SARE water holding capacity calcs

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Data processing

NOTE: Calcs are loosely based on Agron479 lab, although I found that lab hard to follow.

I have my plot-treatment key (called key):

```
head(key)
```

```
## # A tibble: 6 x 8
##
    site_name site_desc
                                                              rep plot code
                                     sys_trt crop_trt cc_trt
##
    <chr>
             <chr>
                                             <chr>
                                                     <chr> <dbl> <dbl> <chr>
## 1 Boyd42 Boyd Farm, soybean plots sil
                                                                     2 B42-p2
                                             soy
                                                     no
                                                               1
                                                                    11 B42-p11
## 2 Boyd42 Boyd Farm, soybean plots sil
                                             soy
                                                     no
                                                               3
## 3 Boyd42
             Boyd Farm, soybean plots sil
                                                                    16 B42-p16
                                             soy
                                                     no
## 4 Boyd42
             Boyd Farm, soybean plots sil
                                             soy
                                                     no
                                                                4
                                                                    20 B42-p20
## 5 Boyd42
             Boyd Farm, soybean plots sil
                                                               5
                                                                    28 B42-p28
                                             soy
                                                     no
## 6 Boyd42
             Boyd Farm, soybean plots sil
                                                                     7 B42-p7
                                             soy
                                                     CC
```

I have my raw data (called *datraw*):

```
head(datraw)
```

```
## # A tibble: 6 x 23
     code satsamp_g satwater_g cell_nu cylinder_g
                                                    atm `10_cm` `25_cm` `50_cm`
##
                         <dbl> <chr>
##
     <chr>
             <dbl>
                                            <dbl> <dbl>
                                                          <dbl>
                                                                   <dbl>
                                                                           <dbl>
## 1 St-1~
              859.
                          4.39 1a
                                             46.1 48.9
                                                           48.5
                                                                   53.8
                                                                           49.2
## 2 St-3~
              880.
                          4.68 2a
                                             46.1 50.7
                                                           51.6
                                                                   57.4
                                                                           49.3
                                                                           49.6
## 3 St-5~
              848.
                          3.26 3a
                                             46.1 50.2
                                                           50.6
                                                                   56.9
## 4 St-7~
               842.
                          6.49 4a
                                             46.1 50.2
                                                           52.0
                                                                   59.0
                                                                           50.7
                          3.92 5a
                                             46.2 50.4
                                                                           48.9
## 5 St-2~
               856.
                                                           51.8
                                                                   55.5
## 6 St-4~
               846.
                         19.3 6a
                                             46.2 51.9
                                                           55.0
                                                                   63.1
                                                                           51.8
## # ... with 14 more variables: `100 cm` <dbl>, `200 cm` <dbl>, `500 cm` <dbl>,
      sampafter500_g <dbl>, drysoil_g <dbl>, ringpluscrap_g <dbl>, ring_g <dbl>,
      site_name <chr>, site_desc <chr>, sys_trt <chr>, crop_trt <chr>,
      cc trt <chr>, rep <dbl>, plot <dbl>
## #
```

1. Calculate the bulk density

```
dat1 <- datraw %>%
    # assume volume of soil sample is 347.50 cm3
    mutate(soilvol_cm3 = 347.5) %>%
    # calc bulk density based on dry weight of soil
    mutate(bulkden_gcm3 = drysoil_g / soilvol_cm3) %>%
    select(code, drysoil_g, bulkden_gcm3)
head(dat1)
## # A tibble: 6 x 3
```

```
## # A tibble: 6 x 3
## code drysoil_g bulkden_gcm3
```

```
<chr>>
                 <dbl>
                               <dbl>
## 1 St-1no
                                1.51
                   526
## 2 St-3no
                   532
                                1.53
## 3 St-5no
                   512
                                1.47
## 4 St-7no
                   498
                                1.43
## 5 St-2cc
                                1.50
                   520
## 6 St-4cc
                                1.39
                   482
```

2. Calculate actual amount of water released at each pressure.

Note that pressure of 999 refers to the amount of water remaining in the soil after the 500 cm pressure was applied. This was determined by weighing the soil, drying it, then reweighing it.

```
datraw %>%
mutate(
    w_0cm_g = satwater_g,
    w_2.5cm_g = atm - cylinder_g,
    w_10cm_g = `10_cm` - cylinder_g,
    w_25cm_g = `25_cm` - cylinder_g,
    w_50cm_g = `50_cm` - cylinder_g,
    w_100cm_g = `100_cm` - cylinder_g,
    w_100cm_g = `100_cm` - cylinder_g,
    w_200cm_g = `200_cm` - cylinder_g,
    w_990cm_g = `s00_cm` - cylinder_g,
    w_999cm_g = sampafter500_g - ringpluscrap_g - drysoil_g
) %>%
    select(code, starts_with("w_"))
```

```
## # A tibble: 6 x 10
##
     code w_0cm_g w_2.5cm_g w_10cm_g w_25cm_g w_50cm_g w_100cm_g w_200cm_g
##
             <dbl>
                        <dbl>
                                  <dbl>
                                           <dbl>
                                                     <dbl>
                                                                <dbl>
                                                                           <dbl>
     <chr>>
## 1 St-1~
              4.39
                         2.73
                                   2.35
                                            7.63
                                                                 4.08
                                                                           4.68
                                                      3.08
## 2 St-3~
              4.68
                         4.59
                                   5.46
                                           11.2
                                                      3.15
                                                                 3.07
                                                                           3.92
## 3 St-5~
              3.26
                         4.12
                                   4.53
                                           10.8
                                                      3.5
                                                                 3.26
                                                                           3.92
## 4 St-7~
              6.49
                         4.1
                                   5.89
                                           12.8
                                                      4.57
                                                                 4.62
                                                                           5.45
## 5 St-2~
               3.92
                         4.2
                                   5.67
                                            9.34
                                                      2.76
                                                                 3.73
                                                                           4.46
## 6 St-4~
                         5.70
                                   8.79
             19.3
                                           16.9
                                                      5.55
                                                                 3.63
                                                                           3.89
## # ... with 2 more variables: w_500cm_g <dbl>, w_999cm_g <dbl>
```

3. Gather into long form to get cumulatives.

```
dat3 <-
  dat2 %>%
  gather(w_0cm_g:w_999cm_g, key = press_cm, value = water_g) %>%
  separate(press_cm, into = c("water", "press_cm", "grams"), sep = "_") %>%
  select(-water, -grams) %>%
  # get pressure as a numeric value
  mutate(press_cm = parse_number(press_cm)) %>%
  # arrange within a sample
  group_by(code) %>%
  arrange(code, -press_cm) %>%
```

```
# get cumulative water retained in soil at that pressure point (is this right?)
  mutate(cumwater_g = cumsum(water_g))
head(dat3)
## # A tibble: 6 x 4
## # Groups: code [1]
     code
           press_cm water_g cumwater_g
##
     <chr>
              <dbl>
                        <dbl>
                                  <dbl>
## 1 B42-p10
                 999 110.
                                    110.
## 2 B42-p10
                 500
                         4.56
                                    115.
## 3 B42-p10
                 200
                                    128.
                     12.6
## 4 B42-p10
                 100
                      5.93
                                    134.
## 5 B42-p10
                  50
                         8.03
                                    142.
## 6 B42-p10
                  25
                         9.82
                                    151.
```

4. Calculate water content of soil.

Divide the cumulative water released at each pressure by the weight of the dry soil (gravimetric water content). The gravimetric water content is converted to volumetric using the bulk density.

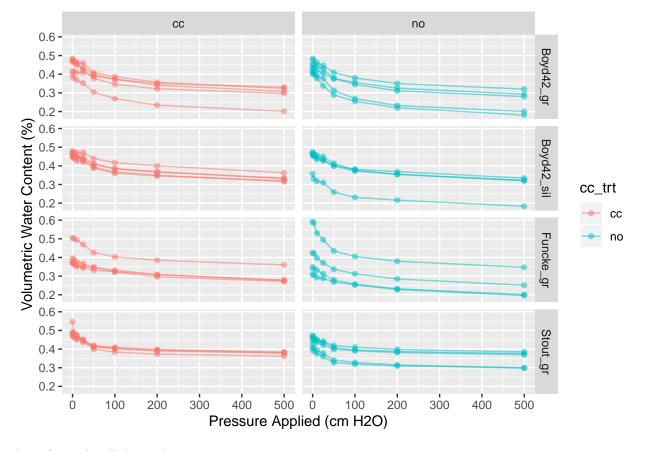
```
dat4 <-
  dat3 %>%
  left join(dat1) %>%
  mutate(gtheta = cumwater_g / drysoil_g,
         vtheta = gtheta * bulkden_gcm3) %>%
  ungroup() %>%
  mutate_if(is.numeric, round, 3)
head(dat4)
## # A tibble: 6 x 8
             {\tt press\_cm\ water\_g\ cumwater\_g\ drysoil\_g\ bulkden\_gcm3\ gtheta\ vtheta}
##
     code
##
     <chr>>
                <dbl>
                        <dbl>
                                    <dbl>
                                              <dbl>
                                                            <dbl> <dbl> <dbl>
## 1 B42-p10
                  999
                      110.
                                     110.
                                               541.
                                                            1.56 0.204 0.318
## 2 B42-p10
                  500
                         4.56
                                     115.
                                               541.
                                                            1.56 0.213 0.331
## 3 B42-p10
                  200
                       12.6
                                     128.
                                               541.
                                                             1.56 0.236 0.367
## 4 B42-p10
                  100
                         5.93
                                     134.
                                               541.
                                                             1.56 0.247 0.384
## 5 B42-p10
                   50
                         8.03
                                     142.
                                               541.
                                                             1.56 0.262 0.407
## 6 B42-p10
                   25
                         9.82
                                     151.
                                               541.
                                                             1.56 0.28
                                                                          0.436
```

5. Join the data with the key so I have treatment info.

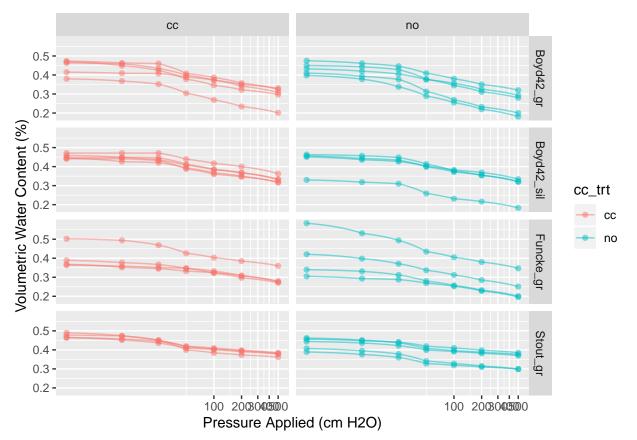
```
## 1 B42-p10 Boyd42
                                                                       0.204 0.318
                        sil
                                 СС
                                            2
                                                       1.56
                                                                  999
## 2 B42-p10 Boyd42
                        sil
                                            2
                                                       1.56
                                                                  500
                                                                       0.213
                                                                              0.331
                                 СС
## 3 B42-p10 Boyd42
                                            2
                                                                               0.367
                        sil
                                 СС
                                                       1.56
                                                                  200
                                                                       0.236
## 4 B42-p10 Boyd42
                                            2
                                                       1.56
                                                                  100
                                                                       0.247
                                                                               0.384
                        sil
                                 СС
                                            2
## 5 B42-p10 Boyd42
                        sil
                                 СС
                                                       1.56
                                                                   50
                                                                       0.262
                                                                               0.407
## 6 B42-p10 Boyd42
                                            2
                                                       1.56
                                                                   25
                                                                       0.28
                                                                               0.436
                        sil
                                 СС
```

Figures

Data for each cell, no transformation



Data for each cell, log-scale



Data averaged for each treatment, no transformation

