Negative Binomial Regression Fit Report

20250610

Project Summary

- We are using data from the American Society of Microbiology's (ASM) 12 published journals to investigate the relationship between the number of citations (variable 'is.referenced.by.count') a published scientific article receives and if the authors have included access to their raw sequencing data (variable 'da', data availability) in the manuscript.
- We are trying to understand if publishing raw data helps to improve citation metrics. We have data from 2000-2024, and will also adjust for time published (variable 'age.in.months'), as older papers have had the opportunity to accumulate more citations over time.

```
opts <- options(knitr.kable.NA = " ")
knitr::kable(all_journals, digits = 3, col.names = gsub("_", " ", names(all_journals)), cap = "log(time) = log(age.in.months), shortened for the state of the
```

Table 1: $\log(\text{time}) = \log(\text{age.in.months})$, shortened for ease of reading

			time	at	at one	at	at five		five	at	at ten		ten
	full	$_{ m time}$	adj	one	year	five	years	five	years	ten	years	$_{ m ten}$	years
coefficients	full pvalue	adj	pvalue	year	pvalue	years	pvalue	years	pvalue	years	pvalue	years	pvalue
rsquared	0.678	0.681		0.355		0.516		0.660		0.223		0.680	
(Intercept)	- 0.000	-	0.000	-	0.100	2.539	0.000	-	0.000	3.995	0.000	-	0.000
	0.946	0.580		2.334				3.473				2.696	
da_Yes	- 0.000	-	0.000	1.929	0.089	0.258	0.628	-	0.763	-	0.840	-	0.448
	1.508	0.972						0.178		0.075		0.267	
$\log(\text{time})$	$0.993\ 0.000$	0.928	0.000					1.614	0.000			1.398	0.000
Applied and Environmental	- 0.000	-	0.000	2.334	0.166	0.198	0.685	-	0.828	0.046	0.866	-	0.079
Microbiology	1.034	0.885						0.121				0.565	
Genome Announcements	- 0.002	-	0.001									-	0.001
	4.530	4.495										8.881	

				1.	-	1	-			C				
		full	time	time	at	at one	at five	at five	five	five	$^{\rm at}$	at ten	4	ten
coefficients	full		adj	adj pvalue	one	year pvalue		years pvalue		years pvalue	ten	years pvalue	ten	years pvalue
			auj		year		years		years		years		years	
Infection and Immunity		0.002	-	0.001	-	0.995	0.234	0.780	-	0.168	-	0.399	-	0.024
	0.79		0.795		13.968				1.329		0.337		1.180	
Journal of Bacteriology		0.000	-	0.000	2.894	0.057	-	0.453	-	0.635	-	0.390	-	0.552
	1.17		1.077				0.460		0.336		0.344		0.250	
Journal of Clinical	0.27	30.196	0.222	0.261	3.433	0.048	-	0.782	-	0.353	-	0.001	1.976	0.000
Microbiology							0.188		0.678		0.846			
Journal of Microbiology &		0.900	-	0.863					1.022	0.701			1.542	0.385
Biology Education	0.22		0.221											
Journal of Virology		0.013	_	0.030	2.922	0.052	-	0.946	_	0.755	-	0.826	0.201	0.526
	0.43		0.355				0.033		0.173		0.061			
mBio	-	0.000	-	0.000	4.819	0.001			0.012	0.983			-	0.402
	1.88		1.114										0.266	
Microbiology Resource	-	0.0-0	-	0.055	-	0.170	-	0.000	-	0.992			-	0.347
Announcements	3.08		2.174	0.000	1.215	0.400	1.557		0.017	0.050			1.325	0.000
Microbiology Spectrum		0.000	- -	0.000	2.359	0.100			-	0.059			-	0.000
0.1	3.55		1.706	0.000	2 224	0.40=			1.002				1.791	0.040
mSphere		0.000	-	0.000	2.334	0.127			-	0.727			-	0.049
G	2.57		1.512	0.000	1 000	0.404			0.222	0.004			0.827	0.050
mSystems	-	0.000	-	0.000	1.236	0.164			-	0.964			-	0.053
1 37 4 1: 1 1	2.62		1.281	0.000		0.000	0.154	0.000	0.029	0.500	0.000	0.540	0.870	0.011
da_Yes:Applied and	0.86	9 0.002	0.744	0.003	1 500	0.290	0.154	0.800	- 0.101	0.783	0.269	0.548	0.100	0.811
Environmental					1.592				0.191					
Microbiology	1.00	0.0.050	1 100	0.407									7.010	0.011
da_Yes:Genome	1.68	8 0.259	1.168	0.407									7.219	0.011
Announcements	1.00	F 0 011	0.000	0.000	15 005	0.004		0.001	0.004	0.400			0.000	0.001
da_Yes:Infection and	1.08	5 0.011	0.890	0.023	15.067	0.994	1 000	0.301	0.964	0.430			0.860	0.221
Immunity	1 50	c o ooo	1 110	0.001		0.055	1.239	0.100	0.005	0.050			0.005	0.707
da_Yes:Journal of	1.52	6 0.000	1.118	0.001	0.007	0.655	1.074	0.180	0.885	0.358			0.205	0.737
Bacteriology	0.70	0.0.007	0.700	0.020	0.697	0.000		0.000	1 110	0.000	0.260	0.501		0.000
da_Yes:Journal of Clinical	0.79	9 0.027	0.709	0.030	- 0.111	0.209	0.045	0.966	1.110	0.232	0.360	0.521	1 694	0.002
Microbiology					2.111		0.045						1.634	
da_Yes:Journal of														
Microbiology & Biology														
Education														

				time	at	at one	at	at five		five	at	at ten		ten
		full	$_{ m time}$	adj	one	year	five	years	five	years	ten	years	ten	years
coefficients	full	pvalue	adj	pvalue	year	pvalue	years	pvalue	years	pvalue	years	pvalue	years	pvalue
da_Yes:Journal of Virology	0.80	2 0.009	0.630	0.023	_	0.226	-	0.521	_	0.907	0.155	0.784	-	0.440
					1.669		0.428		0.086				0.347	
da_Yes:mBio	1.15	$61\ 0.000$	0.901	0.001	-	0.002			-	0.728			-	0.721
					3.880				0.234				0.149	
$da_Yes:Microbiology$	0.62	$21\ 0.664$	0.131	0.909					-	0.304			-	0.666
Resource Announcements									1.738				0.618	
$da_Yes:Microbiology$	1.02	270.009	0.728	0.017	-	0.189			-	0.648			-	0.647
Spectrum					1.525				0.302				0.214	
$da_Yes:mSphere$	1.36	660.003	1.036	0.009	-	0.255			-	0.881			0.137	0.794
					1.523				0.116					
$da_Yes:mSystems$	0.88	$31\ 0.062$	0.538	0.164					-	0.375			-	0.500
									0.671				0.358	
$da_Yes:log(time)$	0.36	670.000	0.249	0.000					0.066	0.685			0.096	0.249
log(time):Applied and	0.20	060.000	0.179	0.000					-	0.919			0.109	0.145
Environmental									0.016					
Microbiology														
$\log(\text{time})$:Genome	0.57	80.057	0.572	0.047									1.508	0.012
Announcements														
log(time):Infection and	0.08	$31\ 0.098$	0.083	0.073					0.248	0.355			0.195	0.113
Immunity														
$\log(\text{time})$:Journal of	0.15	670.000	0.141	0.000					-	0.960			-	0.565
Bacteriology									0.010				0.057	
$\log(\text{time})$:Journal of	-	0.023	-	0.030					0.310	0.128			-	0.000
Clinical Microbiology	0.09	3	0.083										0.481	
$\log(\text{time})$:Journal of	-	0.251	-	0.112					-	0.390			-	0.042
Microbiology & Biology	0.54	3	0.552						0.717				0.952	
Education														
log(time):Journal of	0.06	690.046	0.054	0.098					0.071	0.641			-	0.481
Virology													0.053	
$\log(\text{time})$:mBio	0.52	270.000	0.355	0.000					0.098	0.507			0.161	0.033
$\log(\text{time})$:Microbiology	0.39	030.273	0.170	0.564					-	0.376			-	0.969
Resource Announcements									0.408				0.014	
$\log(\text{time})$:Microbiology	0.93	90.000	0.420	0.000					0.309	0.040			0.530	0.000
Spectrum														
$\log(\text{time})$:mSphere	0.62	$22\ 0.000$	0.372	0.000					0.063	0.720			0.217	0.033

coefficients	full	full pvalue	time adj	time adj pvalue	at one year	at one year pvalue	at five years	at five years pvalue	five years	five years pvalue	at ten years	at ten years pvalue	ten years	ten years pvalue
log(time):mSystems	0.71	9 0.000	0.391	0.000					0.073	0.682			0.312	0.007
da_Yes:log(time):Applied	_	0.005	-	0.010					0.109	0.569			0.009	0.929
and Environmental	0.17	0	0.142											
Microbiology														
$da_Yes:log(time):Genome$	-	0.213	-	0.355									-	0.009
Announcements	0.39		0.277										1.604	
$da_Yes:log(time):Infection$	-	0.000	-	0.020					-	0.519			-	0.248
and Immunity	0.23		0.194						0.219				0.193	
$da_Yes:log(time):Journal$		0.000	-	0.000					-	0.388			-	0.880
of Bacteriology	0.40		0.317						0.230				0.022	
da_Yes:log(time):Journal	-	00-	-	0.208					-	0.209			0.462	0.000
of Clinical Microbiology	0.11	2	0.089						0.325					
da_Yes:log(time):Journal														
of Microbiology & Biology														
Education		0.010		0.000					0.000	0.071			0.000	0.204
da_Yes:log(time):Journal		0.010	- 0.197	0.026					0.033	0.871			0.098	0.364
of Virology	0.17		0.137	0.010					0.100	0.550			0.076	0.450
$da_Yes:log(time):mBio$	0.21	0.003	0.167	0.010					0.109	0.558			0.076	0.452
da Yes:log(time):Microbiolog			0.107	0.640					0.386	0.415			0.026	0.944
Resource Announcements	0.24		0.140	0.040					0.360	0.415			0.020	0.944
da_Yes:log(time):Microbiolog			0.140	0.262					0.148	0.433			0.118	0.354
Spectrum	0.15		0.098	0.202					0.140	0.400			0.110	0.004
da_Yes:log(time):mSphere	-		-	0.018					0.050	0.818			_	0.851
	0.29		0.232	0.010					0.000	0.010			0.024	0.001
da_Yes:log(time):mSystems		0.179	-	0.347					0.213	0.315			0.109	0.418
	0.16		0.094						J.219	0.010			3.100	0.110

How well do the models fit (by Cragg-Uhler pseduo R-squared metric)

- Model format for all data from all journals MASS::glm.nb(is.referenced.by.count~ da_factor + log(age.in.months) + container.title + container.title*da_factor + log(age.in.months)*da_factor + container.title*log(age.in.months) + log(age.in.months)*da_factor*container.title, data = nsd_yes_metadata, link = log)

- Use model format for data from each journal MASS::glm.nb(is.referenced.by.count~da_factor + log(age.in.months) + log(age.in.months)*da_factor, data = <each journal>, link = log)
- Overall model fit with all data from all journals:
 - $R^2 \text{ value} = 0.678$
 - Removal of top 1% of data: R^2 value = 0.682
 - Truncate data to last 5 years: R^2 value = 0.660
 - Truncate data to last 10 years: R^2 value = 0.680
 - Summary: Model fit by R² metric does not change by removing the top 1% of data or truncating to data from the last 5 or 10 years.
- Overall model fit for data from EACH journal individually:
 - -4/12 journals have **overall model fit** with $R^2 > 0.5$
 - -4/12 journals have fit with $R^2 > 0.5$ with top 1% of data removed
 - -10/11 journals have model fits >0.5 when **truncated to the last 5 years**, so they are better than their fit overall (one journal has no data from this period)
 - -8/12 journals have model fits >0.5 when **truncated to the last 10 years**, so they are better than their fit overall
 - Summary: Data fits negative binomial model better with only more recent data considered.

All journal model is resistant to changes from removing top 1% of data, but less resistant to changes from truncating at 5 and 10 years.

- When working across the columns in the second table, we have coefficients on the left, followed by their values under the following conditions
 - full model value = all data included in the model
 - no_1percent_value = top 1% of data removed
 - five_years_value = data truncated at 5 years in age of paper
 - ten_years_value = data truncated at 10 years in age of paper
 - Note: Journal of Microbiology and Biology Education(jmbe) has N=7 papers with new sequence data and has been excluded for these analyses, but is a part of the model, and appears as NAs in the table above.

Each journal model are semi-resistant to changes from removing top 1% of data, and even less resistant to changes from truncating at 5 and 10 years.

• See above for mutations on these columns, but these models look less resistant to the transformation of removing the top 1% of data, and even less resistant to changes in coefficients from truncating at 5 and 10 years of data.