Archived Soil Incubations Project

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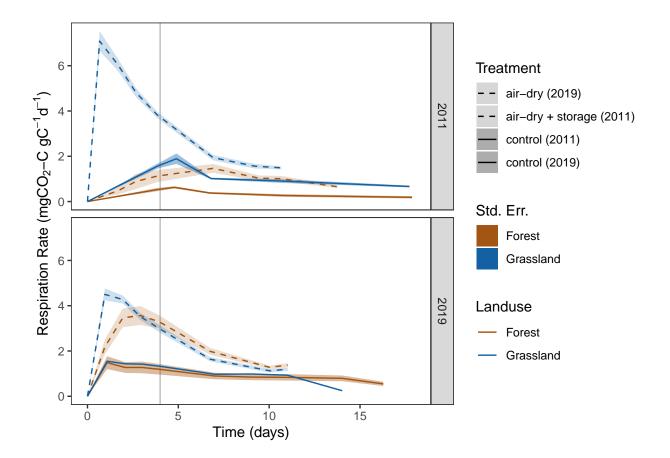
Notes:

- This workbook is intended to load and prepare the key data for analysis for the archive incubation project.
- in general, this is an updated version of script "./src/arc_inc_master.R"
- all code chunk options are set to "echo = FALSE"; see raw .Rmd file for data wrangling code.

CO₂ fluxes and soil data

- 1. Load flux data from air-dry + storage control samples, and convert from "wide" to "long" format so as to match other data.
- 2. Load flux data from air-dry + storage samples and from air-dry experiment (ctl & treatment), C & N data for all the Exploratories samples (measured in 2011), and soil mass and moisture data for all experiments.
- 3. Combine and summarize data in long format to calculate respiration rates and plot over time.
- 4. Plot of CO_2 fluxes over time. Note that the final measurement points for a few samples which took >18 days to reach CO_2 targets are excluded for display reasons. Respiration rates for those samples remained flat.

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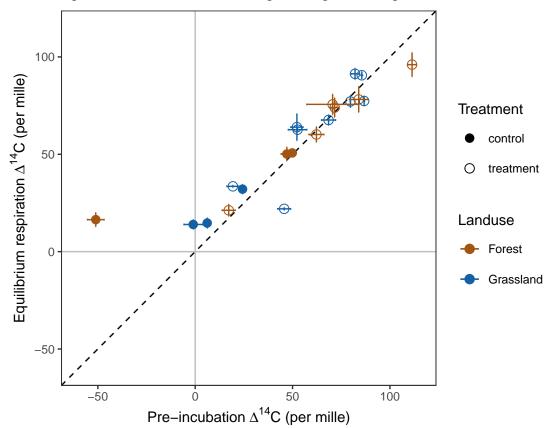
Isotope data

- 1. Read in isotope data from various sources. First load helper function 'read_jena_ams_results.R"
- 2. Next read in data from the appropriate directories in 'data/raw'.
- 3. Create a "tidy" style template for the data, i.e. variables in columns.
- Key variables are as follows:
 - SampleName (incorporates lab rep and treatment, e.g. "HEG10-1_dry")
 - ID (plot IDs, e.g. for "HEG10" for Exploratory samples)
 - Treatment (3 treatments: air-dry, air-dry + storage, storage duration; + controls)
 - Type (2 levels: F = forest, G = grassland)
 - Period (incubation period, 2 levels: pre = preincubation, inc = equilibrium incubation)
 - Experiment (3 levels: arc = air-dry + storage, rewet = air-dry/rewet, time = storage duration)
- Observational columns include:
 - $d14c (\Delta^{14}C-CO_2)$
 - $d13c (\delta^{13}C-CO_2)$
 - C_g_kg (C content)
 - dw_g (dry weight)
 - mgCO2.C_gS (mg CO₂-C respired g⁻¹ soil Period⁻¹)
 - time d (days in incubation period prior to measurement)
 - h2o_grav (gravimetric water content, percent)
 - h2o who field (percent of water holding capacity, field-moist)
- 4. Summarize observational data from timeseries data by unique IDs (SampleName).

- 5. Create helper functions for decay correction, converting Δ^{14} C to fraction modern, and cleaning up extraneous values in raw 14 C data. Archived sample Δ^{14} C data should to be corrected for decay since the year of collection. (Although the correction is very small and likely insignificant, I will do it anyway).
- decay correction formula is:

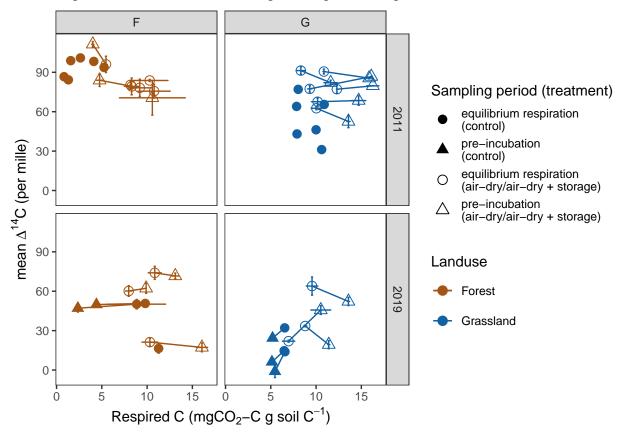
$$1000 \cdot \left((FM \cdot e^{\frac{-year_{sampled} + 1950}{8267}}) - 1 \right)$$

- 6. Clean up ¹⁴C data.
- 7. Combine data.
- 8. Count number of ¹⁴C observations for checking plots.
- 9. Plot pre-incubation period Δ^{14} C against equilibrium respiration period Δ^{14} C.
- Relative outlier point is the very negative (mean = -51.1%) HEW22 pre-incubation control samples from the 2019 air-dry experiment.
- Data from both experiments plotted together, i.e. "treatment" = air-dry/air-dry + storage
- Pre-incubation $\Delta^{14}\mathrm{C}$ was not measured for the 2011 control samples.
- Samples from three of the forest plots of the 2011 treatment samples failed to accumulate enough $\rm CO_2$ to measure $^{14}\rm C$.
- ## Warning: Removed 15 rows containing missing values (geom_point).
- ## Warning: Removed 15 rows containing missing values (geom_errorbar).
- ## Warning: Removed 15 rows containing missing values (geom_errorbarh).

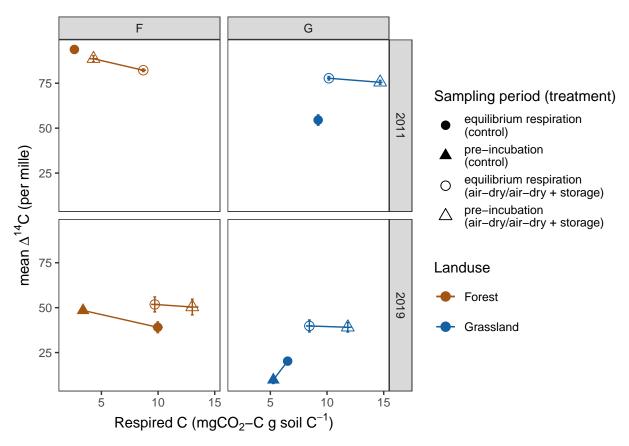


10. Plot Δ^{14} C against proportion of soil C respired by experiment, land cover, and sampling period.

- Note that data are first shown averaged by plot (Fig. 1), and then averaged by land use and treatment within sampling periods (Fig. 2)
- Pre-incubation $\Delta^{14}\mathrm{C}$ was not measured for the 2011 control samples.
- Outlier data excluded (HEW22 control pre-incubation)
- ## Warning: Removed 14 rows containing missing values (geom_point).
- ## Warning: Removed 9 rows containing missing values (geom_path).
- ## Warning: Removed 14 rows containing missing values (geom_errorbar).
- ## Warning: Removed 14 rows containing missing values (geom_errorbarh).



- ## Warning: Removed 2 rows containing missing values (geom_point).
- ## Warning: Removed 2 rows containing missing values (geom_errorbar).
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- 11. Same plot as in step 10, but including outlier data.
- ## Warning: Removed 2 rows containing missing values (geom_point).
- ## Warning: Removed 2 rows containing missing values (geom_errorbar).
- $\hbox{\tt \#\# Warning: Removed 2 rows containing missing values (geom_errorbarh).}$

