

Calculated Info

Marlin

This package contains a couple tools to create useful views of the data not by default in the dataframe. These mostly come in the form of the default case and wastewater dataframe augment function.

Wastewater augmeanting function

The wastewater function does three things. first it replaces Concentrations below the level of detection (LOD) with LOD / 2 which is considered best practice. Second it calculates a geometric average of the two gene concentrations. Finally it gets a flow and population normalized logged version of the data which previous work has shown the best results.

site	date	pop	N1	N2	flow	geoMean	sars_cov2_adj_load_log10	n
Algoma	2020-10-06	3171	10000	NA	0.498	NA	NA	34
Algoma	2020-10-13	3171	10000	16500	0.499	12845.23	0.3056463	34
Algoma	2020-10-20	3171	10000	16500	0.402	12845.23	0.2117718	34
Algoma	2020-10-27	3171	10000	16500	0.670	12845.23	0.4336205	34
Algoma	2020-11-03	3171	10000	16500	0.489	12845.23	0.2968546	34
Algoma	2020-11-10	3171	10000	16500	0.463	12845.23	0.2731267	34

Case augmeanting function

The case function does two things. first it normalizes the data by population. Second it calculates a rolling sum and average of the data. This needs the population data contained in its own data frame.

site	date	tests	prob_case	casef_case	prob_death	death	population	FirstConfirmed	per100k	casespastday	per100k	casesperday	Per100k
Algoma	2020-01-22	0	0	0	0	0	3171	0	NA	NA	NA	NA	NA
Algoma	2020-01-23	0	0	0	0	0	3171	0	NA	NA	NA	NA	NA
Algoma	2020-01-24	0	0	0	0	0	3171	0	NA	NA	NA	NA	NA
Algoma	2020-01-25	0	0	0	0	0	3171	0	NA	NA	NA	NA	NA
Algoma	2020-01-26	0	0	0	0	0	3171	0	NA	NA	NA	NA	NA

site	date	tests	prob_case	casef_case	prob_death	deathf_death	population	FirstConfirmed	per100K	casespastday	per100K	casesperday	Per100K
Algonquin	2020-01-27	0	0	0	0	0	3171	0	NA	NA	NA	NA	NA

These functions are slightly out of date and not used in many new analysis projects