an introduction to R for epidemiologists categorical variables, time and regular expressions

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## Outline



coding categorical variables





### Outline

#### categorical variables

coding categorical variables

#### 2 dates and times

#### B) regular expressions

# creating intervals cut()

- want 7 age categories
  - less than 1, 1 to 4, 5 to 14, 15 to 24, 25 to 44, 45 to 64, older 65

- creates factor with 7 levels
- notation (15, 25] interval "open" (parenthesis) on left (> 15) and "closed" (bracket) on the right boundary ( $\leq$  25)
- "right = FALSE" closed on the left and open on the right: [a, b) use

# add labels to intervals "labels="

clarify the (a, b] interval bracket notation

# assigning intervals

create a categorical character vector

```
agecat2<-character(0)
agecat2[ages<1] <- "<1"
agecat2[ages>=1 & ages<5] <- "1-4"
agecat2[ages>=5 & ages<15] <- "5-14"
agecat2[ages>=15 & ages<25] <- "15-24"
agecat2[ages>=25 & ages<45] <- "25-44"
agecat2[ages>=45 & ages<65] <- "45-64"
agecat2[ages>=65] <- "65+"
table(agecat2)</pre>
```

### about factors the good, the bad, and the ugly

- factors are default R categorical variables
  - integers with "names"
  - optimized for modeling functions (e.g. lmer) and graphics (ggplot)
  - preclude having to keep track of assigned levels (vs. data dictionaries)
- but can be a pain
  - behave weirdly
  - get in way of data manipulations (e.g. merges)
  - probably not necessary most of the time
- read.table automatically converts all strings to factors
  - make "stringsAsFactors=F" part of your vocabulary (should be default...)
  - unclass() returns integers (as.integer does not preserve levels)

### Outline



• coding categorical variables



#### regular expressions

# setting a referent category relevel()

```
names <- c("White", "Black", "Latino", "Asian")
ethnic <- sample(names, 100, replace = T)
ethnic <- factor(ethnic, levels = names)
ethnic
levels(ethnic)
ethnic2 <- relevel(ethnic, ref = "Latino")
levels(ethnic)
unclass(ethnic)
unclass(ethnic2)</pre>
```

#### factors as dummy variables

- generally create k 1 dichotomous variables, each coded 0 or 1
- in R, just create a single factor with the desired number of levels and set the reference level

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### Outline

# categorical variables coding categorical variables

### 2 dates and times

#### B regular expressions

### over view of date objects

from Tomas Aragon, "Applied Epidemiology Using R"



Fig. 3.4. Displayed are functions to convert calendar date and time data into R date-time classes (as.Date, strptime, as.POSIX1t, as.POSIXct), and the format function converts date-time objects into character dates, days, weeks, months, times,

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### from days to dates as.Date()

- yyyy-mm-dd
- Julian dates number of days since January 1, 1970
- specify the format of the input calendar date myDays <- c("10/11/1945", "8/19/2003", "5/15/1964") myDates<- as.Date(myDays, format = "%m/%d/%Y") myDates
- looks like a character, but is class "date", mode "numeric"
- displayed in a standard format (yyyy-mm-dd)
- help(strptime) to get conversion formats see .

```
as.numeric(myDates)
#calculate age (as of today)
today <- Sys.Date()
today
age <- (today - myDates)/365.25
age
age2 <- trunc(as.numeric(age))
age2</pre>
```

- "%a" Abbreviated weekday name.
- "%A" Full weekday name.
- "%b" Abbreviated month name.
- "%B" Full month name.
- "%d" Day of the month as decimal number (01-31)
- "%j" Day of year as decimal number (001-366).
- "%m" Month as decimal number (01-12).
- "%U" Week of the year as decimal number (00-53) Sunday day 1 "%w" Weekday as decimal number (0-6, Sunday is 0).
- "%W" Week of the year as decimal number (00-53) Monday day 1 "%y" Year without century (00-99). (not recommended)

```
as.Date("December 8, 1989", format = "%B %d, %Y")
as.Date("12/8/1989", format = "%m/%d/%Y")
as.Date("12/8/89", format = "%m/%d/%y") #caution 2-digit year
as.Date("08Dec1989", format = "%d%b%Y")
as.Date("08Dec89", format = "%d%b%y")
#standard does not require format
as.Date("1989-12-08")
```

extractions

# extracting info from date objects: weekdays(), months(), quarters(), julian()

weekdays(myDates)
months(myDates)
quarters(myDates)
julian(myDates)

#### date objects as integers

as.Date("2004-01-15"):as.Date("2004-01-23") seq(as.Date("2004-01-15"), as.Date("2004-01-18"), by = 1)

# from dates to time strptime()

• strptime() - accepts both dates and times (HH:MM:SS)

# POSIXIt objects

Portable Operating System Interface, legible time

- strptime() produces named list, class POSIXIt
- POSIXIt vector objects:
  - 'sec' (0-61), 'min' (0-59), 'hour' (0-23), 'mday' (1-31), 'mon' (0-11), 'year' (since 1900), 'wday' (0-6, starting on Sunday), 'yday' (0-365), and 'isdst' (DST flag, + if in force, 0 if not, if unknown)

#### working with elements of POSIXIt date time list

dateTime\$min dateTime\$hour dateTime\$mon dateTime\$wday

- POSIXct based on continuous time in seconds
- as.POSIXct()/as.POSIXct() to coerce

### date-time outputs format()

decjan <- seq(as.Date("2003-12-15"), as.Date("2004-01-15"), by =1)

disease.week <- format(decjan, "%U")</pre>

- %U for weeks starting on Sunday,
- %W for weeks starting on Monday

# working with time read in time data

- read in 1000 motor vehicle crash times (NYPD data)
- note, even if correct Julian, still need to specify format

daytime<-read.csv("/~daytime.csv", stringsAsFactors=F)</pre>

```
head(daytime)
daytime$x[1:20]
class(daytime$x)
```

dateTime<-strptime(daytime\$x, "%Y-%M-%d %H:%M:%S")
class(dateTime)</pre>

# working with time

extract specific times

- create logical vectors and index
- use weekdays(), months()
- o format() for times

```
weekdays(dateTime)=="Sunday"
months(dateTime)=="July"
day<- !weekdays(dateTime)=="Sunday" &
    !weekdays(dateTime)=="Saturday"
month<-!months(dateTime)=="July" &
    !months(dateTime)=="August"
hours <- format(dateTime, "%H") #extract hours from dateTime
sch.hours<-(hours>="07" & hours<="09" ) |
    (hours>="02" & hours<="04")</pre>
```

### Outline



#### 2 dates and times

#### 3 regular expressions

- search for and return index location of character or string
- search and replace, e.g. code ICD-9 into fewer, mutually exclusive categories
- grep("x", dataObject)
- flexible (bewildering?) array of "meta" characters to specify search patterns
- google and StackOverflow are your friends...

## 5 kinds of searchs

and some meta characters

- character .
- class group of characters []
- concatenation string together, e.g. a word
- repetition ? (optional or once) \* (absent or any number) + (at least once)
- alternation Or statement |

## some "metacharacters" (to control matches)

- ^ 2 possible uses, beginning of a string vs. NOT if inside character class brackets
- \$ end of a string
- . any single character (except a newline)
- ? occurring once, or not at all
- + at least once
- \* any number of times
- .\* any number of characters other than a newline
- 0-9, a-z, A-Z shortcuts for ranges of numbers or letters
- \ escape, returns literal for meta-characters

## match a single character

metacharacters

```
vec1 <- c("x", "xa bc", "abc", "ax bc", "ab xc", "ab cx")
grep("x", vec1) # returns integer vector matches
vec1[grep("x", vec1)] #index by position
grep("^x", vec1) # caret ^ matches beginning of line
vec1[grep("^x", vec1)]
vec1[grep("x$", vec1)] # $ metacharacter for end of line
# front space for beginning of a word, but not beginning of a line
vec1[grep(" x", vec1)]
# back space for end of a word, but not end of a line
vec1[grep("x ", vec1)]
```

```
# period matches any single character, including a space.
vec1[grep(".bc", vec1)]
```

# match any single character from among a list

enclose in brackets

"[fhr]" matches f, h, or r
combine with metacharacters for more specificity
"^ [fhr]" f, h, or r at beginning of a line vec2 <- c("fat", "bar", "rat", "elf", "mach", "hat") vec2[grep("^[fhr]", vec2)]
^ inside brackets (character class) is a "not" operator
"^ [^ fhr]" any single character at the beginning of a line except f, h, or r vec2[grep("^[^fhr]", vec2)]
any first character, followed by any character except a, and followed by any character one or more times

vec2[grep("^.[^a].+", vec2)]

### ranges and predefined character classes

```
[0-9] single digit from 0 to 9
[A-Z] single letter from A to Z ([a-z] lower case)
[0-9A-Za-z] any single alphanumeric character
[[:lower:]] - lower-case letters ([a-z])
[[:upper:]] - upper-case letters ([A-Z])
[[:alpha:]] - alphabetic characters ([A-Za-z])
[[:digit:]] - digits ([0-9])
[[:alnum:]] - alphanumeric ([A-Za-z0-9])
[[:punct:]] - punctuation: ! " # $ % & ' ( ) * + , - . /
              :; < = > ? @ [ \ ] ^ { | } ~
[[:graph:]] - graphical characters ([[:alnum:][:punct:]])
[[:space:]] - space characters: tab, newline, vertical tab,
```

form feed, carriage return, and space

### concatenation

combining single characters to match patterns

- find the words "fat", "hat", or "rat"
  vec3 <- c("fat", "bar", "rat", "fat boy", "elf", "mach", "hat")
  vec3[grep("^[fhr]at\$", vec3]]</pre>
- find words that start with "c" or "t" followed by "a" then followed by "b" or "r"

vec4 <- c("cab", "carat","tar","bar","tab","batboy","care")
vec4[grep("[ct]a[br]", vec4)]</pre>

- just three-letter words
   vec4[grep("^...\$", vec4)]
- any word with an "f' and a "t" separated by a single character vec5 <- c("fate", "rat", "fit", "bat", "futbol") vec5[grep("f.t", vec5)]

### repetition

specify times single character or match pattern repeated

- ? optional pattern, matched at most once, or not at all
- + matched one or more times
- \* any number (zero or more) times
- {n} exactly n times vec4[grep("^[[:alpha:]]{3}\$", vec4)]
- {n,} n or more times
- $\{n, m\}$  at least n times, but not more than m times
- match single, isolated words that start with "f" or "F", followed by one or more of any character, and ending with "t"

vec6 <- c("fat","fate","feat","bat","Fahrenheit","bat","foot")
vec6[grep("^[fF].+t\$", vec6)]</pre>

# alternation ("OR")

infix operator | to match from among two or more regular expressions

- ICD-10 hepatitis B codes: B16, B160, B161, B162, B163, B164, B165, B167, B168, B169, B17, B170, B172, B178, B18, B180, B181, B188, B189
- grep ICD10 hepatitis B deaths: dx1 <- c("B16", "B160", "B161", "B162", "B163", "B164", "B165", "B167", "B168", "B169", "B17", "B170", "B172", "B178", "B18", "B180", "B181", "B188", "B189")

grep("^B16[0-9]?\$|^B17[0,2,8]?\$|^B18[0,1,8,9]?\$", dx)

```
    create an indicator
    deathDat$hepB <- "No" #create new field set to no</li>
    get.hepB<-grep("^B16[0-9]?$|^B17[0,2,8]?$|^B18[0,1,8,9]?$",</li>
    deathDat$icd10)
    deatgDat$hepB[get.hepB] <- "Yes" # use index to assign yes</li>
```

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## other regular expression functions

explore on your own...

- regexpr() similar to grep, returns integer vectors with detailed information for the first occurrence of a pattern match within text string elements of a character vector
- gregexpr() similar to regexpr but returns a list with detailed information for the multiple occurrences of a pattern match within text string elements of a character vector
- sub() searches and replaces the first occurrence of a pattern match within text string elements of a character vector
- gsub() searches and replaces multiple occurrences of a pattern match within text string elements of a character vector