

# Creating an R data set from STAR

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## Abstract

A substantial portion of the data from Tennessee's Student Teacher Achievement Ratio (STAR) project, a large-scale, four-year study of reduced class size, has been made available to the public at <http://www.heros-inc.org/data.htm>. We describe the creation of an R(<http://www.r-project.org>) data set from these data.

## 1 Introduction

The data from the STAR project are available in several different forms from the web site <http://www.heros-inc.org/data.htm>. The most convenient form for creation of an R data set is the tab-delimited text file. Download the archive file <http://www.heros-inc.org/text-star.zip> containing two files: `readme.txt`, a description of the data, and `webstar.txt`, the data themselves.

## 2 Reading the data

From the data description file we can see that there are 53 columns in the data set and most of these columns are coded values. Such columns should be represented as factors in R but many of these columns will need to be combined before we can work with them. We will convert the first 5 columns to factors and leave the remaining 48 columns as integers.

```
> str(orig <- read.delim(unz(system.file("original/text-star.zip",  
+                               package = "mlmRev"),  
+                               "webstar.txt"),  
+                               colCl = rep(c("factor", "integer"), c(5, 48))))
```

```
'data.frame':      11598 obs. of  53 variables:  
 $ NEWID   : Factor w/ 11598 levels "100017","100028",...: 839 943 986 1104 1263 1346 1485 1547 1701 1783 ...  
 $ SSEX    : Factor w/ 3 levels "1","2","9": 2 2 2 1 1 1 1 2 1 1 ...  
 $ SRACE   : Factor w/ 7 levels "1","2","3","4",...: 2 1 2 1 2 1 2 1 1 1 ...  
 $ SBIRTHQ : Factor w/ 5 levels "1","2","3","4",...: 3 1 4 4 1 3 1 4 2 3 ...  
 $ SBIRTHY : Factor w/ 7 levels "1977","1978",...: 3 4 3 3 4 3 3 3 3 3 ...  
 $ STARK   : int    2 1 1 2 1 2 2 2 2 2 ...  
 $ STAR1   : int    2 1 1 2 2 2 2 1 1 1 ...  
 $ STAR2   : int    2 1 1 2 2 1 2 1 1 1 ...  
 $ STAR3   : int    1 1 1 1 2 1 1 1 1 1 ...  
 $ CLTYPEK : int    9 1 1 9 3 9 9 9 9 9 ...  
 $ CLTYPE1 : int    9 1 1 9 9 9 9 3 2 2 ...  
 $ CLTYPE2 : int    9 1 3 9 9 2 9 3 2 2 ...
```

```

$ CLTYPE3 : int 2 1 3 1 9 2 3 3 2 2 ...
$ SCHTYPEK: int 9 3 2 9 1 9 9 9 9 9 ...
$ HDEGK   : int 9 2 2 9 2 9 9 9 9 9 ...
$ CLADK   : int 9 1 1 9 6 9 9 9 9 9 ...
$ TOTEXPK : int 99 7 21 99 0 99 99 99 99 99 ...
$ TRACEK  : int 9 1 1 9 1 9 9 9 9 9 ...
$ TREADSSK: int 999 447 450 999 439 999 999 999 999 999 ...
$ TMATHSSK: int 999 473 536 999 463 999 999 999 999 999 ...
$ SESK    : int 9 2 2 9 1 9 9 9 9 9 ...
$ SCHTYPE1: int 9 3 2 9 9 9 9 3 3 3 ...
$ TRACE1  : int 9 1 2 9 9 9 9 1 1 1 ...
$ HDEG1   : int 9 1 2 9 9 9 9 2 2 1 ...
$ CLAD1   : int 9 4 3 9 9 9 9 2 4 4 ...
$ TOTEXP1 : int 99 7 32 99 99 99 99 8 13 7 ...
$ TREADSS1: int 999 507 579 999 999 999 999 475 999 651 ...
$ TMATHSS1: int 999 538 592 999 999 999 999 512 999 532 ...
$ SES1    : int 9 1 9 9 9 9 9 2 2 2 ...
$ SCHTYPE2: int 9 3 2 9 9 3 9 3 3 3 ...
$ TRACE2  : int 9 1 2 9 9 1 9 2 1 1 ...
$ HDEG2   : int 9 1 1 9 9 1 9 2 1 1 ...
$ CLAD2   : int 9 2 4 9 9 1 9 4 4 4 ...
$ TOTEXP2 : int 99 3 4 99 99 13 99 13 6 8 ...
$ TREADSS2: int 999 568 588 999 999 999 999 573 999 596 ...
$ TMATHSS2: int 999 579 579 999 999 999 999 550 999 590 ...
$ SES2    : int 9 2 2 9 9 2 9 2 2 2 ...
$ SCHTYPE3: int 2 3 2 3 9 3 1 3 3 3 ...
$ TREADSS3: int 580 587 644 686 999 644 999 599 999 626 ...
$ TMATHSS3: int 564 593 639 667 999 648 999 583 999 618 ...
$ SES3    : int 1 1 2 2 9 2 1 2 2 2 ...
$ TRACE3  : int 1 1 1 1 9 1 2 1 1 1 ...
$ HDEG3   : int 1 1 1 1 9 1 1 1 1 2 ...
$ CLAD3   : int 4 2 4 4 9 4 1 6 4 4 ...
$ TOTEXP3 : int 30 1 4 10 99 15 17 23 8 8 ...
$ SYSIDKN : int 999 30 11 999 11 999 999 999 999 999 ...
$ SYSID1N : int 999 30 11 999 999 999 999 4 40 21 ...
$ SYSID2N : int 999 30 11 999 999 6 999 4 40 21 ...
$ SYSID3N : int 22 30 11 6 999 6 11 4 40 21 ...
$ SCHIDKN : int 999 63 20 999 19 999 999 999 999 999 ...
$ SCHID1N : int 999 63 20 999 999 999 999 5 77 50 ...
$ SCHID2N : int 999 63 20 999 999 8 999 5 77 50 ...
$ SCHID3N : int 54 63 20 8 999 8 31 5 77 50 ...

```

In the call to `read.delim` we read directly from the zip archive and avoided expanding the much larger text file. The call to `system.file` determines the name of a file that is part of a package. In practice it is often more convenient to use the `file.choose` function which brings up a file chooser panel.

We also check the form of the original data by calling `str` on the object

```

> str(orig)

'data.frame':      11598 obs. of  53 variables:
 $ NEWID   : Factor w/ 11598 levels "100017","100028",...: 839 943 986 1104 1263 1346 1485 1547 1701 1783 ...
 $ SSEX    : Factor w/ 3 levels "1","2","9": 2 2 2 1 1 1 1 2 1 1 ...
 $ SRACE   : Factor w/ 7 levels "1","2","3","4",...: 2 1 2 1 2 1 2 1 1 1 ...

```

```

$ SBIRTHQ : Factor w/ 5 levels "1","2","3","4",...: 3 1 4 4 1 3 1 4 2 3 ...
$ SBIRTHY : Factor w/ 7 levels "1977","1978",...: 3 4 3 3 4 3 3 3 3 ...
$ STARK   : int    2 1 1 2 1 2 2 2 2 2 ...
$ STAR1   : int    2 1 1 2 2 2 2 1 1 1 ...
$ STAR2   : int    2 1 1 2 2 1 2 1 1 1 ...
$ STAR3   : int    1 1 1 1 2 1 1 1 1 1 ...
$ CLYPEK  : int    9 1 1 9 3 9 9 9 9 9 ...
$ CLYPE1  : int    9 1 1 9 9 9 9 3 2 2 ...
$ CLYPE2  : int    9 1 3 9 9 2 9 3 2 2 ...
$ CLYPE3  : int    2 1 3 1 9 2 3 3 2 2 ...
$ SCHYPEK : int    9 3 2 9 1 9 9 9 9 9 ...
$ HDEGK   : int    9 2 2 9 2 9 9 9 9 9 ...
$ CLADK   : int    9 1 1 9 6 9 9 9 9 9 ...
$ TOTEXPK : int   99 7 21 99 0 99 99 99 99 ...
$ TRACEK  : int    9 1 1 9 1 9 9 9 9 9 ...
$ TREADSSK: int   999 447 450 999 439 999 999 999 999 ...
$ TMATHSSK: int   999 473 536 999 463 999 999 999 999 ...
$ SESK    : int    9 2 2 9 1 9 9 9 9 9 ...
$ SCHYPE1 : int    9 3 2 9 9 9 9 3 3 3 ...
$ TRACE1  : int    9 1 2 9 9 9 9 1 1 1 ...
$ HDEG1   : int    9 1 2 9 9 9 9 2 2 1 ...
$ CLAD1   : int    9 4 3 9 9 9 9 2 4 4 ...
$ TOTEXP1 : int   99 7 32 99 99 99 99 8 13 7 ...
$ TREADSS1: int   999 507 579 999 999 999 999 475 999 651 ...
$ TMATHSS1: int   999 538 592 999 999 999 999 512 999 532 ...
$ SES1    : int    9 1 9 9 9 9 9 2 2 2 ...
$ SCHYPE2 : int    9 3 2 9 9 3 9 3 3 3 ...
$ TRACE2  : int    9 1 2 9 9 1 9 2 1 1 ...
$ HDEG2   : int    9 1 1 9 9 1 9 2 1 1 ...
$ CLAD2   : int    9 2 4 9 9 1 9 4 4 4 ...
$ TOTEXP2 : int   99 3 4 99 99 13 99 13 6 8 ...
$ TREADSS2: int   999 568 588 999 999 999 999 573 999 596 ...
$ TMATHSS2: int   999 579 579 999 999 999 999 550 999 590 ...
$ SES2    : int    9 2 2 9 9 2 9 2 2 2 ...
$ SCHYPE3 : int    2 3 2 3 9 3 1 3 3 3 ...
$ TREADSS3: int   580 587 644 686 999 644 999 599 999 626 ...
$ TMATHSS3: int   564 593 639 667 999 648 999 583 999 618 ...
$ SES3    : int    1 1 2 2 9 2 1 2 2 2 ...
$ TRACE3  : int    1 1 1 1 9 1 2 1 1 1 ...
$ HDEG3   : int    1 1 1 1 9 1 1 1 1 2 ...
$ CLAD3   : int    4 2 4 4 9 4 1 6 4 4 ...
$ TOTEXP3 : int   30 1 4 10 99 15 17 23 8 8 ...
$ SYSIDKN : int   999 30 11 999 11 999 999 999 999 999 ...
$ SYSID1N : int   999 30 11 999 999 999 999 4 40 21 ...
$ SYSID2N : int   999 30 11 999 999 6 999 4 40 21 ...
$ SYSID3N : int   22 30 11 6 999 6 11 4 40 21 ...
$ SCHIDKN : int   999 63 20 999 19 999 999 999 999 999 ...
$ SCHID1N : int   999 63 20 999 999 999 999 5 77 50 ...
$ SCHID2N : int   999 63 20 999 999 8 999 5 77 50 ...
$ SCHID3N : int   54 63 20 8 999 8 31 5 77 50 ...

```

## 2.1 Missing value codes

All the columns except the first column have missing values present. Typically the missing value code is "9" but "99", "999" and "9999" are also used. We convert these to R's missing value code `NA` column by column.

```
> mv <- rep("9", 53)
> mv[c(4,17,26,34,45)] <- "99"
> mv[c(19,20,27,28,35,36,39,40,46:53)] <- "999"
> mv[5] <- "9999"
> mv[1] <- "999999"
> for (i in seq(a = orig)) orig[[i]][orig[[i]] == mv[i]] <- NA
> summary(orig[1:5])
```

```
      NEWID      SSEX      SRACE      SBIRTHQ
100017 :    1    1    :6122    1    :7193    1    :2836
100028 :    1    2    :5456    2    :4173    2    :2851
100045 :    1    9     :   0    3     :   32    3    :3422
100064 :    1  NA's:   20    4     :   21    4    :2423
100070 :    1                    6     :   20   99   :   0
100096 :    1                (Other):  14  NA's:  66
(Other):11592                NA's   : 145

      SBIRTHY
1980   :6886
1979   :3915
1978   : 645
1977   :  58
1981   :  24
(Other):  1
NA's   :  69
```

Notice that level "9" is still present for the `SSEX` variable even after all the observations at that level have been replaced by the missing value code. To remove these unused levels from this and all the other columns, we loop over the columns selecting all the values but using the optional argument `drop = TRUE`.

```
> for (i in seq(a = orig)) orig[[i]] <- orig[[i]][drop = TRUE]
> summary(orig[1:5])
```

```
      NEWID      SSEX      SRACE      SBIRTHQ      SBIRTHY
100017 :    1    1    :6122    1    :7193    1    :2836  1977:  58
100028 :    1    2    :5456    2    :4173    2    :2851  1978: 645
100045 :    1  NA's:   20    3     :   32    3    :3422  1979:3915
100064 :    1                    4     :   21    4    :2423  1980:6886
100070 :    1                    5     :   14  NA's:  66  1981:  24
100096 :    1                    6     :   20          1982:  1
(Other):11592                NA's: 145          NA's:  69
```

For convenience we convert the names of the columns to lower case.

```
> names(orig) <- tolower(names(orig))
```

## 3 Setting factor levels

In R the levels of a factor can be given meaningful labels instead of numeric codes and in most cases this eliminates the need for a separate codebook. For example storing the labels of `sex` as "M" and "F" makes the

coding self-explanatory. When used in a model a factor is automatically converted to a set of “contrasts” (there is a technical definition of the term “contrast” in linear models that is not always fulfilled by these derived variables) and the corresponding coefficients are given meaningful names.

When there is a natural ordering of the levels of a factor it can be created as an ordered factor that will preserve this ordering.

The labels can be set after the factor is created or as part of the creation of the factor. Below we will create a “long form” of the data where each row corresponds to a combination of student and grade. In doing this we will need to concatenate related columns of the original data frame. For example, the columns `cltypek`, `cltype1`, `cltype2` and `cltype3` will be concatenated to form a single column `cltype`. If the coding is consistent across the grades then it is easiest to concatenate the integer codes and set the labels on the “long” version of the variable.

However there are two groups of variables, `hdeg` and `clad`, that are not coded consistently. In each case the codes used for kindergarten teachers are different from those used for teachers of grades 1 to 3 classes. The codes for kindergarten teachers are a superset of those for the other teachers but the numbering is not consistent; a bachelor’s degree is coded as 2 for kindergarten but 1 for the others. Thus we cannot combine the numeric values - we must create the labels for each column and then concatenate the labels and convert to a factor.

```
> orig$hdegk <- ordered(orig$hdegk, levels = 1:6,
+                       labels = c("ASSOC", "BS/BA", "MS/MA/MEd", "MA+", "Ed.S", "Ed.D/Ph.D"))
> orig$hdeg1 <- ordered(orig$hdeg1, levels = 1:4,
+                       labels = c("BS/BA", "MS/MA/MEd", "Ed.S", "Ed.D/Ph.D"))
> orig$hdeg2 <- ordered(orig$hdeg2, levels = 1:4,
+                       labels = c("BS/BA", "MS/MA/MEd", "Ed.S", "Ed.D/Ph.D"))
> orig$hdeg3 <- ordered(orig$hdeg3, levels = 1:4,
+                       labels = c("BS/BA", "MS/MA/MEd", "Ed.S", "Ed.D/Ph.D"))
> orig$cladk <- factor(orig$cladk, levels = c(1:3,5:8),
+                     labels = c("1", "2", "3", "APPR", "PROB", "NOT", "PEND"))
> orig$clad1 <- factor(orig$clad1, levels = 1:6,
+                     labels = c("NOT", "APPR", "PROB", "1", "2", "3"))
> orig$clad2 <- factor(orig$clad2, levels = 1:6,
+                     labels = c("NOT", "APPR", "PROB", "1", "2", "3"))
> orig$clad3 <- factor(orig$clad3, levels = 1:6,
+                     labels = c("NOT", "APPR", "PROB", "1", "2", "3"))
```

## 4 Creating separate data frames

These data are represented in a “wide” format where each row corresponds to a student. Some of the columns, such as `ssex`, are indeed a property of the student; some, such as `hdegk` are properties of teachers; some, such as `schttypek` are properties of schools or classes in schools; and some are unique to a student/grade combination. We will create separate frames for each of these types.

The first 5 columns are student-level data

```
> student <- orig[1:5]
> names(student) <- c("id", "sx", "eth", "birthq", "birthy")
> levels(student$sx) <- c("M", "F")
> levels(student$eth) <- c("W", "B", "A", "H", "I", "O")
> student$birthy <- ordered(student$birthy)
> student$birthq <- ordered(paste(student$birthy, student$birthq, sep=":"))
> summary(student)
```

	id		sx		eth		birthq		birthy
100017 :	1	M	:6122	W	:7193	1980:3	:2304	1977:	58

```

100028 :    1  F    :5456  B    :4173  1980:1 :2221  1978: 645
100045 :    1  NA's: 20  A    :   32  1980:2 :2190  1979:3915
100064 :    1                    H    :   21  1979:4 :1879  1980:6886
100070 :    1                    I    :   14  1979:3 :   923  1981:   24
100096 :    1                    O    :   20  1979:2 :   586  1982:    1
(Other):11592                    NA's: 145  (Other):1495  NA's: 69

```

The other columns refer to a combination of the student and grade. We first create an expanded or “long” version of the table with a row for each student/grade combination.

To create the long version of the table we repeat the student ids four times and add a column for the grade level. Related groups of columns, such as `cltypek`, `cltype1`, `cltype2` and `cltype3`, are concatenated then converted to a factor. However, there are two groups, `hdeg` and `clad`, for which this approach will not work because these groups are not encoded consistently.

```

> long <- data.frame(id = rep(orig$newid, 4),
+                   gr = ordered(rep(c("K", 1:3), each = nrow(orig)),
+                               levels = c("K", 1:3)),
+                   star = factor(unlist(orig[6:9])),
+                   cltype = factor(unlist(orig[10:13])),
+                   schtype = factor(unlist(orig[c(14,22,30,38)])),
+                   hdeg = ordered(unlist(lapply(orig[c(15,24,32,43)], as.character)),
+                                 levels = c("ASSOC", "BS/BA", "MS/MA/MEd", "MA+", "Ed.S", "Ed.D/Ph.D")),
+                   clad = factor(unlist(lapply(orig[c(16,25,33,44)], as.character)),
+                                 levels = c("NOT", "APPR", "PROB", "PEND", "1", "2", "3")),
+                   exp = unlist(orig[c(17,26,34,45)]),
+                   trace = factor(unlist(orig[c(18,23,31,42)]), levels=1:6,
+                                 labels=c("W", "B", "A", "H", "I", "O")),
+                   read = unlist(orig[c(19,27,35,39)]),
+                   math = unlist(orig[c(20,28,36,40)]),
+                   ses = factor(unlist(orig[c(21,29,37,41)]), labels=c("F", "N")),
+                   sch = factor(unlist(orig[50:53])))

```

We can now eliminate the combinations that are completely missing. Checking

```
> summary(long)
```

id	gr	star	cltype	schtype
100017 : 4	K:11598	1:26796	1 : 8015	1 : 5624
100028 : 4	1:11598	2:19596	2 : 9192	2 : 6428
100045 : 4	2:11598		3 : 9589	3 :12561
100064 : 4	3:11598		NA's:19596	4 : 2183
100070 : 4				NA's:19596
100096 : 4				
(Other):46368				
hdeg	clad	exp	trace	
ASSOC : 0	1 :18303	Min. : 0.00	W :21550	
BS/BA :16586	APPR : 2030	1st Qu.: 5.00	B : 5005	
MS/MA/MEd: 9587	PROB : 1961	Median :11.00	A : 14	
MA+ : 161	NOT : 1757	Mean :12.04	H : 0	
Ed.S : 237	3 : 1059	3rd Qu.:17.00	I : 0	
Ed.D/Ph.D: 58	(Other): 877	Max. :42.00	O : 0	
NA's :19763	NA's :20405	NA's :19789	NA's:19823	
read	math	ses	sch	

```

Min. :315.0 Min. :288.0 F :13111 51 : 826
1st Qu.:467.0 1st Qu.:505.0 N :12858 27 : 562
Median :552.0 Median :557.0 NA's:20423 9 : 543
Mean :539.9 Mean :553.7 22 : 534
3rd Qu.:604.0 3rd Qu.:603.0 63 : 534
Max. :775.0 Max. :774.0 (Other):23797
NA's :22130 NA's :21779 NA's :19596

```

indicates that fewest missing values are in the `sch`, `cltype`, and `sctype` columns. They are also consistent

```
> with(long, all.equal(is.na(sctype), is.na(sch)))
```

```
[1] TRUE
```

```
> with(long, all.equal(is.na(cltype), is.na(sch)))
```

```
[1] TRUE
```

hence we use these to subset the data frame

```
> long <- long[!is.na(long$sch),]
```

It turns out that we could have used the `star` column as this simply indicates if the student was in the study that year.

```
> summary(long[1:5])
```

```

      id      gr      star      cltype      sctype
100173 :    4 K:6325  1:26796  1:8015  1: 5624
100201 :    4 1:6829  2:    0  2:9192  2: 6428
10023  :    4 2:6840           3:9589  3:12561
100236 :    4 3:6802           4: 2183
100302 :    4
100361 :    4
(Other):26772

```

Because it now contains no information we will drop it.

```
> long$star <- NULL
```

For convenience we set the row names of this data frame to be a combination of the student id and the grade.

```
> rownames(long) <- paste(long$id, long$gr, sep = '/')
```

We can extract the school-level data from this table.

```
> school <- unique(long[, c("sch", "sctype")])
```

```
> length(levels(school$sch)) == nrow(school)
```

```
[1] TRUE
```

```
> row.names(school) <- school$sch
```

```
> school <- school[order(as.integer(as.character(school$sch))),]
```

```
> long$sctype <- NULL
```

```
> levels(school$sctype) <- c("inner", "suburb", "rural", "urban")
```

```
> levels(long$cltype) <- c("small", "reg", "reg+A")
```

We can create a merged data set with

```
> star <- merge(merge(long, school, by = "sch"), student, by = "id")
```

```
> star$time <- as.integer(star$gr) - 1
```

## 5 Assigning teacher ids

There are no teacher id numbers available but we can obtain a reasonably accurate surrogate by determining the unique combinations of all the variables associated with the teacher.

```
> teacher <- unique(star[, c("cltype", "trace", "exp", "clad", "gr",
+                           "hdeg", "sch")])
> teacher <- teacher[with(teacher, order(sch, gr, cltype, exp, hdeg,
+                                       clad, trace)), ]
```

To generate the correspondence between the observations and the teacher we create labels that incorporate the levels of each of the variables that defined the unique combinations.

```
> row.names(teacher) <- tch <- teacher$tch <- seq(nrow(teacher))
> names(tch) <- do.call("paste", c(teacher[,1:7], list(sep=":")))
> star$tch <- tch[do.call("paste", c(star[c("cltype", "trace", "exp",
+                                       "clad", "gr", "hdeg", "sch")],
+                                   list(sep = ":")))]
```

We can check if this is successful by generating tables of class sizes.

```
> table(table(star$tch))

 1  2  3 11 12 13 14 15 16 17 18 19 20 21 22 23 24
45  4  1  2 17 68 81 116 111 106 28 41 49 103 152 137 138
25 26 27 28 29 30 32 44 46
83 47 29 13 11  2  1  1  1
```

```
> table(table(subset(star, cltype == "small")$tch))

 1  2 11 12 13 14 15 16 17 18 19 20 32
12  1  2 17 68 81 114 108 100 22  9  1  1
```

```
> table(table(subset(star, cltype == "reg")$tch))

 1  2 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30
21  1  1  2  5  4 16 31 47 84 64 70 38 20 11  4  5  1
```

```
> table(table(subset(star, cltype == "reg+A")$tch))

 1  2  3 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 44 46
12  2  1  1  1  1  2 16 17 56 68 73 68 45 27 18  9  6  1  1  1
```

We see that there are three classes with sizes greater than 30 and that one of these is labelled as a “small” class. It is likely that each of these represents two or more classes but we do not have enough information to distinguish them.

## 6 Initial model fits

Some initial model fits are

```
> library(lme4)
> (mm1 <- lmer(math ~ gr + sx + eth + cltype + (1/id) + (1/sch), star))
```



```

Linear mixed model fit by REML ['lmerMod']
Formula: math ~ gr + sx + eth + cltype + (1 | id) + (1 | sch)
Data: star
REML criterion at convergence: 245140
Random effects:
Groups   Name          Std.Dev.
id       (Intercept) 31.35
sch      (Intercept) 11.07
Residual                26.20
Number of obs: 24578, groups: id, 10732; sch, 80
Fixed Effects:
(Intercept)      gr.L      gr.Q      gr.C      sxF
      560.498      96.412     -4.580     -3.460      2.953
      ethB      ethA      ethH      ethI      eth0
     -22.891      2.128      1.173     -34.783      2.419
cltypereg cltypereg+A
     -7.111      -5.910

> (rm1 <- lmer(read ~ gr + sx + eth + cltype + (1|id) + (1|sch), star))

```

```

Linear mixed model fit by REML ['lmerMod']
Formula: read ~ gr + sx + eth + cltype + (1 | id) + (1 | sch)
Data: star
REML criterion at convergence: 241465
Random effects:
Groups   Name          Std.Dev.
id       (Intercept) 30.72
sch      (Intercept) 10.46
Residual                26.31
Number of obs: 24226, groups: id, 10621; sch, 80
Fixed Effects:
(Intercept)      gr.L      gr.Q      gr.C      sxF
      541.701      131.403     -28.207     -1.630      9.061
      ethB      ethA      ethH      ethI      eth0
     -18.807      8.503      1.942     -32.098      7.635
cltypereg cltypereg+A
     -7.828      -4.745

```

## 7 Session Info

```

> toLatex(sessionInfo())

```

- R version 3.5.2 (2018-12-20), x86\_64-w64-mingw32
- Locale: LC\_COLLATE=C, LC\_CTYPE=English\_Canada.1252, LC\_MONETARY=English\_Canada.1252, LC\_NUMERIC=C, LC\_TIME=English\_Canada.1252
- Running under: Windows 10 x64 (build 17763)
- Matrix products: default
- Base packages: base, datasets, grDevices, graphics, methods, stats, utils
- Other packages: Matrix 1.2-15, lattice 0.20-38, lme4 1.1-20, mlmRev 1.0-8

- Loaded via a namespace (and not attached): MASS 7.3-51.1, Rcpp 1.0.3, compiler 3.5.2, grid 3.5.2, minqa 1.2.4, nlme 3.1-137, nloptr 1.2.1, splines 3.5.2, tools 3.5.2