

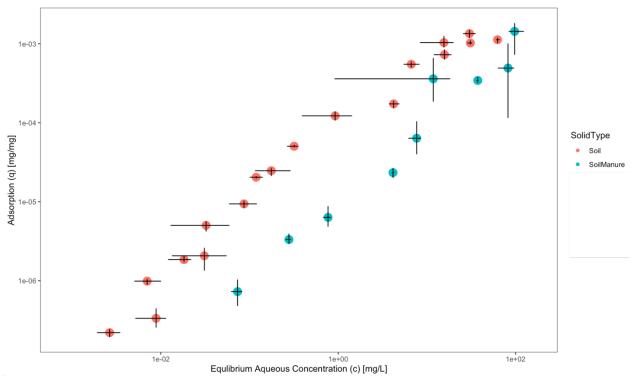
**Figure 2**: Soil-only (S) equilibration time between erythromycin and Caneseraga soil (blue) compared to control reactors (yellow) with no soil. Error bars represent the range of the 3 replicates used to obtain the points plotted.

Table 3: Converted model parameters from linearized and non-linear regressions. Eq. 2 is the non-
linear regression and Eq. 3 the linearization are Langmuir isotherms with K and $q_{max}$ parameters.

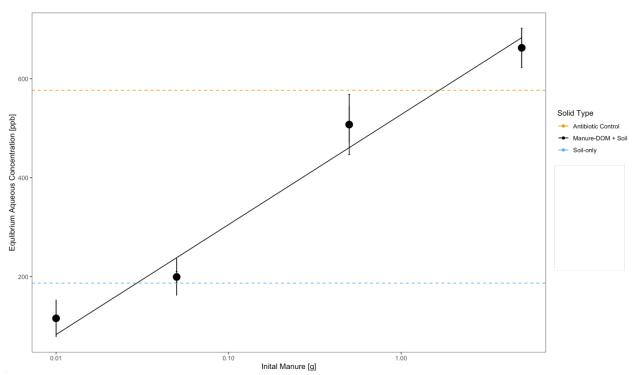
Model	S Parameters		SM Parameters		
	K	$q_{max}$	K	$q_{max}$	
Eq. 2	8.01x10 <sup>-2</sup> *	1.53x10 <sup>-3</sup> *	$\mathrm{NA}^\dagger$	$\mathrm{NA}^\dagger$	
Eq. 3	6.38x10 <sup>-2</sup> *	1.52x10 <sup>-3</sup> *	1.99x10 <sup>-4</sup> *	4.63x10 <sup>-2</sup>	

\* Statistically significant model parameters (p-value < 0.05)

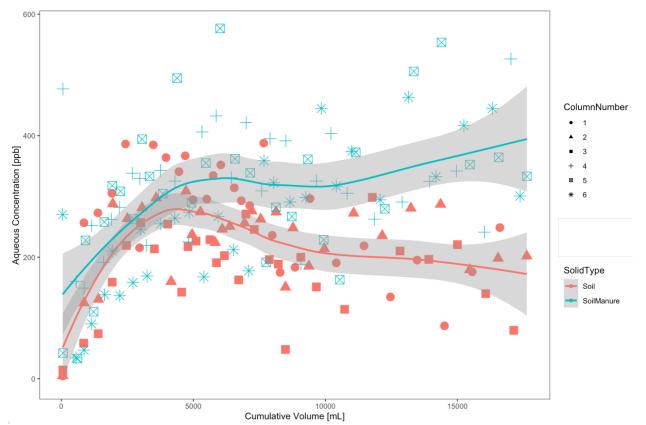
<sup>†</sup>Non-linear regression modeling (Eq. 2) with the SM data did not generate a stable result.



**Figure 3**: Soil-only (S) and soil + manure (SM) isotherms. Error bars represent the range in the 3 replicates averaged for each point.



**Figure 4**: Filtered manure-DOM influence on erythromycin aqueous concentration after 72 hr reaction with 5 g soil. Error bars represent the range of the data. Fitted model follows eq.4 (black line). Antibiotic control (yellow line) and 72 hr S sample (blue line) are provided for reference.



**Figure 5:** Aqueous concentrations of erythromycin ( $C_o = 500$  ppb) after passage through S (red) or SM (blue) columns. Unique shapes indicate samples from the same columns. Columns 1-3 were S, columns 4-6 were SM. Trend lines are fitted with a loess smoothing model.