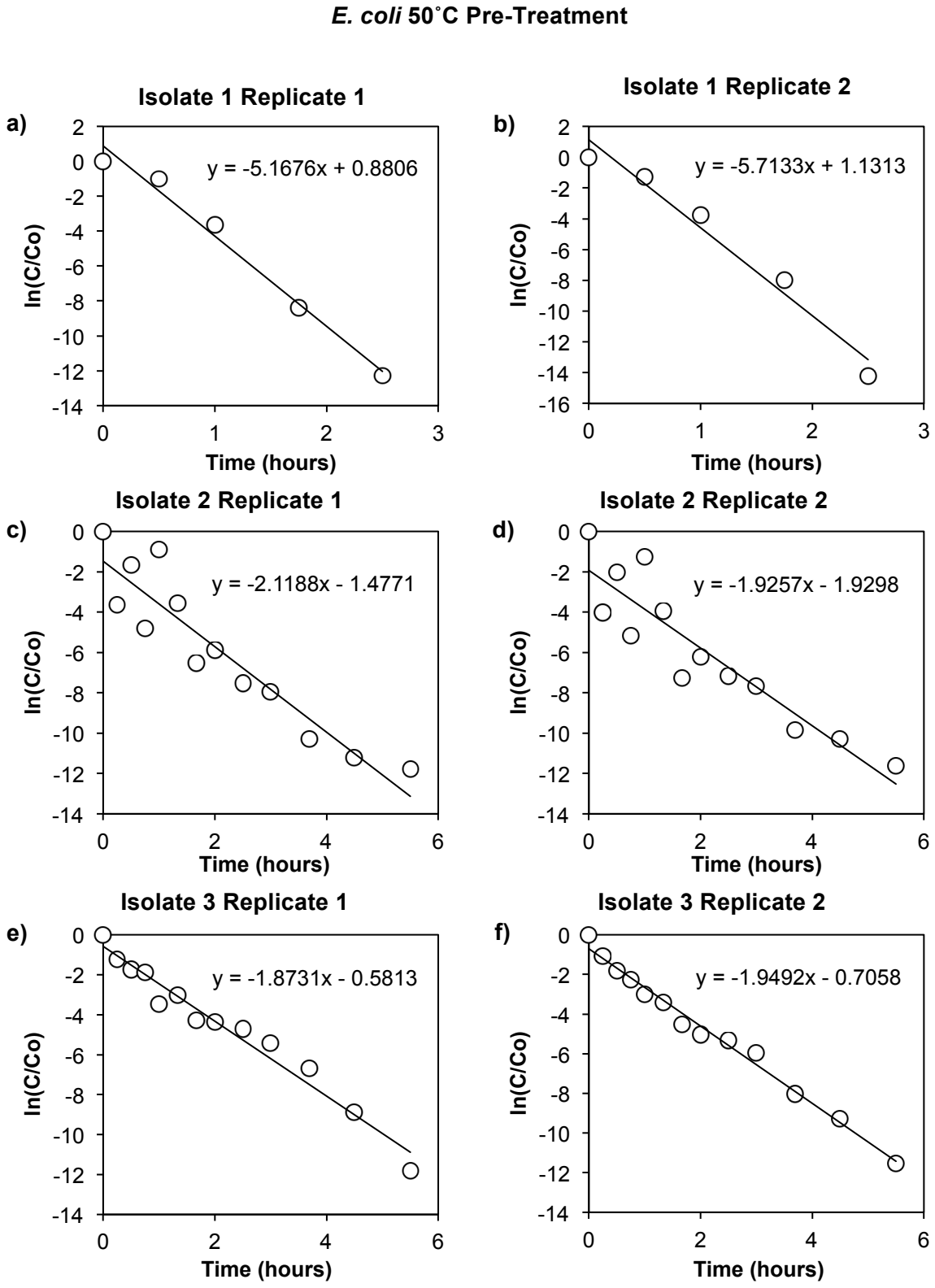

Figure S2. Inactivation profiles at 50˚C for pre-treatment *E. coli* isolates. Each isolate is tested in duplicate. Testing was conducted in PBS.

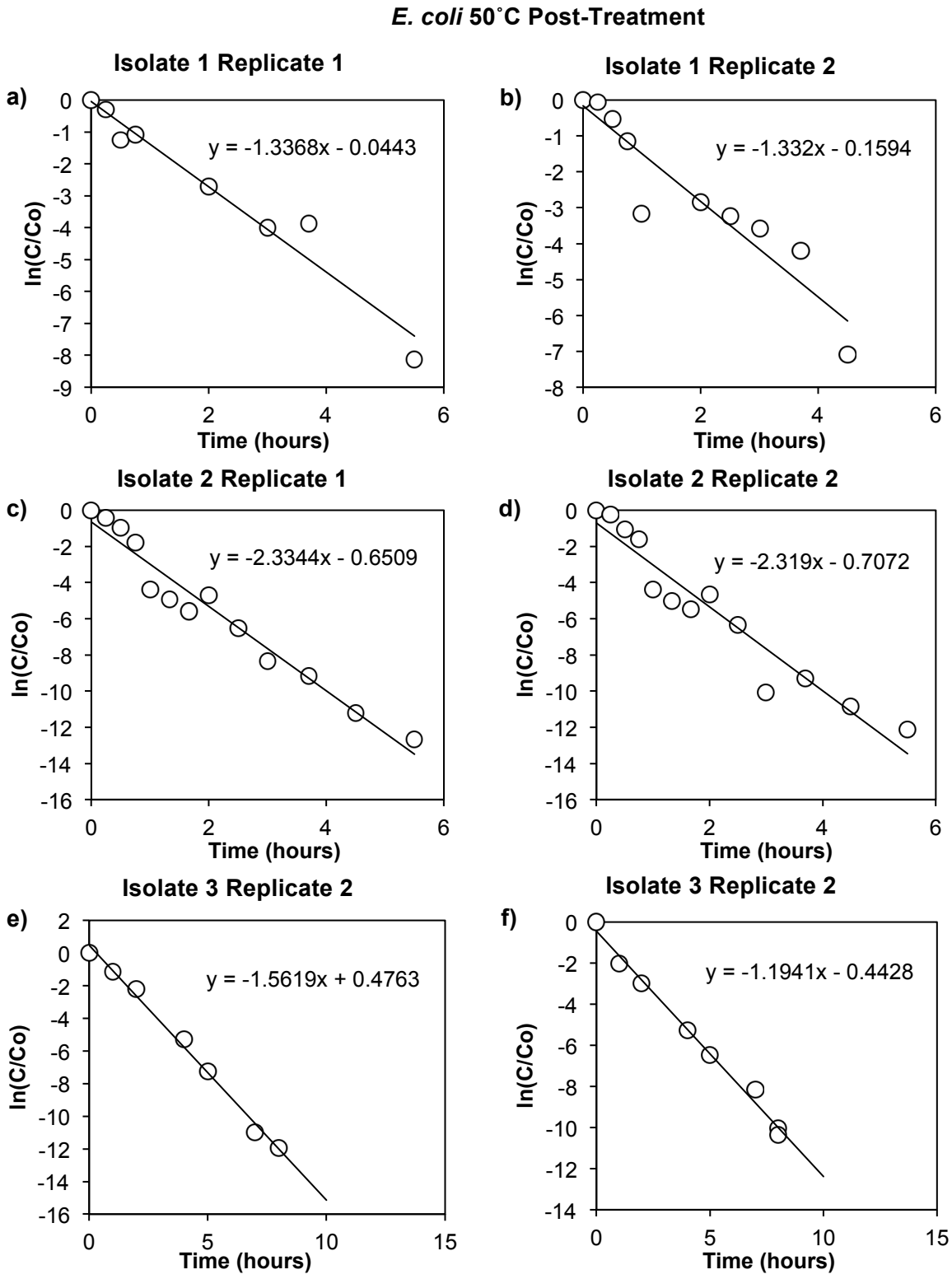

Figure S3. Inactivation profiles at 50˚C for post-treatment *E. coli* isolates. Each isolate is tested in duplicate. Testing was conducted in PBS.

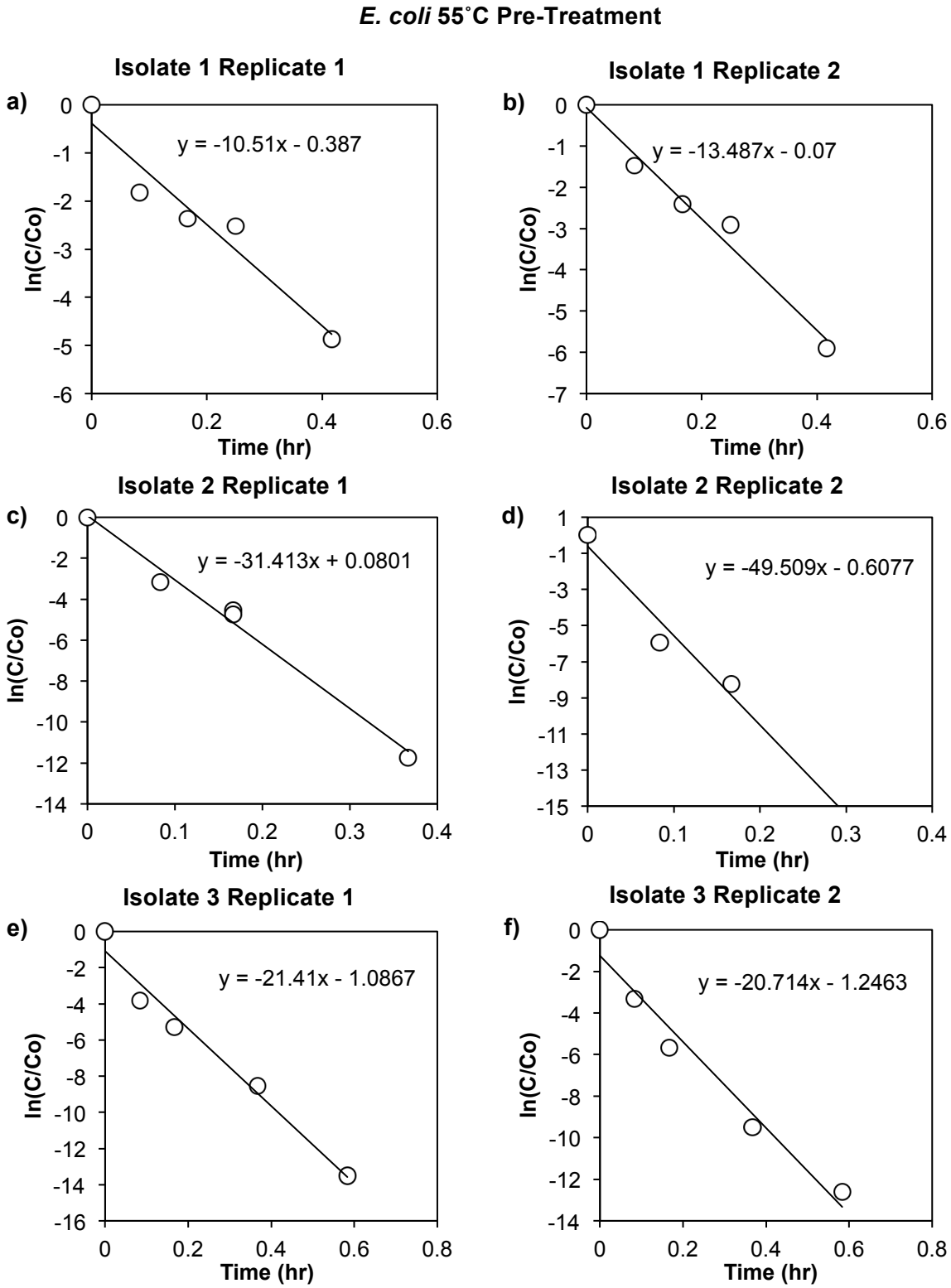

Figure S4. Inactivation profiles at 55˚C for pre-treatment *E. coli* isolates. Each isolate is tested in duplicate. Testing was conducted in PBS.

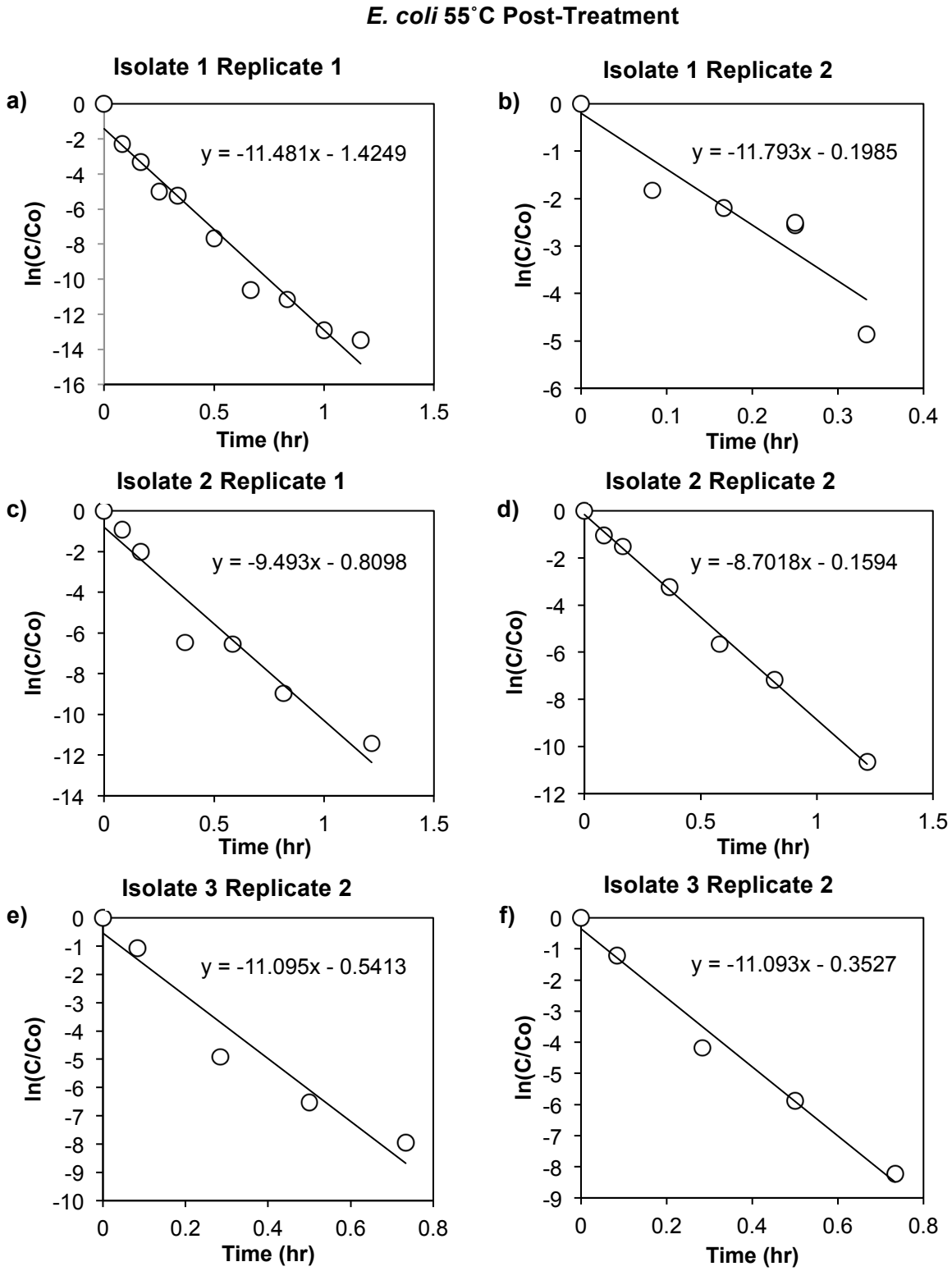

Figure S5. Inactivation profiles at 55˚C for post-treatment *E. coli* isolates. Each isolate is tested in duplicate. Testing was conducted in PBS.

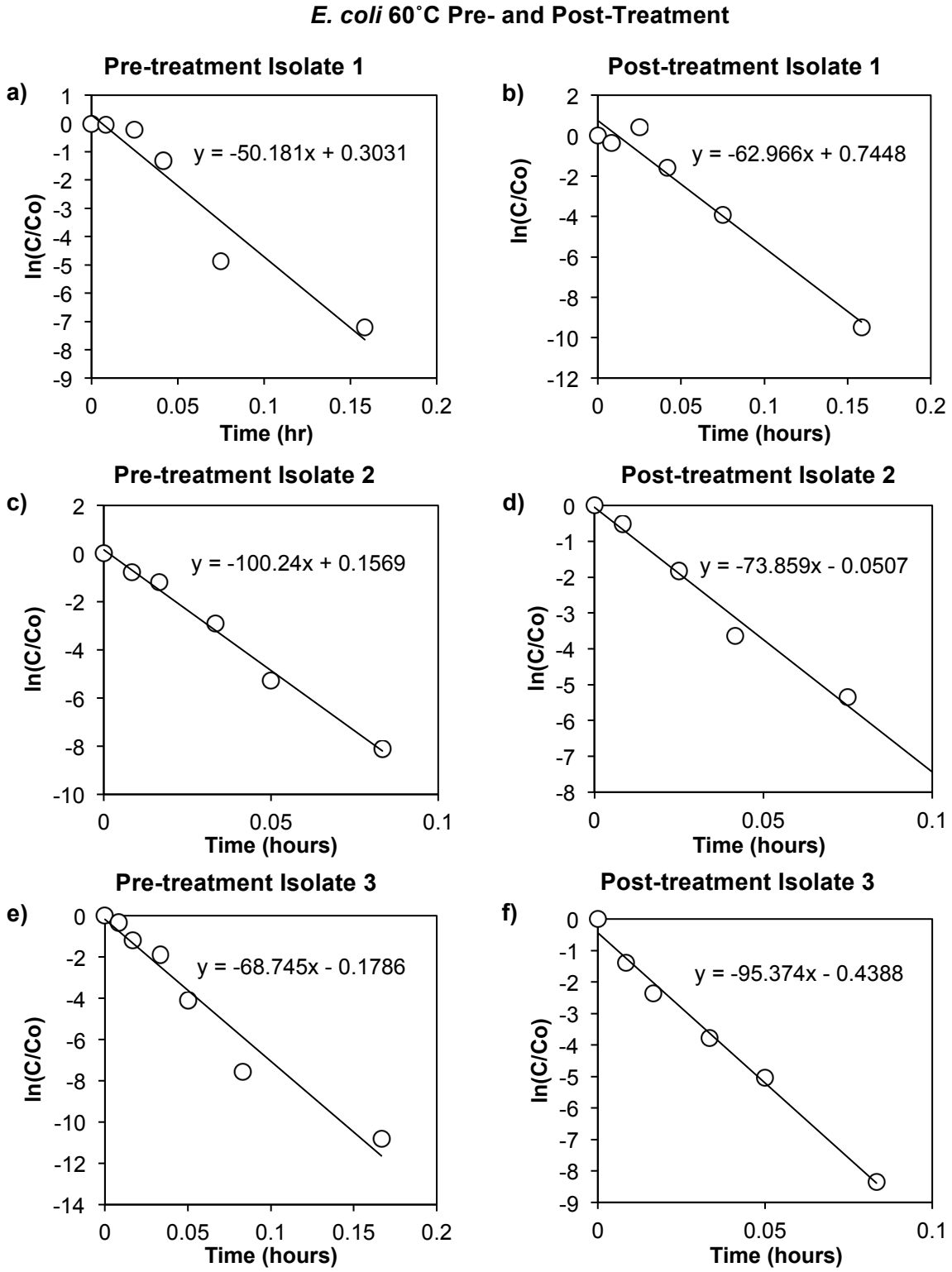

Figure S6. Inactivation profiles at 60˚C for pre-treatment and post-treatment *E. coli* isolates. Testing was conducted in PBS.

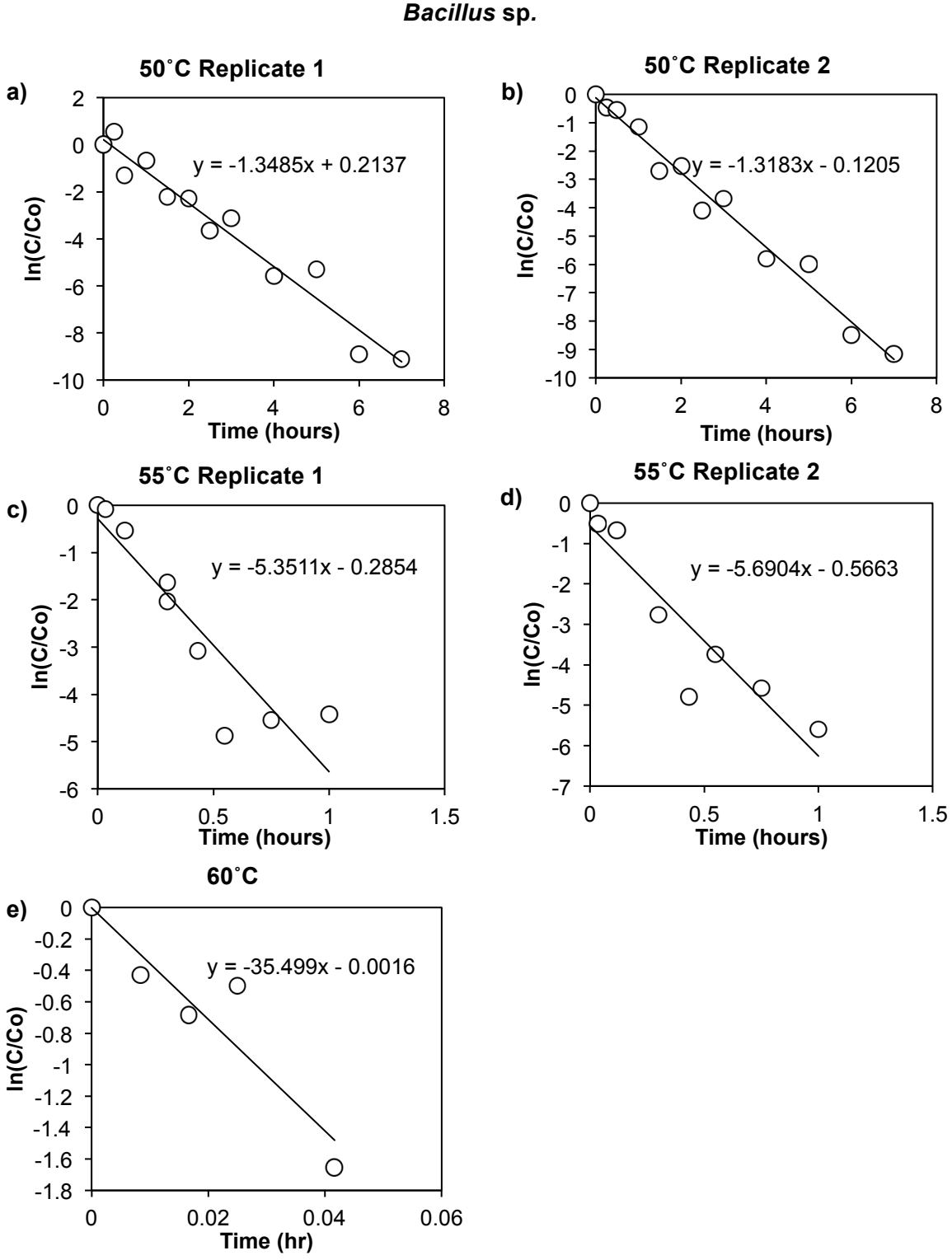

Figure S7. Inactivation profiles of *Bacillus* sp. in PBS at various temperatures. Inset equations and lines represent linear best fit.

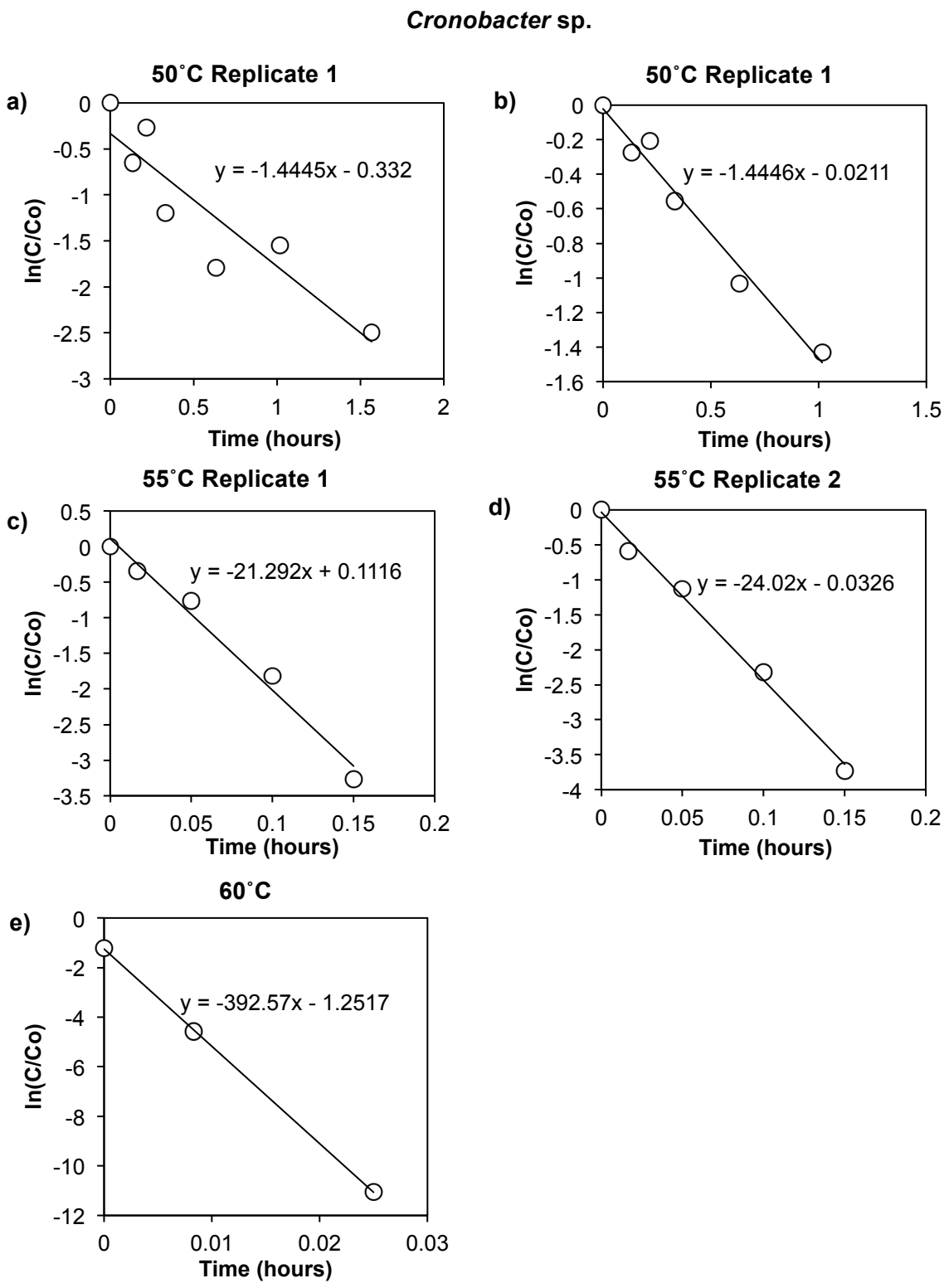

Figure S8. Inactivation profiles of *Cronobacter* sp. in PBS at various temperatures. Inset equations and lines represent linear best fit.

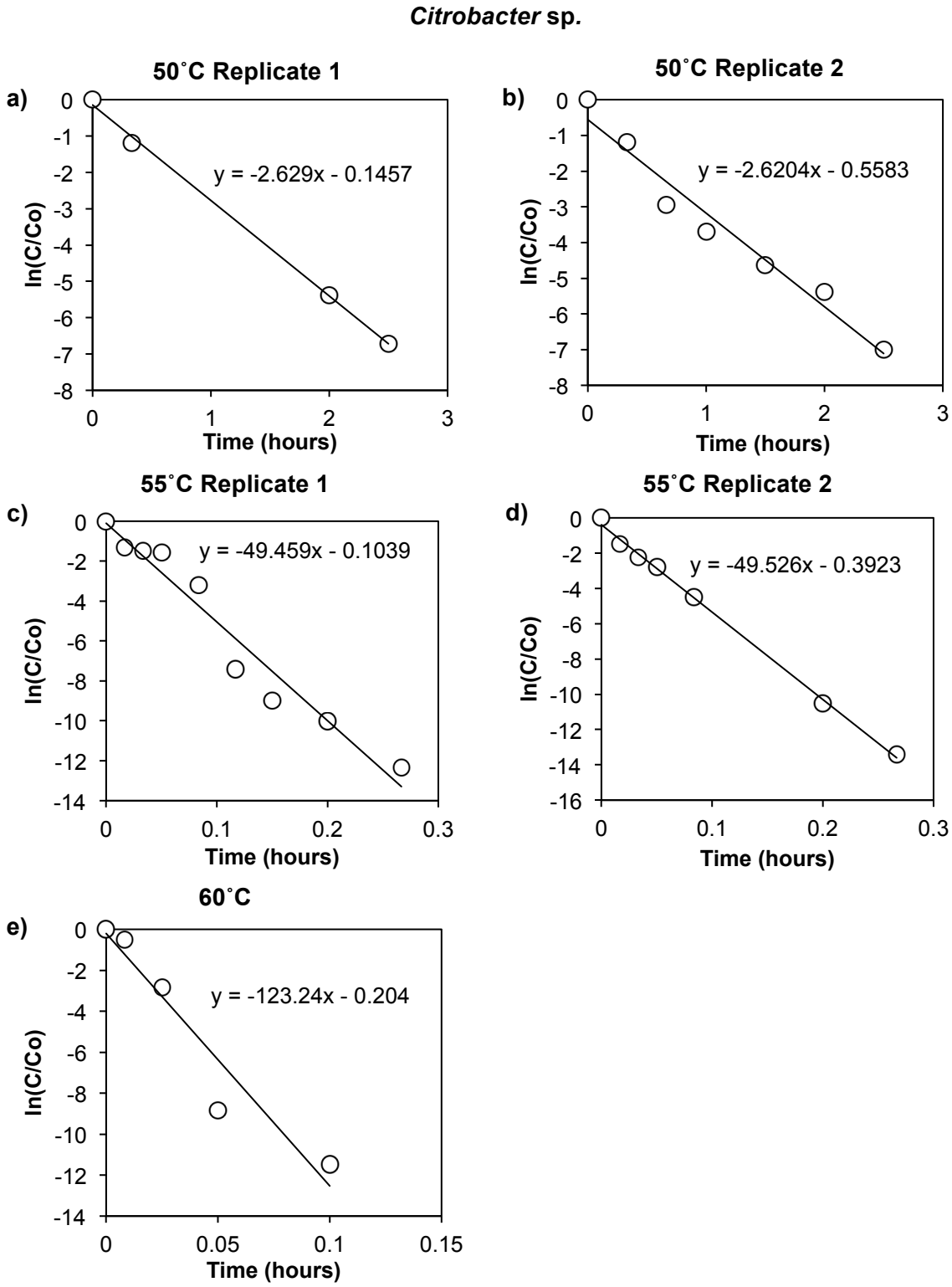

Figure S9. Inactivation profiles of *Citrobacter* sp. in PBS at various temperatures. Inset equations and lines represent linear best fit.

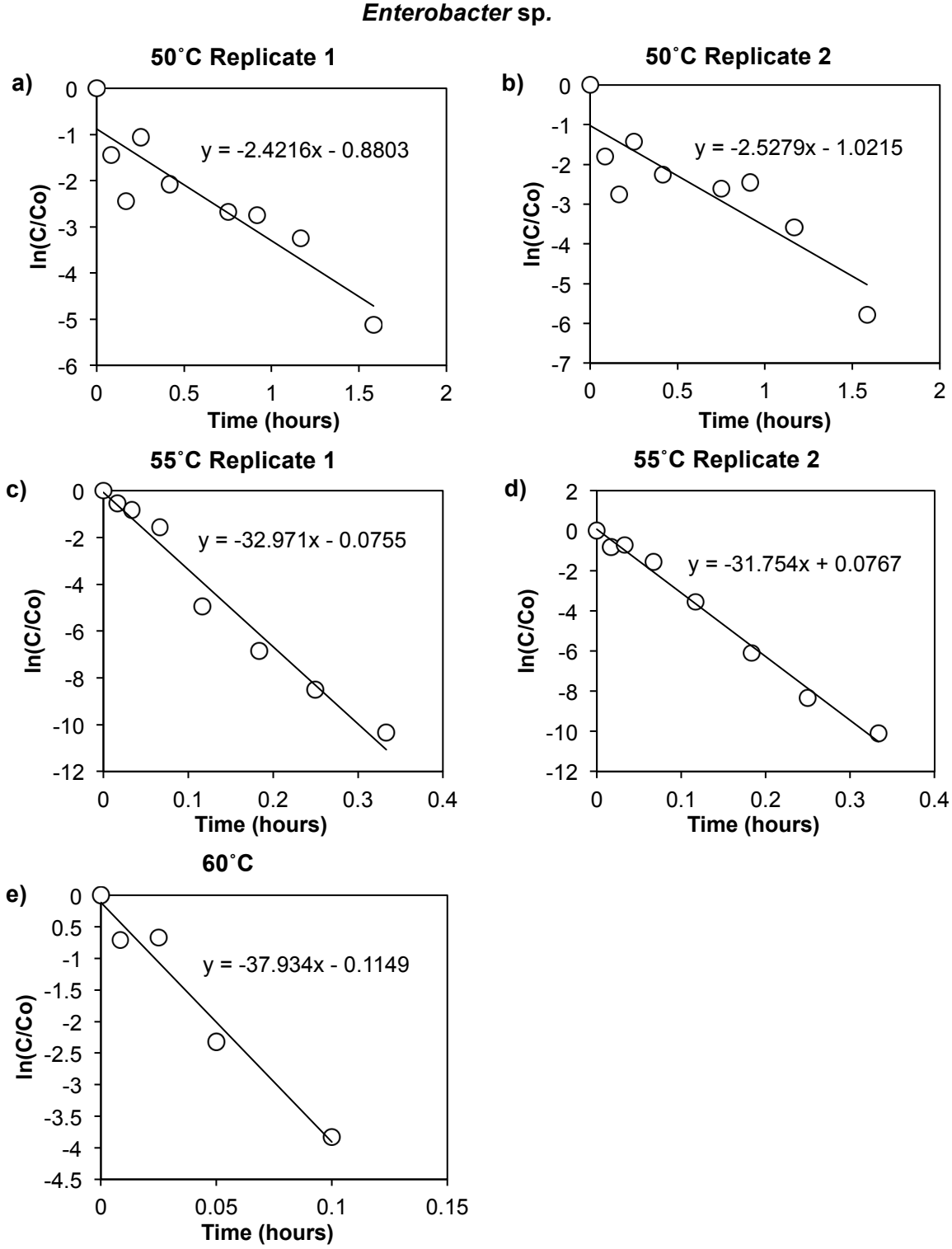

Figure S10. Inactivation profiles of *Enterobacter* sp. in PBS at various temperatures. Inset equations and lines represent linear best fit.

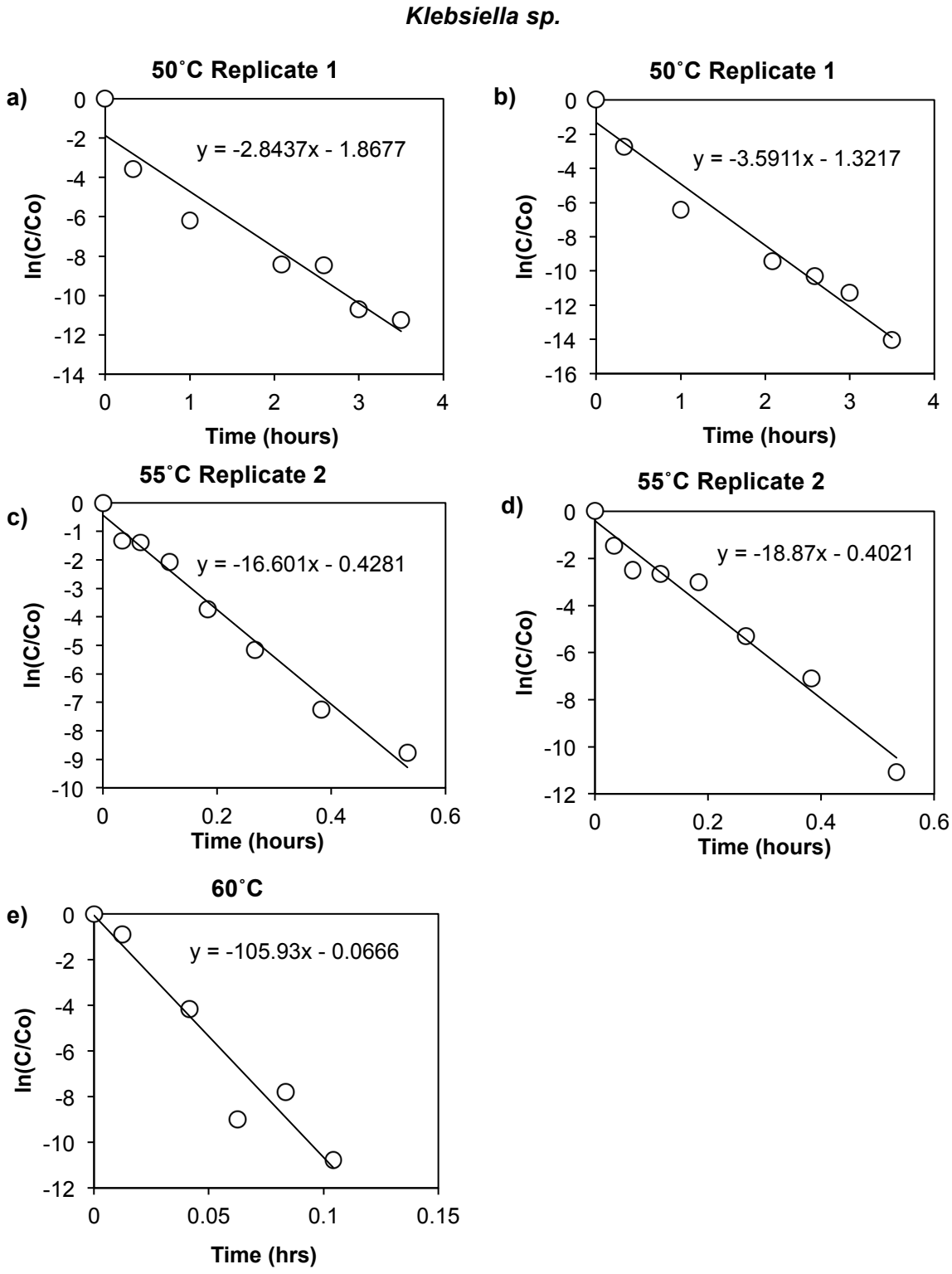

Figure S11. Inactivation profiles of *Klebsellia* sp. in PBS at various temperatures. Inset equations and lines represent linear best fit.

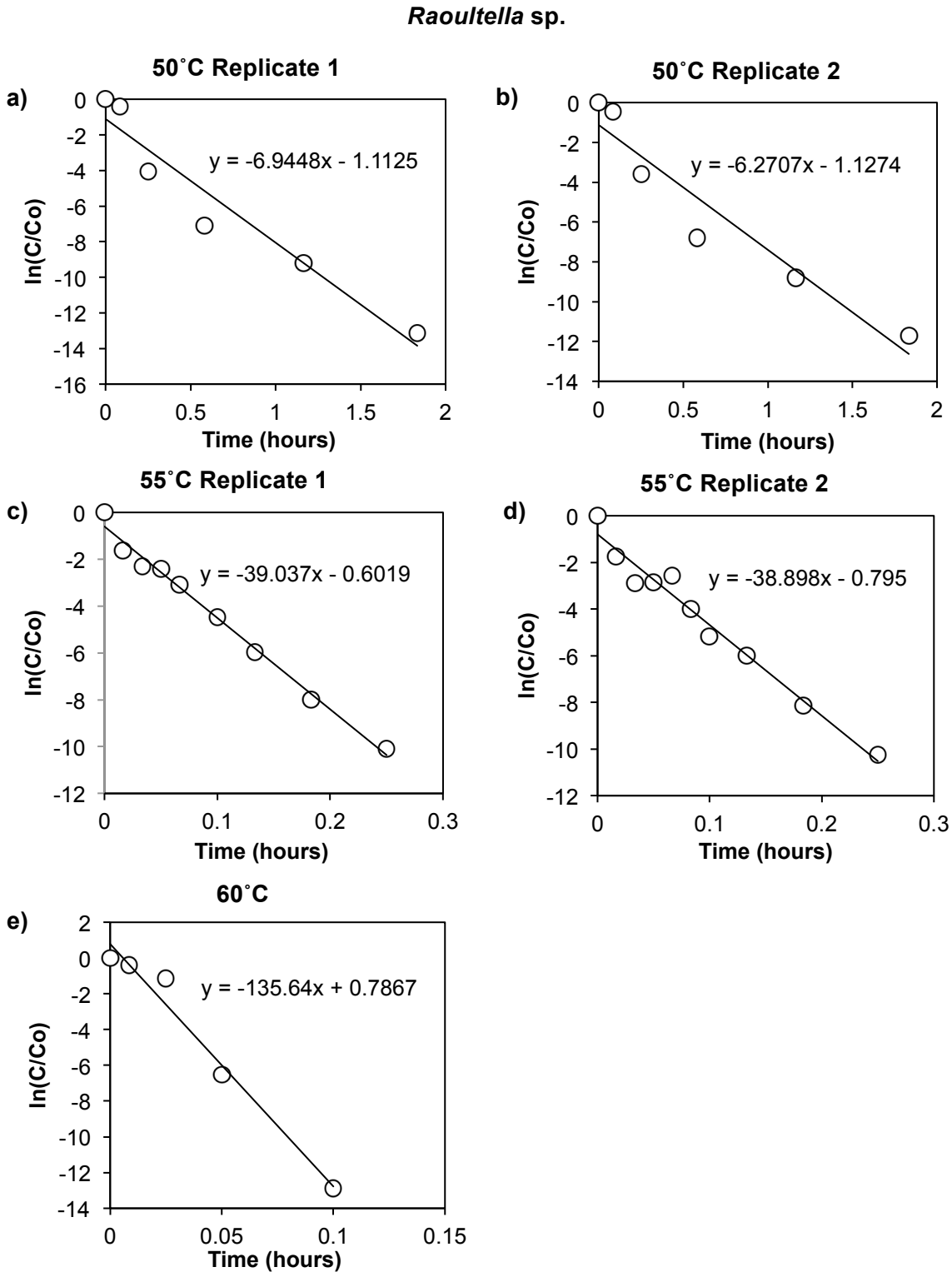

Figure S12. Inactivation profiles of *Raoultella* sp. in PBS at various temperatures. Inset equations and lines represent linear best fit.