ReadMe Processed Data

General guide to data:

Column Headings: RR1,RR4,RR5 etc... are each gel trap samples. 30 total. Row Indices: names of organisms found as solo sinking cells in gel trap samples

Metadata:

Fields - number viewed either 1 for entire gel, or 15 or 20 for 2mmx2mm subfields of gel **m2per field**-size of Field unit: m²

Elapsed_Time_(days) number of days gel trap was out collecting sinking particles. unit: days
True Depth (m) record of actual depth trap drifted at. unit: m

Trap_Type: type of sediment trap platform: either neutrally bouyant(NBST) or surface tethered(STT)\ **Epoch:** Unit of ~8day deployment cycle. Consecutive in time, Epoch 1 then Epoch 2 then Epoch 3

.csv files

/EXPORTS_Gel_Fluxes_115x_25x_7x.csv

-cell fluxes from all magnifications Source: A. Bodel

/EXPORTS Gel Fluxes Uncertainties 115x 25x 7x .csv

-cell flux uncertainties(based on counting uncertainty) Source: A. Bodel

/EXPORTS_Carbon_per_Cell_.csv

Estimation of amount of POC per cell for each organism found in gels

V_um3 Volume in cubic micrometers: Average Simulated Biovolume of each organism (See /Extended methods)

pgC picograms of carbon: Estimated value (See /Extended methods)

ugC micrograms of carbon:

Source: A. Bodel

/functional group fluxes.csv

-fluxes of cells grouped together by major taxonomic functional groups, Source: A. Bodel

/From colleen All EXPORTS flux summmed by Trap csv.csv

-fluxes of different particle types. Source: Colleen Durkin

.png files

/cells in gels.png Unpublished © 2020 A. Bodel

/mechanisms.png Unpublished © 2020 A. Bodel

/EXPORTS pathways.png © 2017 DK Steinberg