Fluid Data

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November 11, 2017

Introduction

The cdata R package provides a powerful extension of the "fluid data" (or "coordinatized data") concept (please see here for some notes) that goes way beyond the concepts of pivot/un-pivot.

The fluid data concept is:

- 1) Data cells have coordinates, and the dependence of these coordinates on a given data representation (a table or map) is an inessential detail to be abstracted out.
- 2) There may not be one "preferred" shape (or frame of reference) for data: you have to anticipate changing data shape many times to adapt to the tasks and tools (data relativity).

cdata supplies two general operators for fluid data work at database scale (and Spark big data scale):

- 1) moveValuesToRows*(): operators centered around SQL cross-join semantics. un-pivot, tidyr::gather(), and cdata::unpivotValuesToRows() are special cases of this general operator.
- 2) moveValuesToColumns*(): an operator centered around SQL group by semantics. pivot, tidyr::spread(), cdata::pivotValuesToColumns(), transpose, and one-hot-encoding are special cases of this general operator.

Because these operators are powerful, they are fairly general, and at first hard to mentally model (especially if you insist on think of them in only in terms of more a specialized operator such as pivot, instead of more general relational concepts such as "cross join" and "group by"). These operators are thin wrappers populating and enforcing a few invariants over a large SQL statement. That does not mean that these operators are trivial, they are thin because SQL is powerful and we have a good abstraction.

Due to the very detailed and explicit controls used in these operatorsthey are very comprehensible once studied. We will follow-up later with additional training material to make quicker comprehension available to more readers. This document is limiting itself to being a mere concise statement of and demonstration of the operators.

Data coordinate notation theory

We are going to introduce a explicit, dense, and powerful data coordinate notation.

Consider the following table that we call a "control table":

```
suppressPackageStartupMessages(library("cdata"))
packageVersion("cdata")
## [1] '0.5.1'
suppressPackageStartupMessages(library("dplyr"))
options(width = 160)
tng <- cdata::makeTempNameGenerator('fdexample')</pre>
controlTable <- dplyr::tribble(~group, ~col1, ~col2,</pre>
                                 'aa', 'c1', 'c2',
                                 'bb', 'c3', 'c4')
knitr::kable(controlTable)
                                    col2
                       group
                              col1
                       aa
                              c1
                                    c2
                       bb
                              c3
                                    c4
```

Control tables partially specify a change of data shape or change of data cell coordinates.

The specification is interpreted as follows:

The region controlTable[, 2:ncol(controlTable)] specifies partial coordinates of data cells in another table. In our example these partial coordinates are "c1", "c2", "c3", and "c4" treated as column names. For example if our data is:

ID	c1	c2	c3	c4
id1	val_id1_c1	val_id1_c2	val_id1_c3	val_id1_c4
id2	val_id2_c1	val_id2_c2	val_id2_c3	val_id2_c4
id3	val_id3_c1	val_id3_c2	val_id3_c3	val_id3_c4

Then each data cell in dat1 (excluding the key-columns, in this case "ID") is named by the row-id (stored in the ID column) plus the column-name ("c1", "c2", "c3", and "c4"). Knowing ID plus the column name unique identifies the data-caring cell in table dat1.

However, there is an alternate cell naming available from the controlTable notation. Each name in the region controlTable[
, 2:ncol(controlTable)] is itself uniquely named by the group entry and column name of the control table itself. This means we have the following correspondence from the partial names "c1", "c2", "c3", and "c4" to a new set of partial names:

```
namePairings <- expand.grid(seq_len(nrow(controlTable)),</pre>
                     2:ncol(controlTable))
colnames(namePairings) <- c("controlI", "controlJ")</pre>
namePairings$coords style1 <-
  vapply(seq_len(nrow(namePairings)),
         function(ii) {
           as.character(paste("column:",
                               controlTable[namePairings$controlI[[ii]],
                                            namePairings$controlJ[[ii]]]))
         },
         character(1))
namePairings$coords_style2 <-
  vapply(seq_len(nrow(namePairings)),
         function(ii) {
           paste("group:",
                 controlTable$group[[namePairings$controlI[[ii]]]],
                 ", column:",
                 colnames(controlTable)[[namePairings$controlJ[[ii]]]])
         },
         character(1))
as.matrix(namePairings[ , c("coords_style1", "coords_style2")])
        coords_style1 coords_style2
##
## [1,] "column: c1"
                      "group: aa , column: col1"
## [2,] "column: c3"
                      "group: bb , column: col1"
                      "group: aa , column: col2"
## [3,] "column: c2"
## [4,] "column: c4"
                      "group: bb , column: col2"
```

The idea is the control table is a very succinct description of the pairing of the namePairings\$coords_style1 cell partial coordinates and the namePairings\$coords_style2 cell partial coordinates. As we have said the namePairings\$coords_style1 cell partial coordinates become full cell coordinates for the data cells in dat1 when combined with the dat1 ID column. The namePairings\$coords_style2 are part of a natural naming for the data cells in the following table:

ID	group	col1	col2
id1	aa	val_id1_c1	val_id1_c2
id1	bb	val_id1_c3	val_id1_c4
id2	aa	val_id2_c1	val_id2_c2
id2	bb	val_id2_c3	val_id2_c4
id3	aa	val_id3_c1	val_id3_c2
id3	bb	val_id3_c3	val_id3_c4

For dat2 the composite row-key (ID, group) plus the column name (one of col1 or col2) gives us the positions of the data carrying cells.

So essentially the two readings of controlTable are a succinct representation of the explicit pairing of data cell coordinates shown in the namePairings table.

The Operators

In terms of the above notation/theory our two operators moveValuesToRows*() and moveValuesToColumns*() are (in principle) easy to describe:

- moveValuesToRows*() reshapes data from style 1 to style 2
- moveValuesToColumns*() reshapes data from style 2 to style 1.

The above is certainly succinct, but carries a lot of information and allows for a lot of different possible applications. Many important applications are derived from how these two operators interact with row-operations and column-operations.

We give simple examples of each of the operators below.

```
moveValuesToRows*()
wideTableName <- 'dat'
d <- dplyr::copy_to(my_db,</pre>
```

```
dplyr::tribble(
        ~ID,
                      ~c1,
                                    ~c2,
                                                  ~c3.
                                                                 ~c4,
      'id1', 'val_id1_c1', 'val_id1_c2', 'val_id1_c3', 'val_id1_c4',
      'id2', 'val_id2_c1', 'val_id2_c2', 'val_id2_c3', 'val_id2_c4',
      'id3', 'val_id3_c1', 'val_id3_c2', 'val_id3_c3', 'val_id3_c4' ),
             wideTableName, overwrite = TRUE, temporary=TRUE)
controlTable <- dplyr::tribble(~group, ~col1, ~col2,</pre>
                                  'aa', 'c1', 'c2',
                                  'bb', 'c3', 'c4')
columnsToCopy <- 'ID'</pre>
cdata::moveValuesToRowsN(wideTable = wideTableName,
                         controlTable = controlTable,
                         my_db = my_db,
                         columnsToCopy = columnsToCopy,
                         tempNameGenerator = tng) %>%
  dplyr::tbl(my_db, .) %>%
  arrange(ID, group) %>%
  knitr::kable()
             ID
                        col1
                                     col2
                  group
             id1
                         val id1 c1 val id1 c2
                  aa
             id1
                  bb
                         val id1 c3 val id1 c4
             id2
                         val_id2_c1 val_id2_c2
                  aa
             id2
                  bb
                         val_id2_c3 val_id2_c4
             id3
                  aa
                         val_id3_c1
                                    val_id3_c2
             id3
                 bb
                         val id3 c3 val id3 c4
moveValuesToColumns*()
tallTableName <- 'dat'
d <- dplyr::copy_to(my_db,</pre>
 dplyr::tribble(
          ~group, ~col1,
                                      ~col2,
   ~ID,
   "id1", "aa",
                  "val_id1_gaa_col1", "val_id1_gaa_col2",
   "id1", "bb",
                  "val_id1_gbb_col1", "val_id1_gbb_col2",
```

"val_id2_gaa_col1", "val_id2_gaa_col2",

"id2", "aa",

ID	c1	c2	c3	c4
id1	val_id1_gaa_col1	$val_id1_gaa_col2$	$val_id1_gbb_col1$	$val_id1_gbb_col2$
id2	$val_id2_gaa_col1$	$val_id2_gaa_col2$	$val_id2_gbb_col1$	$val_id2_gbb_col2$
id3	$val_id3_gaa_col1$	$val_id3_gaa_col2$	$val_id3_gbb_col1$	$val_id3_gbb_col2$

Pivot/Un-Pivot

Pivot and un-pivot (or tidyr::spread() and tidyr::gather()) are special cases of the moveValuesToColumns*() and moveValuesToRows*() operators. Pivot/un-pivot are the cases where the control table has two columns.

Pivot

```
d <- data.frame(
  index = c(1, 2, 3, 1, 2, 3),
  meastype = c('meas1', 'meas1', 'meas1', 'meas2', 'meas2', 'meas2'),
  meas = c('m1_1', 'm1_2', 'm1_3', 'm2_1', 'm2_2', 'm2_3'),
  stringsAsFactors = FALSE)
knitr::kable(d)</pre>
```

 index
 meastype
 meas

 1
 meas1
 m1_1

 2
 meas1
 m1_2

 3
 meas1
 m1_3

 1
 meas2
 m2_1

 2
 meas2
 m2_2

 3
 meas2
 m2_3

```
# the cdata::pivotValuesToColumns version
# equivalent to tidyr::spread(d, 'meastype', 'meas')
```

index	meastype_meas1	meastype_meas2
1	m1_1	m2_1
2	$m1_2$	$m2_2$
3	$m1_3$	$m2_3$

knitr::kable(controlTable)

meastype	meas
meas1	meastype_meas1
meas2	meastype_meas2

index	$meastype_meas1$	meastype_meas2
1	m1_1	m2_1
2	$m1_2$	$m2_2$
3	m1_3	m2_3

```
Un-Pivot
```

```
d <- data.frame(
  index = c(1, 2, 3),
  info = c('a', 'b', 'c'),
  meas1 = c('m1_1', 'm1_2', 'm1_3'),
  meas2 = c('2.1', '2.2', '2.3'),
  stringsAsFactors = FALSE)
knitr::kable(d)</pre>
```

index	info	meas1	meas2
1	a	m1_1	2.1
2	b	$m1_2$	2.2
3	\mathbf{c}	$m1_3$	2.3

index	info	meastype	meas
1	a	meas1	m1_1
1	a	meas2	2.1
2	b	meas1	$m1_2$
2	b	meas2	2.2
3	\mathbf{c}	meas1	$m1_3$
3	\mathbf{c}	meas2	2.3

meastype	meas
meas1	meas1

index	info	meastype	meas
1	a	meas1	m1_1
1	a	meas2	2.1
2	b	meas1	$m1_2$
2	b	meas2	2.2
3	\mathbf{c}	meas1	$m1_3$
3	\mathbf{c}	meas2	2.3

Additional Interesting Applications

Interesting applications of cdata::moveValuesToRows*() and cdata::moveValuesToColumns*() include situations where tidyr is not available (databases and Spark) and also when the data transformation is not obviously a single pivot or un-pivot.

Row-parallel dispatch

A particularly interesting application is converting many column operations into a single operation using a row-parallel dispatch.

Suppose we had the following data in the following format in our system of record (but with many more column groups and columns):

Common tasks might include totaling columns and computing rates between columns. However, sometimes that is best done in a row-oriented representation (though outside systems may need column oriented, or more denormalized results).

With fluid data the task is easy:

group	purchases	rebates
Q1	Q1purchases	Q1rebates
Q2	Q2purchases	Q2rebates

ID	group	purchases	rebates
1	Q1	20	5
1	Q2	10	3
2	Q1	5	10
2	Q2	6	12

```
# perform the calculation in one easy step
calc <- purchasesTall %>%
  mutate(purchasesPerRebate = purchases/rebates) %>%
  compute(name = "purchasesTallC")
knitr::kable(calc)
```

ID	group	purchases	rebates	purchasesPerRebate
1	Q1	20	5	4.000000
1	Q2	10	3	3.333333
2	Q1	5	10	0.500000
2	Q2	6	12	0.500000

```
# move what we want back
controlTable <- controlTable %>%
 mutate(purchasesPerRebate =
           paste0(group, "purchasesPerRebate"))
knitr::kable(controlTable)
```

Rebate Rebate

```
# notice the step back is not a single
# pivot or un-pivot
# due to the larger controlTable
# (especially if there were more quarters)
result <- moveValuesToColumnsN(tallTable = "purchasesTallC",</pre>
                                keyColumns = "ID",
                                controlTable = controlTable,
                                my_db = my_db,
                                tempNameGenerator = tng) %>%
  dplyr::tbl(my_db, .)
knitr::kable(result)
```

ID	Q1purchases	Q1rebates	Q1purchasesPerRebate	Q2purchases	Q2rebates	Q2purchasesPerRebate
1	20	5	4.0	10	3	3.333333
2	5	10	0.5	6	12	0.500000

The point is: the above can work on a large number of rows and columns (especially on a system such as Spark where row operations are performed in parallel).

The above work pattern is particularly powerful on big data systems when the tall table is built up in pieces by appending data (so only the pivot style step is required).

One-hot encoding

Adding indicators or dummy variables (by one-hot encoding, or other methods) are essentially special cases of the pivot flavor of cdata::moveValuesToColumns*().

Transpose

Transpose is a special case of these operators. In fact the key-columns behave like group specifiers, meaning we can transpose many similarly structured tables at once.

```
group_by/aggregate
```

Many operations that look like a complicated pivot in column format are in fact a simple row operation followed a group_by/aggregate (and optional format conversion).

Some fun

The structure of the control table is so similar to the data expected by moveValuesToColumns*() that you can actually send the control table through moveValuesToColumns*() to illustrate the kernel of the transformation.

```
controlTable <- dplyr::tribble(~group, ~col1, ~col2,</pre>
                                   'aa', 'c1', 'c2',
                                   'bb', 'c3', 'c4')
tallTableName <- 'dc'
d <- dplyr::copy_to(my_db, controlTable, tallTableName)</pre>
keyColumns <- NULL
wideTableName <- moveValuesToColumnsN(tallTable = tallTableName,</pre>
                             controlTable = controlTable,
                             keyColumns = keyColumns,
                             my_db = my_db)
dw <- dplyr::tbl(my_db, wideTableName)</pre>
knitr::kable(dw)
                        c1
                            c2 c3
                                     c4
                        c1
                            c2
                                 c3
                                     c4
```

The transformed table is essentially an example row of the wideform.

And we can, of course, map back.

This "everything maps to a row" form means the control table is

essentially a graphical representation of the desired data transform. It also helps make clear that just about any even more general shape to shape transform can be achieved by a moveValuesToColumnsN() followed by a moveValuesToRowsN().1

```
moveValuesToRowsN(wideTable = wideTableName,
                  controlTable = controlTable,
                  my_db = my_db) \%>\%
  dplyr::tbl(my_db, .) %>%
  arrange(group) %>%
  knitr::kable()
```

group	col1	col2
aa	c1	c2
bb	c3	c4

Conclusion

cdata::moveValuesToRows*() and cdata::moveValuesToColumns*() represent two very general "fluid data" or "coordinatized data" operators that have database scale (via DBI/dbplyr/dplyr) and big data scale implementations (via Sparklyr). Some very powerful data transformations can be translated into the above explicit control table terminology. The extra details explicitly managed in the control table notation makes for clear calling interfaces.

```
for(ti in tng(dumpList = TRUE)) {
  dplyr::db_drop_table(my_db, ti)
DBI::dbDisconnect(my_db)
```

¹ Or by the a moveValuesToRowsN() followed by a moveValuesToColumnsN(). One direction is storing all intermediate values in a single denormalized column, the other is storing in many RDF-triple like rows.