Introducción a R (II)

FSC

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Antes de empezar...

Los materiales de esta clase han sido preparados utilizando dos libros que están disponibles bajo licencia OUP y que os invito a explorar:

Hands-on programming with R https://rstudio-education.github.io/hopr/

Data Science with R https://rafalab.github.io/dsbook/

Paquetes en R

Ya hemos hablado acerca de una de las principales características de R: las funciones se construyen a partir del propio lenguaje y juntas conforman diferentes paquetes, que no son mas que conjuntos de funciones especificas para un cierto tema. Un paquete contiene código, documentación, ejemplos de uso y datasets.

Los paquetes de R pueden ser de uso personal o compartidos con la comunidad a traves fundamentalmente de 3 tipos de repositorios:

- CRAN: Es el repositorio oficial y consiste en una red de ftp y web servers mantenidos por la comunidad de R en todo el mundo. Para que un paquete pueda ser distribuido a través de CRAN necesita pasar una serie de requerimientos.
- Bioconductor: Es una colección de paquetes especificos de análisis de datos biomédicos y también mantenidos y testados por un grupo específico de expertos.
- Github: Aqui no hay control sobre los paquetes, como en los dos anteriores. Pero hoy en dia es una de las formas más comunes de compartir paquetes.

Instalar paquetes de CRAN

Para ello usamos el comando:

```
## Loading required package: sm
## Package 'sm', version 2.2-5.6: type help(sm) for summary information
## Loading required package: zoo
##
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
## as.Date, as.Date.numeric
```

Este es el paquete más antiguo de R. Suele ser útil usar la opción dependencies = T porque bajará automáticamente todos los paquetes de los que el paquete de interés depende.

Se pueden instalar varios paquetes a la vez usando la notación vectorial.

Instalar paquetes de Bioconductor

Instalar paquetes via devtools

El paquete devtools forma parte de CRAN, por lo tanto podemos instalarlo usando:

A partir de ahi se pueden instalar paquetes de distintas fuentes usando distintas funciones:

```
install_bioc() from Bioconductor,
install_bitbucket() from Bitbucket,
install_cran() from CRAN,
install_git() from a git repository,
install_github() from GitHub,
install_local() from a local file,
install_svn() from a SVN repository,
install_url() from a URL, and
install_version() from a specific version of a CRAN package.
```

Lo usaremos mucho durante el resto del curso porque algunos de los datasets que necesitamos vienen en paquetes contribuidos pero fuera de CRAN.

Entornos en R (environment)

Para seguir avanzando es importante que entendamos cómo R guarda los objetos. En realidad funciona de una forma parecida al sistema de ficheros de un ordenador. Cada objeto se guarda dentro de un environment

(o entorno) que a su vez puede estar dentro de otro, con una estructura jerárquica. Vamos a ver la estructura de nuestro entorno en esta sesion:

```
#install.packages("pryr")
library(pryr)
## Warning: package 'pryr' was built under R version 3.6.2
## Registered S3 method overwritten by 'pryr':
##
     method
                 from
##
     print.bytes Rcpp
##
## Attaching package: 'pryr'
##
  The following objects are masked from 'package:purrr':
##
##
       compose, partial
##
  The following object is masked from 'package:data.table':
##
       address
parenvs(all = TRUE)
##
      label
                                         name
## 1
      <environment: R GlobalEnv>
## 2 <environment: package:pryr>
                                         "package:pryr"
## 3 <environment: package:vioplot>
                                         "package:vioplot"
     <environment: package:zoo>
                                         "package:zoo"
## 4
      <environment: package:sm>
                                         "package:sm"
## 5
## 6
     <environment: package:rvest>
                                         "package:rvest"
     <environment: package:xml2>
                                         "package:xm12"
## 7
## 8 <environment: package:usethis>
                                         "package:usethis"
## 9 <environment: package:devtools>
                                         "package:devtools"
## 10 <environment: package:dslabs>
                                         "package:dslabs"
## 11 <environment: package:forcats>
                                         "package:forcats"
## 12 <environment: package:stringr>
                                         "package:stringr"
## 13 <environment: package:purrr>
                                         "package:purrr"
## 14 <environment: package:tibble>
                                         "package:tibble"
## 15 <environment: package:ggplot2>
                                         "package:ggplot2"
## 16 <environment: package:tidyverse>
                                         "package:tidyverse"
## 17 <environment: package:data.table>
                                         "package:data.table"
## 18 <environment: package:tidyr>
                                         "package:tidvr"
## 19 <environment: package:readr>
                                         "package:readr"
## 20 <environment: package:xlsx>
                                         "package:xlsx"
## 21 <environment: package:dplyr>
                                         "package:dplyr"
## 22 <environment: package:stats>
                                         "package:stats"
## 23 <environment: package:graphics>
                                         "package:graphics"
## 24 <environment: package:grDevices>
                                         "package:grDevices"
## 25 <environment: package:utils>
                                         "package:utils"
## 26 <environment: package:datasets>
                                         "package:datasets"
## 27 <environment: package:methods>
                                         "package:methods"
## 28 <environment: 0x000000013069828>
                                         "Autoloads"
## 29 <environment: base>
## 30 <environment: R_EmptyEnv>
```

```
x <- 5
where("x")
## <environment: R_GlobalEnv>
#> <environment: R_GlobalEnv>
where("mean")
## <environment: base>
R\_EmptyEnv es el único entorno que no tiene ningún padre. Los entornos de R no se almacenan físicamente
en tu ordenador, están en memoria RAM.
Para explorar la jerarquía de nuestro entorno tenemos algunas funciones:
as.environment("package:stats")
## <environment: package:stats>
## attr(,"name")
## [1] "package:stats"
## attr(,"path")
## [1] "C:/Users/fscabo/Documents/R/R-3.6.1/library/stats"
globalenv()
## <environment: R_GlobalEnv>
baseenv()
## <environment: base>
emptyenv()
## <environment: R_EmptyEnv>
parent.env(globalenv())
## <environment: package:pryr>
## attr(,"name")
## [1] "package:pryr"
## attr(,"path")
## [1] "C:/Users/fscabo/Documents/R/R-3.6.1/library/pryr"
search()
    [1] ".GlobalEnv"
                              "package:pryr"
                                                     "package:vioplot"
##
   [4] "package:zoo"
                              "package:sm"
                                                     "package:rvest"
## [7] "package:xml2"
                              "package:usethis"
                                                    "package:devtools"
## [10] "package:dslabs"
                              "package:forcats"
                                                     "package:stringr"
## [13] "package:purrr"
                              "package:tibble"
                                                    "package:ggplot2"
## [16] "package:tidyverse"
                              "package:data.table"
                                                    "package:tidyr"
## [19] "package:readr"
                              "package:xlsx"
                                                    "package:dplyr"
## [22] "package:stats"
                              "package:graphics"
                                                     "package:grDevices"
## [25] "package:utils"
                              "package:datasets"
                                                    "package:methods"
## [28] "Autoloads"
                              "package:base"
```

globalenv(), baseenv(), los environments en el search path, y emptyenv() se conectan como se muestra en la figura. Cada vez que se abre un nuevo paquete con library() se inserta entre el global environment y el ultimo entorno de paquete generado.

Los objetos guardados en un entorno pueden ser vistos con:

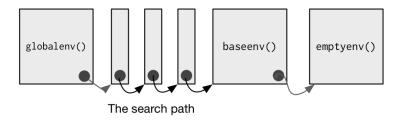


Figure 1: Jerarquía de entornos

```
ls(emptyenv())
## character(0)
ls(globalenv())
## [1] "x"
En cualquier momento podemos saber cual es el entorno con el que está trabajando R:
environment()
## <environment: R_GlobalEnv>
new<-"Hello Global"
Si guardamos algo en ese objeto new R lo sobreescribira:
new <- "Hello Active"
new
## [1] "Hello Active"
## "Hello Active"
Algunas funciones guardan objetos temporales.
roll <- function() {</pre>
  die <- 1:6
  dice <- sample(die, size = 2, replace = TRUE)</pre>
  sum(dice)
  return(environment())
}
```

Para no sobreescribir un posible objeto que se llame "dice", R crea un nuevo entorno cada vez que se evalua una funcion. De ahi las dependencias que hemos visto al principio. Al finalizar la evaluación, R vuelve al entorno global:

```
show_env <- function(){
  list(ran.in = environment(),
    parent = parent.env(environment()),
    objects = ls.str(environment()))
}
show_env()

## $ran.in
## <environment: 0x0000000182845d8>
##
```

```
## $parent
## <environment: R_GlobalEnv>
##
## $objects
```

R ha creado un nuevo runtime environment durante el tiempo que ha estado ejecutando la función.

Programando en R

Motivational example: Máquina Tragaperras

Para simular estos datos necesitamos hacer dos cosas:

1. Generar combinaciones de tres elementos de entre los siguientes símbolos: diamonds (DD), sevens (7), triple bars (BBB), double bars (BB), single bars (B), cherries (C), and zeroes (0).

Cada símbolo aparece según su probabilidad en la rueda.

2. Asignar un premio a cada combinación

Las máquinas tragaperras de la marca Manitoba tienen el siguiente esquema de premios:

Combination	Prize(\$)
DD DD DD	100
7 7 7	80
BBB BBB BBB	40
BB BB BB	25
В В В	10
C C C	10
Any combination of bars	5
C C *	5
C * C	5
* C C	5
C * *	2
* C *	2
* * C	2

Cada vez que jugamos a la maquina necesitamos pagar 1 dollar. Queremos crear un programa que nos de un score cada vez que jugamos. Podriamos hacer nuestro programa como un script o como una función

Estrategia general en programación

Hay tres recomendaciones que pueden ayudaros a hacer programas por muy complejos que sean:

- Romper el problema en subproblemas mas pequeños
- Escribir en lenguaje natural las instrucciones que se necesitarán para cada parte
- Usar ejemplos concretos

Un programa en R tiene dos tipos de tareas: paralelas y secuenciales

Instrucciones secuenciales

Un ejemplo de instrucciones secuenciales sería la siguiente función:

```
play <- function() {
    # step 1: generate symbols
    symbols <- get_symbols()</pre>
```



Figure 2: Esquema juego

```
# step 2: display the symbols
print(symbols)
# step 3: score the symbols
score(symbols)
}
```

Genera los simbolos al azar, los muestra y les da una puntuación de acuerdo con las reglas del juego.

Vamos a escribir una función para la primera tarea:

```
get_symbols <- function() {
    wheel <- c("DD", "7", "BBB", "BB", "C", "0")
    sample(wheel, size = 3, replace = TRUE,
        prob = c(0.03, 0.03, 0.06, 0.1, 0.25, 0.01, 0.52))
}
get_symbols()
## [1] "B" "DD" "B"
get_symbols()
## [1] "0" "DD" "0"
get_symbols()</pre>
```

Casos paralelos

Una vez que se eligen los tres símbolos hay que decidir como se puntua. Aqui tenemos que mirar varias cosas en paralelo como muestra la figura.

Cómo testamos cada una de esas opciones?

• Opción 1: ¿Son los tres iguales? Por ejemplo,

```
symbols<-c("7","7","7")
symbols[1] == symbols[2] & symbols[2] == symbols[3]

## [1] TRUE

## TRUE
symbols[1] == symbols[2] & symbols[1] == symbols[3]

## [1] TRUE</pre>
```

```
## TRUE
all(symbols == symbols[1])
## [1] TRUE
## TRUE
```

O equivalentemente:

```
length(unique(symbols) == 1)
```

[1] 1

Y tenemos la primera opción programada.

Nota: 😂 y // funcionan como sólo uno excepto porque no evaluan la segunda condicion si la primera es clara

• Opción 2: ¿tenemos todos símbolos de tipo B?

Es decir, entre nuestros símbolos tenemos alguna de estas combinaciones:

```
symbols <- c("B", "BBB", "BB")

symbols[1] == "B" | symbols[1] == "BBB" &
    symbols[2] == "B" | symbols[2] == "BBB" &
    symbols[3] == "B" | symbols[3] == "BBB" | symbols[3] == "BBB" |
## [1] TRUE
## TRUE</pre>
```

Pero no parece una solución muy elegante:

```
all(symbols %in% c("B", "BBB"))
```

```
## [1] TRUE
```

```
## TRUE
```

Hay overlapping entre el caso 1 y el 2: "B-B-B"

```
symbols <- c("B", "B", "B")
all(symbols %in% c("B", "BB", "BBB"))</pre>
```

```
## [1] TRUE
```

```
## TRUE
```

Nuestro programa ya contendría:

```
same <- symbols[1] == symbols[2] && symbols[2] == symbols[3]
bars <- symbols %in% c("B", "BBB")</pre>
```

• Opción 3: Si no se da ninguna de las anteriores, cuantas cerezas (C) tenemos?

```
count.c<-sum(symbols == "C")</pre>
```

if

Utilizamos el comando if para seleccionar situaciones que cumplen una determinada condicion. Si se cumple esto, haz aquello:

```
if (this) {
  that
}
```

Error in eval(expr, envir, enclos): object 'this' not found

this tiene que ser el resultado de un test lógico y resultar por tanto en TRUE o FALSE.

Si la condución se cumple (TRUE) se ejecutará el código entre corchetes. Si no se cumple (FALSE) no se hará nada.

Por ejemplo, si un número es negativo, multiplica por (-1) para hacerlo positivo:

```
num < -(-1)
num < -2
if (num < 0) {
  num <- num * -1
}
num <- -2
if (num < 0) {
  num <- num * -1
}
num
## [1] 2
## 2
num <- 4
if (num < 0) {
  num <- num * -1
}
num
## [1] 4
## 4
```

El resultado del test lógico tiene que ser un vector de una sola dimension. Si es un vector de varios TRUE/FALSE se evaluará sólo el primer elemento del vector.

Recuerda: las funciones any y all condensan las entradas de un vector lógico en un solo valor

Puedes incluir tantas lineas de código como quieras entre los corchetes:

```
num <- -1
if (num < 0) {
  print("num is negative.")
  print("Don't worry, I'll fix it.")
  num <- num * -1
  print("Now num is positive.")
}

## [1] "num is negative."
## [1] "Don't worry, I'll fix it."
## [1] "Now num is positive."

## "num is negative."
## "Don't worry, I'll fix it."
## "Now num is positive."
num</pre>
```

```
## [1] 1
## 1
```

Ejercicios:

Qué devuelven los siguientes códigos? (Intenta razonarlo y no ejecutarlo) Caso 1.

```
if (3 == 3) {
 x <- 2
}
х
## [1] 2
Caso 2.
x <- 1
if (TRUE) {
  x <- 2
}
х
## [1] 2
Case 3.
x <- 1
if (x == 1) {
 x <- 2
  if (x == 1) {
    x <- 3
  }
}
```

[1] 2

else

Hasta ahora si no se cumple la condicion impuesta, R no hace nada. Pero puede ser interesante que R haga una cosa distinta si se cumple que si no se cumple la condición. Para eso usamos else

```
if (this) {
  Plan A
} else {
  Plan B
}

## Error: <text>:2:8: unexpected symbol
## 1: if (this) {
## 2: Plan A
##
```

Vamos a intentar hacer un código que redondee un decimal al entero más cercano:

```
a <- 3.14
```

la función trunc nos da el entero más cercano:

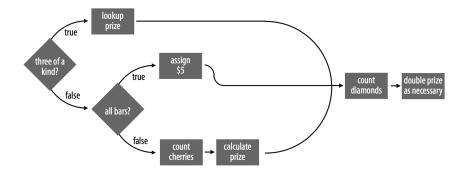


Figure 3: Esquema juego

```
dec <- a - trunc(a)
dec
## [1] 0.14
## 0.14
if (dec >= 0.5) {
  a \leftarrow trunc(a) + 1
} else {
  a <- trunc(a)
}
a
## [1] 3
## 3
Si hay más de dos casos, podemos usar else if:
a <- 1
b <- 1
if (a > b) {
  print("A wins!")
} else if (a < b) {</pre>
  print("B wins!")
} else {
  print("Tie.")
## [1] "Tie."
## "Tie."
```

Volviendo a la función que calcula el score para cada combinación de la máquina tragaperras, tenemos ocho posibles escenarios excluyentes, que se resumen en la figura 2.

Ya teniamos como testar cada una de las opciones y ahora sabemos como unirlas con if/else. Vamos a rellenar el cuerpo de la funcion score()

```
score <- function(symbols) {
    # calculate a prize
    prize</pre>
```

```
}
```

Donde calculate de prize va a ser:

```
score <- function (x) {</pre>
  # identify case
  same <- x[1] == x[2] & x[2] == x[3]
  bars <- x %in% c("B", "BB", "BBB")</pre>
  # get prize
  if (same & x[1]!=0) {
    payouts <- c("DD" = 100, "7" = 80, "BBB" = 40, "BB" = 25,
      "B" = 10, "C" = 10, "0" = 0)
    prize <- unname(payouts[x[1]])</pre>
  } else if (all(bars)) {
    prize <- 5
  } else {
    cherries <- sum(x == "C")</pre>
    prize \leftarrow c(0, 2, 5)[cherries + 1]
  }
  # adjust for diamonds
  diamonds <- sum(x == "DD")
 prize * 2 ^ diamonds
score(symbols)
```

[1] 10

Finalmente nuestro programa entero es asi:

```
play <- function() {
   symbols <- get_symbols()
   print(symbols)
   score(symbols)
}</pre>
```

Juguemos:

[1] 0

```
play()
## [1] "B" "DD" "O"
## [1] 0
play()
## [1] "0" "0" "B"
## [1] 0
play()
## [1] "7" "0" "0"
```

for and while loops

Vamos a intentar ver si estamos ante una tragaperras fraudulenta. Generamos un data frame que contenga todas las posibles combinaciones de valores que podemos obtener. Para eso usamos la función expand.grid

```
wheel <- c("DD", "7", "BBB", "B", "C", "0")
combos <- expand.grid(wheel, wheel, stringsAsFactors = FALSE)
View(combos)</pre>
```

Vamos a calcular la probabilidad de obtener cada una de estas combinaciones usando las probabilidades que hemos utilizado en get symbols

```
prob <- c("DD" = 0.03, "7" = 0.03, "BBB" = 0.06,
   "BB" = 0.1, "B" = 0.25, "C" = 0.01, "0" = 0.52)

combos$prob1 <- prob[combos$Var1]
combos$prob2 <- prob[combos$Var2]
combos$prob3 <- prob[combos$Var3]
View(combos)</pre>
```

Si las tiradas son independientes:

```
combos$prob <- combos$prob1 * combos$prob2 * combos$prob3
sum(combos$prob)</pre>
```

```
## [1] 1
View(combos)
```

Vamos a añadir una columna a nuestro data.frame combos que contenga el premio para cada combinación que hemos creado:

```
for (i in 1:nrow(combos)) {
   symbols <- c(combos[i, 1], combos[i, 2], combos[i, 3])
   combos*prize[i] <- score(symbols)
}
View(combos)</pre>
```

Ahora podemos calcular el premio esperado:

```
sum(combos$prize * combos$prob)
```

```
## [1] 0.538014
```

Por lo tanto esta máquina paga mas o menos el 54% de lo que juegas...

Importación de datos en R

En la clase anterior hemos leido archivos que estaban en datasets almacenados en librerias de R. Sin embargo, es comun tener la necesidad de leer datos a partir de diferentes tipos de archivos (.txt, .cvs, .xlsx) o incluso de otras fuentes como paginas web o bases de datos.

Importacion de datos desde archivos

Cuando se trata de importar datos a partir de un archivo en una unidad de disco a la que tenemos acceso en local o en remoto lo primero que tenemos que hacer es identificar dicha unidad y decirle a R cual es el lugar que queremos usar como directorio de trabajo working_directory. Es el lugar desde el que se leeran y en el que se escribira todo.

```
getwd()
```

[1] "C:/Users/fscabo/Desktop/MasterDataScience_KSchool/Session2_ImportExport_v2"

Para cambiar al $working_directory$ que queremos usar podemos utilizar la funcion setwd() o usar la GUI de RStudio en Session -> Set Working directory -> Choose Path

En esta sesion vamos a usar algunos de los datasets almacenados en el paquete dslabs de R.Irizarry en su libro Data Science Book. Vamos a copiar el archivo de datos el ejemplo murders en nuestro working_directory

```
dir <- system.file(package="dslabs") #extracts the location of package
dir</pre>
```

```
## [1] "C:/Users/fscabo/Documents/R/R-3.6.1/library/dslabs"
```

```
filename <- file.path(dir,"extdata/murders.csv")
file.copy(filename, "murders.csv")</pre>
```

```
## [1] FALSE
```

Ahora podeis comprobar que el archivo "murders.csv" esta en vuestro working directory.

```
list.files()
```

Funciones standard

Se trata de un archivo con extension .csv pequeño y facil de manejar. Hay varias funciones basicas que podemos usar para ello:

```
dat <- read.csv("DataSets/murders.csv")
head(dat)</pre>
```

```
##
          state abb region population total
## 1
                      South
                               4779736
                                          135
        Alabama AL
## 2
         Alaska
                 AK
                       West
                                710231
                                           19
## 3
                               6392017
                                          232
        Arizona AZ
                       West.
## 4
       Arkansas AR
                      South
                               2915918
                                           93
## 5 California CA
                       West
                              37253956
                                         1257
## 6
       Colorado CO
                       West
                               5029196
```

Alternativamente podriamos haber utilizado la funcion *read.delim* tambien del paquete base que es una de las mas flexibles pero también que requiere mas parametrizacion:

```
dat2 <- read.delim("DataSets/murders.csv",sep=",",header=T)
head(dat2)</pre>
```

```
##
          state abb region population total
## 1
                               4779736
                                          135
        Alabama AL
                      South
## 2
         Alaska AK
                       West
                                710231
                                           19
## 3
                               6392017
                                          232
        Arizona AZ
                       West
       Arkansas
                 AR
                      South
                               2915918
                                           93
## 5 California CA
                       West
                              37253956
                                        1257
       Colorado
                 CO
                       West
                               5029196
                                           65
```

Vamos a ver de que clase es el objeto dat

```
class(dat)
```

```
## [1] "data.frame"
```

Es una buena señal, lo ha leido como data.frame. Si recordais podemos acceder cada una de las columnas del data.frame con el símbolo \$

dat\$state

```
##
    [1] Alabama
                              Alaska
                                                    Arizona
    [4] Arkansas
##
                              California
                                                    Colorado
    [7] Connecticut
                              Delaware
                                                    District of Columbia
##
## [10] Florida
                              Georgia
                                                    Hawaii
## [13] Idaho
                              Illinois
                                                    Indiana
## [16] Iowa
                              Kansas
                                                    Kentucky
## [19] Louisiana
                              Maine
                                                    Maryland
## [22] Massachusetts
                              Michigan
                                                    Minnesota
## [25] Mississippi
                              Missouri
                                                    Montana
                                                    New Hampshire
## [28] Nebraska
                              Nevada
## [31] New Jersey
                              New Mexico
                                                    New York
## [34] North Carolina
                              North Dakota
                                                    Ohio
## [37] Oklahoma
                              Oregon
                                                    Pennsylvania
## [40] Rhode Island
                              South Carolina
                                                    South Dakota
## [43] Tennessee
                              Texas
                                                    Utah
## [46] Vermont
                              Virginia
                                                    Washington
## [49] West Virginia
                              Wisconsin
                                                    Wyoming
## 51 Levels: Alabama Alaska Arizona Arkansas California ... Wyoming
```

Sin embargo tenemos un problema porque la variable state se he volcado en R como factor y como ya hemos comentado anteriormente está desaconsejado utilizar los factores en R salvo en ocasiones puntuales dado que internamente para R por eficiencia los factores son realmente almacenados como numericos y transformaciones numerico -> string pueden ser complicadas. Por ello seria mucho mejor no leer los strings como factores. R tiene una opcion para ello:

```
dat <- read.csv("DataSets/murders.csv", stringsAsFactors = FALSE)
class(dat$state)</pre>
```

[1] "character"

Utilizamos una de las funciones fundamentales del paquete dplyr para volver a poner la variable state como un factor por medio de:

```
dat <- mutate(dat, region = as.factor(region))</pre>
```

Abrid el archivo murders.csv y volvedlo a guardar como archivo .xlsx. Vamos a leerlo utilizando el paquete library(xlsx)

```
dat2 <- read.xlsx(file = "DataSets/murders.xlsx", sheetIndex = 1)
class(dat2)</pre>
```

[1] "data.frame"

```
head(dat2)
```

```
##
          state abb region population total
## 1
        Alabama AL
                      South
                               4779736
                                          135
## 2
         Alaska
                       West
                                710231
                                           19
                 AK
## 3
                               6392017
                                          232
        Arizona
                 ΑZ
                       West
       Arkansas
                               2915918
                 AR
                      South
                                           93
## 5 California CA
                       West
                               37253956
                                         1257
       Colorado
                 CO
                       West
                               5029196
                                           65
```

class(dat2\$state)

```
## [1] "factor"
```

```
identical(dat$state,dat2$state)

## [1] FALSE
identical(as.matrix(dat$state),as.matrix(dat2$state))

## [1] TRUE

Other function to read tabulated data into R is read.table:

dat3=read.table(file = "DataSets/murders.csv",header = T,sep=",")
head(dat3)

## state abb region population total

## 1 Alabama AL South 4779736 135
```

```
Alabama AL South
## 1
                             4779736
                                       135
## 2
         Alaska AK
                     West
                              710231
                                        19
                                       232
## 3
       Arizona AZ
                     West
                             6392017
      Arkansas AR South
                             2915918
                                        93
                            37253956
                                      1257
## 5 California CA
                     West
      Colorado CO
                             5029196
                     West
```

NOTA IMPORTANTE: ASCII es un sistema de codificacion que convierte los caracteres en números. ASCII usa 7 bits (0000001) por lo que se pueden generar un total de 128 símbolos. Para lenguages con un gran número de símbolos se puede utilizar otra codificación llamada UTF para la que se pueden escoger combinaciones de 8, 16 o 24 bits. RStudio utiliza UTF-8. NOTA IMPORTANTE 2: Todos los archivos que hemos leido hasta ahora son archivos de texto; i.e. se pueden leer con un editor cualquiera. Archivos xls, html, json son binarios y necesitaremos otro tipo de herramientas para leerlos.

Cómo abrir grandes archivos

El paquete data.table contiene funciones para leer archivos de varios megas de RAM. En particular fread() funciona muy bien y se le pueden ajustar los parámetros: sep, colClasses and nrows. bit64::integer64 types también se detectan y leer automáticamente sin necesidad de transformarlos antes en caracter.También es interesante mostrar el progreso en la lectura del archivo para tener una idea de la duracion de la lectura.Otro parámetro que hereda de las funciones básicas de lectura es stringsAsFactors.

```
counts.rnaseq=fread("DataSets/Counts.genes.DiffAll.genes.limma.random.txt",sep = "\t",showProgress=T)
#head(counts.rnaseq)
class(counts.rnaseq)
```

[1] "data.table" "data.frame"

Esta función devuelve un objeto data.table en lugar de un data.frame a no ser que se le especifique: data.table = FALSE. Ejemplo de tiempos (del manual de data.table):

```
write.table(DT,
            "DataSets/test.csv",
            sep=",",
            row.names=FALSE,
            quote=FALSE)
cat("File size (MB):",
    round(file.info("test.csv")$size/1024^2),"\n")
## File size (MB): NA
system.time(DF1 <-read.csv("DataSets/test.csv",stringsAsFactors=FALSE))</pre>
##
      user system elapsed
      4.30
##
              0.19
                       4.53
system.time(DF2 <- read.table("DataSets/test.csv",header=TRUE,sep=",",quote="",</pre>
    stringsAsFactors=FALSE,comment.char="",nrows=n,
    colClasses=c("integer","integer","numeric",
                  "character", "numeric", "integer")))
##
      user system elapsed
##
      1.21
              0.07
                       1.28
system.time(DT <- fread("DataSets/test.csv"))</pre>
##
      user system elapsed
              0.06
##
      0.10
                       0.06
  #require(sqldf)
  #require(ff)
identical(DF1,DF2)
## [1] TRUE
all.equal(as.data.table(DF1), DT)
## [1] TRUE
```

tidy verse

Ya hemos visto como el paquete xlsx (y otros) contenian funciones para leer archivos excel en R. El paquete tidyverse también puede utilizarse para ello a traves de las funciones:

Function	Separador	Sufijo
read_table	espacio	txt
$read_csv$,	csv
${\rm read_csv2}$;	csv
$read_tsv$	tab	tsv
${\rm read_delim}$	general	txt

The readxl package provides functions to read-in Microsoft Excel formats:

Function	Format	Typical suffix
read_xls	auto detect the format original format new format	xls, xlsx xls xlsx

Si os fijáis todas estas funciones son parecidas a las que teníamos en el paquete base separadas por un punto como read.csv. Comparemos ambas funciones:

```
library(tidyverse)
dat=read.csv("DataSets/murders.csv")
class(dat)
## [1] "data.frame"
head(dat)
##
          state abb region population total
                               4779736
## 1
        Alabama
                 AL
                      South
                                          135
## 2
         Alaska
                 AK
                       West
                                710231
                                           19
## 3
                       West
                               6392017
                                          232
        Arizona
                 ΑZ
## 4
       Arkansas
                 AR
                      South
                               2915918
                                           93
## 5 California CA
                              37253956
                                         1257
                       West
       Colorado
                 CO
                       West
                               5029196
                                           65
dat2=read_csv("DataSets/murders.csv")
class(dat2)
## [1] "spec_tbl_df" "tbl_df"
                                                    "data.frame"
                                     "tbl"
head(dat2)
## # A tibble: 6 x 5
##
     state
                abb
                       region population total
##
     <chr>
                <chr> <chr>
                                    <dbl> <dbl>
## 1 Alabama
                 AL
                       South
                                  4779736
                                            135
## 2 Alaska
                 AK
                       West
                                  710231
                                             19
## 3 Arizona
                 ΑZ
                       West
                                  6392017
                                            232
## 4 Arkansas
                 AR
                       South
                                  2915918
                                             93
## 5 California CA
                                 37253956
                       West
                                           1257
## 6 Colorado
                       West
                                 5029196
                                             65
```

OMG: What's that? *Tibbles* son nuevos tipos de data.frames que contienen muchas de las buenas prácticas que hemos ido mencionando. Por ejemplo:

• Tibble nunca cambia el tipo de una variable (si es caracter, es caracter, no factor...)

```
tibble(x = letters)
## # A tibble: 26 x 1
##
      x
##
      <chr>>
##
    1 a
##
    2 b
##
    3 c
##
    4 d
##
    5 e
##
    6 f
   7 g
##
```

```
##
    8 h
## 9 i
## 10 j
## # ... with 16 more rows
tibble(x = 1:3, y = list(1:5, 1:10, 1:20))
## # A tibble: 3 x 2
##
         х у
##
     <int> <list>
## 1
          1 <int [5]>
## 2
         2 <int [10]>
## 3
          3 <int [20]>
   • Tibble nunca cambia el nombre de las variables
names(tibble("Old Alphabet" = letters))
## [1] "Old Alphabet"
names(data.frame("Old Alphabet" = letters))
## [1] "Old.Alphabet"
   • Evalua los argumentos secuencialmente
#data.frame(z=1:5, y=z ^2)
tibble(x = 1:5, y = x^2)
## # A tibble: 5 x 2
##
         Х
##
     <int> <dbl>
## 1
          1
                1
          2
## 2
                4
## 3
          3
                9
## 4
          4
               16
  • El objetivo fundamental de tibble es almacenar las variables de una forma consistente, por lo que no
     hace uso de row.names
   • Printing: Por defecto solo se escriben en pantalla las 10 primeras filas y todas las columnas que quepan.
     Se puede cambiar el tipo de letra, color, etc. tibble.print_max = Inf muestra todas las filas
tibble(x = 1:5, y = x^2 2)
## # A tibble: 5 x 2
##
         х
                У
     <int> <dbl>
##
## 1
          1
                1
## 2
         2
                4
## 3
          3
                9
          4
               16
## 4
          5
               25
data.frame(x=1:5, y=(1:5)^2)
     х
        У
## 1 1
        1
```

2 2 4

```
## 3 3 9
## 4 4 16
## 5 5 25
```

• Subsetting siempre devuelve otro objeto tibble, sea de las dimensiones que sea

```
df1 \leftarrow data.frame(x = 1:3, y = 3:1)
class(df1[, 1:2])
## [1] "data.frame"
class(df1[, 1])
## [1] "integer"
df2 \leftarrow tibble(x = 1:3, y = 3:1)
class(df2[, 1:2])
## [1] "tbl_df"
                      "tbl"
                                     "data.frame"
class(df2[, 1])
## [1] "tbl_df"
                      "tbl"
                                     "data.frame"
Si una columna no tiene exactamente el nombre por el que se la llama no se hace partial matching:
df <- data.frame(abc = 1)</pre>
df$a
## [1] 1
df2 <- tibble(abc = 1)
## Warning: Unknown or uninitialised column: 'a'.
## NULL
```

Tidy data

La organización Gapminder trata de desenmascarar falsos mitos acerca del estado del mundo en terminos de pobreza, desigualdad, etc a través del uso de datos. Utilizando el dataset *gapminder* del paquete *dslabs* trataremos de contestar a dos preguntas: * Es cierto que el mundo se divide en paises occidentales ricos y no-occidentales pobres? * Han aumentado las diferencias entre paises en los últimos 40 años?

1. Descargamos los datos:

```
library(dslabs)
data(gapminder)
#si no tienes el paquete instalado
#gapminder=read.csv("DataSets/Gapminder.csv")
```

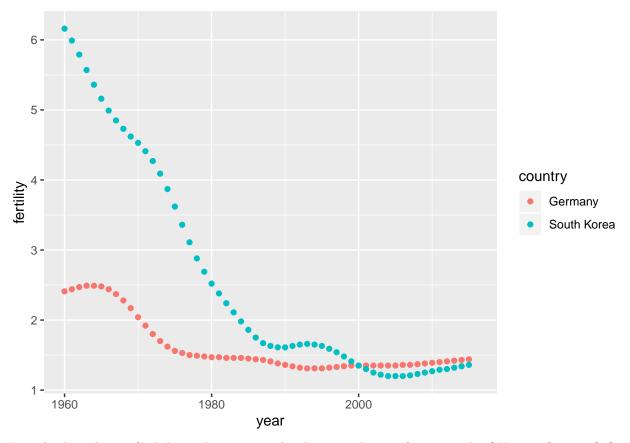
2. Exploramos los datos:

```
View(gapminder)
head(gapminder)
```

```
country year infant_mortality life_expectancy fertility
##
## 1
                 Albania 1960
                                         115.40
                                                           62.87
                                                                       6.19
## 2
                 Algeria 1960
                                         148.20
                                                           47.50
                                                                       7.65
## 3
                  Angola 1960
                                         208.00
                                                           35.98
                                                                      7.32
## 4 Antigua and Barbuda 1960
                                             NA
                                                           62.97
                                                                      4.43
## 5
               Argentina 1960
                                          59.87
                                                           65.39
                                                                      3.11
```

```
## 6
                Armenia 1960
                                                       66.86
                                                                  4.55
##
    population
                        gdp continent
                                              region
## 1
                               Europe Southern Europe
       1636054
                        NA
## 2
      11124892 13828152297
                               Africa Northern Africa
## 3
       5270844
                         NA
                               Africa
                                      Middle Africa
## 4
         54681
                         NA Americas
                                           Caribbean
## 5
      20619075 108322326649 Americas
                                      South America
       1867396
                                        Western Asia
## 6
                         NΑ
                                 Asia
str(gapminder)
## 'data.frame':
                   10545 obs. of 9 variables:
## $ country
                     : Factor w/ 185 levels "Albania", "Algeria", ...: 1 2 3 4 5 6 7 8 9 10 ...
## $ year
                     $ infant_mortality: num 115.4 148.2 208 NA 59.9 ...
## $ life_expectancy : num 62.9 47.5 36 63 65.4 ...
## $ fertility
                    : num 6.19 7.65 7.32 4.43 3.11 4.55 4.82 3.45 2.7 5.57 ...
## $ population
                     : num 1636054 11124892 5270844 54681 20619075 ...
## $ gdp
                     : num NA 1.38e+10 NA NA 1.08e+11 ...
                    : Factor w/ 5 levels "Africa", "Americas", ...: 4 1 1 2 2 3 2 5 4 3 ...
## $ continent
                     : Factor w/ 22 levels "Australia and New Zealand",..: 19 11 10 2 15 21 2 1 22 21
## $ region
  3. Para trabajar con el universo tidyverse necesitamos que los datos esten en formato "tidy"
tidy_data <- gapminder %>%
  filter(country %in% c("South Korea", "Germany")) %>%
  select(country, year, fertility)
head(tidy_data)
##
        country year fertility
        Germany 1960
## 1
                          2.41
## 2 South Korea 1960
                          6.16
## 3
        Germany 1961
                          2.44
## 4 South Korea 1961
                          5.99
        Germany 1962
                          2.47
## 6 South Korea 1962
                          5.79
tidy_data %>%
  ggplot(aes(year, fertility, color = country)) +
 geom_point()
```

Warning: Removed 2 rows containing missing values (geom_point).



Este plot ha sido tan fácil de producir porque los datos estaban en formato *tidy*. QUe significa esto? Que cada medida especifica de pais & fecha ocupa una fila de un *tibble* o *data.frame*. Las columnas serian cada una de las observaciones que queramos mirar (fertilidad, esperanza de vida...)

```
path <- system.file("extdata", package="dslabs")</pre>
filename <- file.path(path, "fertility-two-countries-example.csv")</pre>
wide_data <- read_csv(filename)</pre>
## Parsed with column specification:
## cols(
     .default = col_double(),
     country = col_character()
##
## )
## See spec(...) for full column specifications.
wide_data<-read_csv("DataSets/fertility-two-countries-example.csv")</pre>
## Parsed with column specification:
## cols(
##
     .default = col_double(),
##
     country = col_character()
## )
## See spec(...) for full column specifications.
wide_data
## # A tibble: 2 x 57
     country `1960` `1961` `1962` `1963` `1964` `1965` `1966` `1967` `1968`
```

```
<dbl> <
## 1 Germany
               2.41
                      2.44
                             2.47
                                    2.49
                                           2.49
                                                  2.48
                                                         2.44
                                                                 2.37
                                                                        2.28
## 2 South ~
               6.16
                      5.99
                             5.79
                                    5.57
                                           5.36
                                                  5.16
                                                         4.99
     ... with 47 more variables: `1969` <dbl>, `1970` <dbl>, `1971` <dbl>,
       `1972` <dbl>, `1973` <dbl>, `1974` <dbl>, `1975` <dbl>, `1976` <dbl>,
## #
      `1977` <dbl>, `1978` <dbl>, `1979` <dbl>, `1980` <dbl>, `1981` <dbl>,
      `1982` <dbl>, `1983` <dbl>, `1984` <dbl>, `1985` <dbl>, `1986` <dbl>,
       `1987` <dbl>, `1988` <dbl>, `1989` <dbl>, `1990` <dbl>, `1991` <dbl>,
## #
## #
       `1992` <dbl>, `1993` <dbl>, `1994` <dbl>, `1995` <dbl>, `1996` <dbl>,
## #
       `1997` <dbl>, `1998` <dbl>, `1999` <dbl>, `2000` <dbl>, `2001` <dbl>,
       `2002` <dbl>, `2003` <dbl>, `2004` <dbl>, `2005` <dbl>, `2006` <dbl>,
       `2007` <dbl>, `2008` <dbl>, `2009` <dbl>, `2010` <dbl>, `2011` <dbl>,
## #
       `2012` <dbl>, `2013` <dbl>, `2014` <dbl>, `2015` <dbl>
#seleccionamos las primeras 9 columnas
select(wide_data, country, `1960`: `1967`)
## # A tibble: 2 x 9
##
     country
                 `1960` `1961` `1962` `1963` `1964` `1965` `1966`
                                                                   1967
##
     <chr>
                  <dbl> <dbl> <dbl>
                                       <dbl>
                                              <dbl> <dbl>
                                                             <dbl>
                                                                    <dbl>
## 1 Germany
                   2.41
                          2.44
                                 2.47
                                        2.49
                                               2.49
                                                      2.48
                                                              2.44
                                                                     2.37
## 2 South Korea
                   6.16
                          5.99
                                 5.79
                                               5.36
                                                      5.16
                                                              4.99
                                        5.57
                                                                     4.85
  5. Convertir wide_data en tidy_data
new tidy data <- wide data %>%
  pivot_longer(cols=`1960`:`2015`,names_to="year",values_to = "fertility")
new_tidy_data
## # A tibble: 112 x 3
##
      country year fertility
##
      <chr>
              <chr>
                        <dbl>
## 1 Germany 1960
                         2.41
##
   2 Germany 1961
                         2.44
## 3 Germany 1962
                         2.47
## 4 Germany 1963
                         2.49
## 5 Germany 1964
                         2.49
## 6 Germany 1965
                         2.48
## 7 Germany 1966
                         2.44
## 8 Germany 1967
                         2.37
## 9 Germany 1968
                         2.28
## 10 Germany 1969
                         2.17
## # ... with 102 more rows
class(new_tidy_data$year)
## [1] "character"
new_tidy_data <- wide_data %>%
  pivot longer(cols=`1960`: 2015`,
               names_to="year",
               values_to = "fertility",
               names_ptypes = list(year=integer()))
class(new_tidy_data$year)
```

[1] "integer"

A veces se necesita volver de tidy data a wide data:

```
new_wide_data <- new_tidy_data %>%
  pivot_wider(names_from=year, values_from = fertility)
new_wide_data
## # A tibble: 2 x 57
##
     country `1960` `1961` `1962` `1963` `1964` `1965` `1966` `1967` `1968`
##
              <dbl>
                     <dbl>
                            <dbl>
                                    <dbl>
                                           <dbl>
                                                  <dbl>
                                                          <dbl>
                                                                 <dbl>
## 1 Germany
               2.41
                      2.44
                             2.47
                                     2.49
                                            2.49
                                                   2.48
                                                          2.44
                                                                  2.37
## 2 South ~
               6.16
                      5.99
                                            5.36
                                                   5.16
                                                          4.99
                                                                  4.85
                             5.79
                                     5.57
     ... with 47 more variables: `1969` <dbl>, `1970` <dbl>, `1971` <dbl>,
       `1972` <dbl>, `1973` <dbl>, `1974` <dbl>, `1975` <dbl>, `1976` <dbl>,
## #
       `1977` <dbl>, `1978` <dbl>, `1979` <dbl>, `1980` <dbl>, `1981` <dbl>,
## #
       `1982` <dbl>, `1983` <dbl>, `1984` <dbl>, `1985` <dbl>, `1986`
       `1987` <dbl>, `1988` <dbl>, `1989` <dbl>, `1990` <dbl>, `1991`
## #
       `1992` <dbl>, `1993` <dbl>, `1994` <dbl>, `1995` <dbl>, `1996` <dbl>,
```

`1997` <dbl>, `1998` <dbl>, `1999` <dbl>, `2000` <dbl>, `2001` <dbl>,

'2002' <dbl>, '2003' <dbl>, '2004' <dbl>, '2005' <dbl>, '2006' <dbl>, '2007' <dbl>, '2008' <dbl>, '2009' <dbl>, '2010' <dbl>, '2011' <dbl>,

`2012` <dbl>, `2013` <dbl>, `2014` <dbl>, `2015` <dbl>

Importando datos de bases de datos

Datos estructurados: RMySQL -> RMariaDB

MariaDB es un servidor de bases de datos. Podemos interaccionar con el usando un cliente o bien programaticamente usando APIs (Application Programming Interface). Si interaccionamos a traves de un cliente como mysql empezariamos conectandonos a la base de datos con el comando:

mysql -u root -p -h localhost

#

#

#

Existe un paquete de R para hacer lo mismo y traernos datos:

```
library(RMariaDB)
```

```
## Warning: package 'RMariaDB' was built under R version 3.6.2
```

En primer lugar creamos una objeto de conexion a la base de datos y podemos ver todas las bases de datos que hay dentro de ella:

```
##
                                                    Database
## 1
                                          information schema
## 2
                                              PERCONA SCHEMA
## 3
                      acanthochromis_polyacanthus_core_94_1
## 4
                      acanthochromis_polyacanthus_core_95_1
## 5
                      acanthochromis_polyacanthus_core_96_1
## 6
                      acanthochromis_polyacanthus_core_97_1
## 7
                      acanthochromis_polyacanthus_core_98_1
## 8
                      acanthochromis_polyacanthus_core_99_1
## 9
             acanthochromis_polyacanthus_otherfeatures_94_1
## 10
             acanthochromis_polyacanthus_otherfeatures_95_1
```

```
## 11
             acanthochromis polyacanthus otherfeatures 96 1
##
  12
             acanthochromis_polyacanthus_otherfeatures_97_1
##
  13
             acanthochromis polyacanthus otherfeatures 98 1
             acanthochromis_polyacanthus_otherfeatures_99_1
##
  14
##
  15
                    acanthochromis polyacanthus rnaseq 94 1
  16
                    acanthochromis polyacanthus rnaseq 95 1
##
                    acanthochromis polyacanthus rnaseq 96 1
  17
                    acanthochromis_polyacanthus_rnaseq_97_1
## 18
                    acanthochromis_polyacanthus_rnaseq_98_1
##
  19
                    acanthochromis_polyacanthus_rnaseq_99_1
##
  20
##
  21
                                   accipiter_nisus_core_99_1
                                 accipiter_nisus_rnaseq_99_1
##
  22
##
  23
                                    aedes_aegypti_core_48_1b
## 24
                                    aedes_aegypti_core_49_1b
## 25
                                    aedes_aegypti_core_50_1c
## 26
                                    aedes_aegypti_core_51_1c
## 27
                                    aedes_aegypti_core_52_1d
##
  28
                                    aedes aegypti core 53 1d
                                    aedes_aegypti_core_54_1d
##
  29
##
  30
                                    aedes aegypti core 55 1d
##
  31
                            ailuropoda_melanoleuca_core_60_1
## 32
                           ailuropoda melanoleuca core 61 1a
## 33
                           ailuropoda_melanoleuca_core_62_1b
  34
                            ailuropoda melanoleuca core 63 1
##
                            ailuropoda melanoleuca core 64 1
## 35
  36
                            ailuropoda melanoleuca core 65 1
##
  37
                            ailuropoda_melanoleuca_core_66_1
                            ailuropoda_melanoleuca_core_67_1
##
  38
##
  39
                            ailuropoda_melanoleuca_core_68_1
                            ailuropoda melanoleuca core 69 1
## 40
## 41
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                      caenorhabditis_elegans_funcgen_70_230
## 854
                      caenorhabditis_elegans_funcgen_71_235
## 855
                      caenorhabditis elegans funcgen 72 235
                      caenorhabditis_elegans_funcgen_73_235
## 856
## 857
                      caenorhabditis elegans funcgen 74 235
## 858
                      caenorhabditis_elegans_funcgen_75_240
## 859
                      caenorhabditis elegans funcgen 76 240
## 860
                      caenorhabditis_elegans_funcgen_77_240
## 861
                      caenorhabditis elegans funcgen 78 245
## 862
                      caenorhabditis elegans funcgen 79 245
                      caenorhabditis elegans funcgen 80 245
## 863
## 864
                      caenorhabditis_elegans_funcgen_81_245
## 865
                      caenorhabditis_elegans_funcgen_82_245
## 866
                      caenorhabditis_elegans_funcgen_83_245
## 867
                      caenorhabditis_elegans_funcgen_84_245
## 868
                       caenorhabditis_elegans_funcgen_85_250
## 869
                      caenorhabditis_elegans_funcgen_86_250
## 870
                      caenorhabditis_elegans_funcgen_87_250
## 871
                      caenorhabditis_elegans_funcgen_88_250
## 872
                      caenorhabditis_elegans_funcgen_89_250
                      caenorhabditis_elegans_funcgen_90_250
## 873
## 874
                      caenorhabditis elegans funcgen 91 260
```

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## 875
                      caenorhabditis elegans funcgen 92 260
## 876
                       caenorhabditis_elegans_funcgen_93_260
## 877
                       caenorhabditis elegans funcgen 94 260
## 878
                       caenorhabditis_elegans_funcgen_95_260
## 879
                       caenorhabditis elegans funcgen 96 260
## 880
                      caenorhabditis elegans funcgen 97 269
## 881
                      caenorhabditis elegans funcgen 98 269
## 882
                      caenorhabditis elegans funcgen 99 269
                                   calidris_pugnax_core_96_1
## 883
## 884
                                   calidris_pugnax_core_97_1
## 885
                                   calidris_pugnax_core_98_1
## 886
                                   calidris_pugnax_core_99_1
## 887
                          calidris_pugnax_otherfeatures_96_1
## 888
                          calidris_pugnax_otherfeatures_97_1
## 889
                          calidris_pugnax_otherfeatures_98_1
## 890
                          calidris_pugnax_otherfeatures_99_1
## 891
                                 calidris_pugnax_rnaseq_96_1
## 892
                                 calidris pugnax rnaseg 97 1
## 893
                                 calidris_pugnax_rnaseq_98_1
## 894
                                 calidris_pugnax_rnaseq_99_1
## 895
                                  calidris_pygmaea_core_96_1
## 896
                                  calidris pygmaea core 97 1
## 897
                                  calidris_pygmaea_core_98_1
## 898
                                  calidris pygmaea core 99 1
## 899
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## 900
                                calidris_pygmaea_rnaseq_97_1
## 901
                                calidris_pygmaea_rnaseq_98_1
## 902
                                calidris_pygmaea_rnaseq_99_1
## 903
                                callithrix_jacchus_core_56_3
## 904
                                callithrix_jacchus_core_57_3
## 905
                              callithrix_jacchus_core_58_321
## 906
                             callithrix_jacchus_core_59_321a
## 907
                             callithrix_jacchus_core_60_321b
## 908
                             callithrix_jacchus_core_61_321c
## 909
                             callithrix jacchus core 62 321d
## 910
                              callithrix_jacchus_core_63_321
## 911
                              callithrix jacchus core 64 321
## 912
                              callithrix_jacchus_core_65_321
## 913
                              callithrix jacchus core 66 321
## 914
                              callithrix_jacchus_core_67_321
## 915
                              callithrix jacchus core 68 321
## 916
                              callithrix jacchus core 69 321
                              callithrix jacchus core 70 321
## 917
## 918
                              callithrix_jacchus_core_71_321
## 919
                              callithrix_jacchus_core_72_321
## 920
                              callithrix_jacchus_core_73_321
## 921
                              callithrix_jacchus_core_74_321
## 922
                              callithrix_jacchus_core_75_321
## 923
                              callithrix_jacchus_core_76_321
                              callithrix_jacchus_core_77_321
## 924
## 925
                              callithrix_jacchus_core_78_321
## 926
                              callithrix_jacchus_core_79_321
## 927
                              callithrix_jacchus_core_80_321
## 928
                              callithrix_jacchus_core_81_321
```

```
## 929
                              callithrix jacchus core 82 321
## 930
                              callithrix_jacchus_core_83_321
## 931
                              callithrix jacchus core 84 321
## 932
                              callithrix_jacchus_core_85_321
## 933
                              callithrix_jacchus_core_86_321
## 934
                              callithrix jacchus core 87 321
## 935
                              callithrix jacchus core 88 321
## 936
                              callithrix jacchus core 89 321
## 937
                              callithrix_jacchus_core_90_321
## 938
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## 939
                                callithrix_jacchus_core_92_1
## 940
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## 941
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                                callithrix_jacchus_core_95_1
## 942
## 943
                                callithrix_jacchus_core_96_1
## 944
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## 945
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## 946
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## 947
                             callithrix_jacchus_funcgen_92_1
## 948
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## 949
                             callithrix_jacchus_funcgen_94_1
## 950
                             callithrix jacchus funcgen 95 1
## 951
                             callithrix_jacchus_funcgen_96_1
## 952
                             callithrix jacchus funcgen 97 1
## 953
                             callithrix jacchus funcgen 98 1
## 954
                             callithrix jacchus funcgen 99 1
## 955
                      callithrix_jacchus_otherfeatures_56_3
## 956
                      callithrix_jacchus_otherfeatures_57_3
## 957
                    callithrix_jacchus_otherfeatures_58_321
## 958
                   callithrix_jacchus_otherfeatures_59_321a
## 959
                   callithrix_jacchus_otherfeatures_60_321b
## 960
                   callithrix_jacchus_otherfeatures_61_321c
## 961
                   callithrix_jacchus_otherfeatures_62_321d
## 962
                    callithrix_jacchus_otherfeatures_63_321
## 963
                    callithrix jacchus otherfeatures 64 321
                    callithrix_jacchus_otherfeatures_65_321
## 964
## 965
                    callithrix jacchus otherfeatures 66 321
## 966
                    callithrix_jacchus_otherfeatures_67_321
## 967
                    callithrix jacchus otherfeatures 68 321
## 968
                    callithrix_jacchus_otherfeatures_69_321
## 969
                    callithrix jacchus otherfeatures 70 321
## 970
                    callithrix jacchus otherfeatures 71 321
                    callithrix jacchus otherfeatures 72 321
## 971
                    callithrix_jacchus_otherfeatures_73_321
## 972
## 973
                    callithrix_jacchus_otherfeatures_74_321
## 974
                    callithrix_jacchus_otherfeatures_75_321
## 975
                    callithrix_jacchus_otherfeatures_76_321
## 976
                    callithrix_jacchus_otherfeatures_77_321
## 977
                    callithrix_jacchus_otherfeatures_78_321
## 978
                    callithrix_jacchus_otherfeatures_79_321
                    callithrix_jacchus_otherfeatures_80 321
## 979
## 980
                    callithrix_jacchus_otherfeatures_81_321
## 981
                    callithrix_jacchus_otherfeatures_82_321
## 982
                    callithrix jacchus otherfeatures 83 321
```

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## 983
                    callithrix jacchus otherfeatures 84 321
## 984
                    callithrix_jacchus_otherfeatures_85_321
## 985
                    callithrix jacchus otherfeatures 86 321
## 986
                    callithrix_jacchus_otherfeatures_87_321
## 987
                    callithrix_jacchus_otherfeatures_88_321
## 988
                    callithrix jacchus otherfeatures 89 321
## 989
                    callithrix jacchus otherfeatures 90 321
                    callithrix jacchus otherfeatures 91 321
## 990
## 991
                              callithrix jacchus rnaseg 92 1
## 992
                              callithrix_jacchus_rnaseq_93_1
## 993
                              callithrix_jacchus_rnaseq_94_1
## 994
                              callithrix_jacchus_rnaseq_95_1
## 995
                              callithrix_jacchus_rnaseq_96_1
## 996
                              callithrix_jacchus_rnaseq_97_1
## 997
                              callithrix_jacchus_rnaseq_98_1
## 998
                              callithrix_jacchus_rnaseq_99_1
                             callorhinchus_milii_core_97_613
## 999
## 1000
                             callorhinchus milii core 98 613
## 1001
                             callorhinchus_milii_core_99_613
## 1002
                    callorhinchus milii otherfeatures 97 613
## 1003
                    callorhinchus_milii_otherfeatures_98_613
## 1004
                    callorhinchus milii otherfeatures 99 613
## 1005
                           callorhinchus_milii_rnaseq_97_613
## 1006
                           callorhinchus milii rnaseq 98 613
## 1007
                           callorhinchus_milii_rnaseq_99_613
## 1008
                               camelus dromedarius core 99 2
## 1009
                             camelus_dromedarius_rnaseq_99_2
## 1010
                                 canis_familiaris_core_48_2f
## 1011
                                 canis_familiaris_core_49_2g
## 1012
                                 canis_familiaris_core_50_2h
## 1013
                                 canis_familiaris_core_51_2i
## 1014
                                 canis_familiaris_core_52_2j
## 1015
                                 canis_familiaris_core_53_2k
## 1016
                                 canis_familiaris_core_54_21
## 1017
                                 canis familiaris core 55 2m
## 1018
                                 canis_familiaris_core_56_2m
## 1019
                                 canis familiaris core 57 2m
## 1020
                                 canis_familiaris_core_58_2n
## 1021
                                 canis familiaris core 59 20
## 1022
                                 canis_familiaris_core_60_2p
## 1023
                                 canis familiaris core 61 2q
## 1024
                                 canis familiaris core 62 2r
## 1025
                                  canis familiaris core 63 2
## 1026
                                  canis_familiaris_core_64_2
## 1027
                                  canis_familiaris_core_65_2
## 1028
                                  canis_familiaris_core_66_2
## 1029
                                  canis_familiaris_core_67_2
## 1030
                                 canis_familiaris_core_68_31
## 1031
                                 canis_familiaris_core_69_31
## 1032
                                 canis_familiaris_core_70_31
## 1033
                                 canis_familiaris_core_71_31
## 1034
                                 canis_familiaris_core_72_31
## 1035
                                 canis_familiaris_core_73_31
## 1036
                                 canis familiaris core 74 31
```

## 1037	canis_familiaris_core_75_31
## 1038	canis_familiaris_core_76_31
## 1039	<pre>canis_familiaris_core_77_31</pre>
## 1040	canis_familiaris_core_78_31
## 1041	canis_familiaris_core_79_31
## 1042	canis_familiaris_core_80_31
## 1043	canis_familiaris_core_81_31
## 1044	canis_familiaris_core_82_31
## 1045	canis_familiaris_core_83_31
## 1046	canis_familiaris_core_84_31
## 1047	canis_familiaris_core_85_31
## 1048	canis_familiaris_core_86_31
## 1049	canis_familiaris_core_87_31
## 1050	canis_familiaris_core_88_31
## 1051	canis_familiaris_core_89_31
## 1052	canis_familiaris_core_90_31
## 1053	canis_familiaris_core_91_31
## 1054	canis_familiaris_core_92_31
## 1055	canis_familiaris_core_93_31
## 1056	canis_familiaris_core_94_31
## 1057	canis_familiaris_core_95_31
## 1058	canis_familiaris_core_96_31
## 1059	canis_familiaris_core_97_31
## 1060	canis_familiaris_core_98_31
## 1061	canis_familiaris_core_99_31
## 1062	canis_familiaris_funcgen_55_2m
## 1063	canis_familiaris_funcgen_56_2m
## 1064	canis_familiaris_funcgen_57_2m
## 1065	canis_familiaris_funcgen_58_2n
## 1066	canis_familiaris_funcgen_59_20
## 1067	canis_familiaris_funcgen_60_2p
## 1068	canis_familiaris_funcgen_61_2q
## 1069	canis_familiaris_funcgen_62_2r
## 1070	canis_familiaris_funcgen_63_2
## 1071	canis_familiaris_funcgen_64_2
## 1072	canis_familiaris_funcgen_65_2
## 1073	canis_familiaris_funcgen_66_2
## 1074	canis_familiaris_funcgen_67_2
## 1075	canis_familiaris_funcgen_68_31
## 1076	canis_familiaris_funcgen_69_31
## 1077	canis_familiaris_funcgen_70_31
## 1078	canis_familiaris_funcgen_71_31
## 1079	canis_familiaris_funcgen_72_31
## 1080	canis_familiaris_funcgen_73_31
## 1081	canis_familiaris_funcgen_74_31
## 1082	canis_familiaris_funcgen_75_31
## 1083	canis_familiaris_funcgen_76_31
## 1084	canis_familiaris_funcgen_77_31
## 1085	canis_familiaris_funcgen_78_31
## 1086	canis_familiaris_funcgen_79_31
## 1087	canis_familiaris_funcgen_80_31
## 1087	canis_familiaris_funcgen_81_31
## 1089	canis_familiaris_funcgen_82_31
## 1009	canis_familiaris_funcgen_82_31
" H TOOO	caminalianing in cken 00 01

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## 1091
                              canis familiaris funcgen 84 31
## 1092
                              canis_familiaris_funcgen_85_31
## 1093
                              canis familiaris funcgen 86 31
## 1094
                              canis_familiaris_funcgen_87_31
## 1095
                              canis familiaris funcgen 88 31
## 1096
                              canis familiaris funcgen 89 31
                              canis familiaris_funcgen_90_31
## 1097
## 1098
                              canis familiaris funcgen 91 31
## 1099
                              canis familiaris funcgen 92 31
## 1100
                              canis_familiaris_funcgen_93_31
## 1101
                              canis_familiaris_funcgen_94_31
## 1102
                              canis_familiaris_funcgen_95_31
## 1103
                              canis_familiaris_funcgen_96_31
## 1104
                              canis_familiaris_funcgen_97_31
## 1105
                              canis_familiaris_funcgen_98_31
## 1106
                              canis_familiaris_funcgen_99_31
## 1107
                        canis_familiaris_otherfeatures_48_2f
## 1108
                        canis familiaris otherfeatures 49 2g
## 1109
                        canis_familiaris_otherfeatures_50_2h
## 1110
                        canis familiaris otherfeatures 51 2i
## 1111
                        canis_familiaris_otherfeatures_52_2j
## 1112
                        canis familiaris otherfeatures 53 2k
## 1113
                        canis_familiaris_otherfeatures_54_21
## 1114
                        canis familiaris otherfeatures 55 2m
## 1115
                        canis familiaris otherfeatures 56 2m
## 1116
                        canis familiaris otherfeatures 57 2m
## 1117
                        canis_familiaris_otherfeatures_58_2n
## 1118
                        canis_familiaris_otherfeatures_59_2o
## 1119
                        canis_familiaris_otherfeatures_60_2p
## 1120
                        canis_familiaris_otherfeatures_61_2q
## 1121
                        canis_familiaris_otherfeatures_62_2r
## 1122
                         canis_familiaris_otherfeatures_63_2
## 1123
                         canis_familiaris_otherfeatures_64_2
## 1124
                         canis_familiaris_otherfeatures_65_2
## 1125
                         canis familiaris otherfeatures 66 2
## 1126
                         canis familiaris otherfeatures 67 2
## 1127
                        canis familiaris otherfeatures 68 31
## 1128
                        canis_familiaris_otherfeatures_69_31
## 1129
                        canis familiaris otherfeatures 70 31
## 1130
                        canis_familiaris_otherfeatures_71_31
## 1131
                        canis familiaris otherfeatures 72 31
## 1132
                        canis familiaris otherfeatures 73 31
## 1133
                        canis familiaris otherfeatures 74 31
## 1134
                        canis_familiaris_otherfeatures_75_31
## 1135
                        canis_familiaris_otherfeatures_76_31
## 1136
                        canis_familiaris_otherfeatures_77_31
## 1137
                        canis familiaris otherfeatures 78 31
## 1138
                        canis_familiaris_otherfeatures_79_31
## 1139
                        canis_familiaris_otherfeatures_80_31
## 1140
                        canis_familiaris_otherfeatures_81_31
## 1141
                        canis_familiaris_otherfeatures_82_31
## 1142
                        canis_familiaris_otherfeatures_83_31
## 1143
                        canis_familiaris_otherfeatures_84_31
## 1144
                        canis familiaris otherfeatures 85 31
```

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## 1145
                        canis familiaris otherfeatures 86 31
                        canis_familiaris_otherfeatures_87_31
## 1146
                        canis familiaris otherfeatures 88 31
## 1147
## 1148
                        canis_familiaris_otherfeatures_89_31
## 1149
                        canis familiaris otherfeatures 90 31
                        canis familiaris otherfeatures 91 31
## 1150
## 1151
                        canis familiaris otherfeatures 92 31
                        canis familiaris otherfeatures 93 31
## 1152
## 1153
                        canis familiaris otherfeatures 94 31
## 1154
                        canis_familiaris_otherfeatures_95_31
## 1155
                        canis_familiaris_otherfeatures_96_31
## 1156
                        canis_familiaris_otherfeatures_97_31
## 1157
                        canis familiaris otherfeatures 98 31
## 1158
                        canis_familiaris_otherfeatures_99_31
## 1159
                               canis_familiaris_rnaseq_68_31
## 1160
                               canis_familiaris_rnaseq_69_31
## 1161
                               canis_familiaris_rnaseq_70_31
## 1162
                               canis familiaris rnaseg 71 31
## 1163
                               canis_familiaris_rnaseq_72_31
## 1164
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## 1165
                               canis_familiaris_rnaseq_74_31
## 1166
                               canis familiaris rnaseq 75 31
## 1167
                               canis_familiaris_rnaseq_76_31
                               canis familiaris_rnaseq_77_31
## 1168
## 1169
                               canis familiaris rnaseq 78 31
## 1170
                               canis familiaris rnaseq 79 31
## 1171
                               canis_familiaris_rnaseq_80_31
## 1172
                               canis_familiaris_rnaseq_81_31
## 1173
                               canis_familiaris_rnaseq_82_31
## 1174
                               canis_familiaris_rnaseq_83_31
## 1175
                               canis_familiaris_rnaseq_84_31
## 1176
                               canis_familiaris_rnaseq_85_31
## 1177
                               canis_familiaris_rnaseq_86_31
## 1178
                               canis_familiaris_rnaseq_87_31
## 1179
                               canis familiaris rnaseg 88 31
## 1180
                               canis_familiaris_rnaseq_89_31
## 1181
                               canis familiaris rnaseq 90 31
## 1182
                               canis_familiaris_rnaseq_91_31
## 1183
                               canis familiaris rnaseq 92 31
## 1184
                               canis_familiaris_rnaseq_93_31
## 1185
                               canis familiaris rnaseq 94 31
## 1186
                               canis familiaris rnaseq 95 31
## 1187
                               canis familiaris rnaseq 96 31
## 1188
                               canis_familiaris_rnaseq_97_31
## 1189
                               canis_familiaris_rnaseq_98_31
## 1190
                               canis_familiaris_rnaseq_99_31
## 1191
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## 1192
                            canis_familiaris_variation_49_2g
## 1193
                            canis_familiaris_variation_50_2h
## 1194
                            canis_familiaris_variation_51_2i
## 1195
                            canis_familiaris_variation_52_2j
## 1196
                            canis_familiaris_variation_53_2k
## 1197
                            canis_familiaris_variation_54_21
## 1198
                            canis_familiaris_variation_55_2m
```

```
## 1199
                            canis familiaris variation 56 2m
## 1200
                            canis_familiaris_variation_57_2m
                            canis familiaris variation 58 2m
## 1201
## 1202
                            canis_familiaris_variation_59_20
## 1203
                            canis familiaris variation 60 2p
## 1204
                            canis familiaris variation 61 2q
## 1205
                            canis familiaris variation 62 2r
## 1206
                             canis familiaris variation 63 2
## 1207
                             canis familiaris variation 64 2
## 1208
                             canis_familiaris_variation_65_2
## 1209
                             canis_familiaris_variation_66_2
                             canis_familiaris_variation_67_2
## 1210
## 1211
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                            canis_familiaris_variation_69_31
## 1212
## 1213
                            canis_familiaris_variation_70_31
## 1214
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## 1215
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## 1216
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## 1217
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## 1218
                            canis familiaris variation 75 31
## 1219
                            canis_familiaris_variation_76_31
## 1220
                            canis familiaris variation 77 31
## 1221
                            canis_familiaris_variation_78_31
## 1222
                            canis familiaris variation 79 31
## 1223
                            canis familiaris variation 80 31
## 1224
                            canis familiaris variation 81 31
## 1225
                            canis_familiaris_variation_82_31
## 1226
                            canis_familiaris_variation_83_31
## 1227
                            canis_familiaris_variation_84_31
## 1228
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## 1229
                            canis_familiaris_variation_86_31
## 1230
                            canis_familiaris_variation_87_31
## 1231
                            canis_familiaris_variation_88_31
## 1232
                            canis_familiaris_variation_89_31
## 1233
                            canis familiaris variation 90 31
## 1234
                            canis_familiaris_variation_91_31
## 1235
                            canis familiaris variation 92 31
## 1236
                            canis_familiaris_variation_93_31
## 1237
                            canis_familiaris_variation_94_31
## 1238
                            canis_familiaris_variation_95_31
## 1239
                            canis familiaris variation 96 31
## 1240
                            canis familiaris variation 97 31
## 1241
                            canis familiaris variation 98 31
                            canis_familiaris_variation_99_31
## 1242
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                                 canis_lupus_dingo_core_96_1
## 1244
## 1245
                                 canis_lupus_dingo_core_97_1
## 1246
                                 canis_lupus_dingo_core_98_1
## 1247
                                 canis_lupus_dingo_core_99_1
## 1248
                        canis_lupus_dingo_otherfeatures_95_1
## 1249
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## 1306
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## 1358
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## 1359
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##	1368	cavia_porcellus_core_59_3c
##	1369	cavia_porcellus_core_60_3d
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##	1371	cavia_porcellus_core_62_3f
##	1372	cavia_porcellus_core_63_3
##	1373	cavia_porcellus_core_64_3
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                             chlorocebus_sabaeus_rnaseq_83_1
## 1641
                             chlorocebus_sabaeus_rnaseq_84_1
                             chlorocebus_sabaeus_rnaseq_85_1
## 1642
## 1643
                             chlorocebus sabaeus rnaseg 86 1
## 1644
                             chlorocebus_sabaeus_rnaseq_87_1
## 1645
                             chlorocebus_sabaeus_rnaseq_88_1
## 1646
                             chlorocebus_sabaeus_rnaseq_89_1
## 1647
                             chlorocebus_sabaeus_rnaseq_90_1
## 1648
                             chlorocebus sabaeus rnaseg 91 1
## 1649
                             chlorocebus_sabaeus_rnaseq_92_1
## 1650
                             chlorocebus sabaeus rnaseg 93 1
                             chlorocebus_sabaeus_rnaseq_94_1
## 1651
## 1652
                             chlorocebus sabaeus rnaseg 95 1
                             chlorocebus_sabaeus_rnaseq_96_1
## 1653
## 1654
                             chlorocebus sabaeus rnaseg 97 1
## 1655
                             chlorocebus sabaeus rnaseg 98 1
                             chlorocebus_sabaeus_rnaseq_99_1
## 1656
## 1657
                          chlorocebus_sabaeus_variation_96_1
                          chlorocebus_sabaeus_variation_97_1
## 1658
## 1659
                          chlorocebus_sabaeus_variation_98_1
## 1660
                          chlorocebus_sabaeus_variation_99_1
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## 1662
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## 1663
                              choloepus_hoffmanni_core_55_1a
## 1664
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## 1665
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                              choloepus_hoffmanni_core_58_1b
## 1666
## 1667
                              choloepus hoffmanni core 59 1c
## 1668
                              choloepus_hoffmanni_core_60_1d
## 1669
                              choloepus hoffmanni core 61 1e
## 1670
                              choloepus_hoffmanni_core_62_1f
## 1671
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## 1672
                               choloepus hoffmanni core 64 1
                               choloepus hoffmanni core 65 1
## 1673
                               choloepus_hoffmanni_core_66_1
## 1674
## 1675
                               choloepus_hoffmanni_core_67_1
## 1676
                               choloepus_hoffmanni_core_68_1
## 1677
                               choloepus_hoffmanni_core_69_1
## 1678
                               choloepus_hoffmanni_core_70_1
## 1679
                               choloepus_hoffmanni_core_71_1
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## 1681
                               choloepus_hoffmanni_core_73_1
## 1682
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## 1683
                               choloepus_hoffmanni_core_75_1
## 1684
                               choloepus hoffmanni core 76 1
```

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## 1685
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## 1686
                               choloepus_hoffmanni_core_78_1
## 1687
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## 1688
                               choloepus_hoffmanni_core_80_1
## 1689
                               choloepus hoffmanni core 81 1
## 1690
                               choloepus hoffmanni core 82 1
                               choloepus hoffmanni_core_83_1
## 1691
## 1692
                               choloepus hoffmanni core 84 1
## 1693
                               choloepus hoffmanni core 85 1
## 1694
                               choloepus_hoffmanni_core_86_1
## 1695
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## 1696
                               choloepus_hoffmanni_core_88_1
## 1697
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## 1698
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## 1699
                               choloepus_hoffmanni_core_91_1
## 1700
                               choloepus_hoffmanni_core_92_1
## 1701
                               choloepus_hoffmanni_core_93_1
## 1702
                               choloepus hoffmanni core 94 1
## 1703
                               choloepus_hoffmanni_core_95_1
## 1704
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## 1706
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## 1707
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## 1708
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## 1709
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## 1710
                          chrysemys picta bellii core 97 303
## 1711
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## 1714
                chrysemys_picta_bellii_otherfeatures_96_303
## 1715
                chrysemys_picta_bellii_otherfeatures_97_303
## 1716
                chrysemys_picta_bellii_otherfeatures_98_303
## 1717
                chrysemys_picta_bellii_otherfeatures_99_303
## 1718
                       chrysemys_picta_bellii_rnaseq_95_303
## 1719
                       chrysemys picta bellii rnaseg 96 303
## 1720
                        chrysemys_picta_bellii_rnaseq_97_303
## 1721
                        chrysemys picta bellii rnaseq 98 303
## 1722
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## 1724
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## 1725
## 1726
                               ciona intestinalis core 49 2i
                               ciona intestinalis core 50 2j
## 1727
## 1728
                               ciona_intestinalis_core_51_2k
## 1729
                               ciona_intestinalis_core_52_21
## 1730
                               ciona_intestinalis_core_53_21
## 1731
                               ciona_intestinalis_core_54_21
## 1732
                               ciona_intestinalis_core_55_2m
## 1733
                               ciona_intestinalis_core_56_2m
## 1734
                               ciona_intestinalis_core_57_2m
## 1735
                               ciona_intestinalis_core_58_2n
## 1736
                               ciona intestinalis core 59 20
## 1737
                               ciona_intestinalis_core_60_2p
## 1738
                               ciona intestinalis core 61 2q
```

```
## 1739
                               ciona intestinalis core 62 2r
## 1740
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## 1741
                                ciona intestinalis core 64 2
## 1742
                                ciona_intestinalis_core_65_2
## 1743
                                ciona intestinalis core 66 3
## 1744
                                ciona intestinalis core 67 3
                                ciona_intestinalis core 68 3
## 1745
## 1746
                                ciona intestinalis core 69 3
## 1747
                                ciona intestinalis core 70 3
## 1748
                                ciona_intestinalis_core_71_3
## 1749
                                ciona_intestinalis_core_72_3
## 1750
                                ciona_intestinalis_core_73_3
## 1751
                                ciona_intestinalis_core_74_3
## 1752
                                ciona_intestinalis_core_75_3
## 1753
                                ciona_intestinalis_core_76_3
## 1754
                                ciona_intestinalis_core_77_3
## 1755
                                ciona_intestinalis_core_78_3
## 1756
                                ciona intestinalis core 79 3
## 1757
                                ciona_intestinalis_core_80_3
## 1758
                                ciona intestinalis core 81 3
## 1759
                                ciona_intestinalis_core_82_3
## 1760
                                ciona intestinalis core 83 3
## 1761
                                ciona_intestinalis_core_84_3
## 1762
                                ciona intestinalis core 85 3
## 1763
                                ciona intestinalis core 86 3
## 1764
                                ciona intestinalis core 87 3
## 1765
                                ciona_intestinalis_core_88_3
## 1766
                                ciona_intestinalis_core_89_3
## 1767
                                ciona_intestinalis_core_90_3
## 1768
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## 1769
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## 1771
                                ciona_intestinalis_core_94_3
## 1772
                                ciona_intestinalis_core_95_3
## 1773
                                ciona intestinalis core 96 3
## 1774
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## 1775
                                ciona intestinalis core 98 3
## 1776
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## 1778
                            ciona_intestinalis_funcgen_56_2m
## 1779
                            ciona intestinalis funcgen 57 2m
## 1780
                            ciona intestinalis funcgen 58 2n
                            ciona intestinalis funcgen 59 20
## 1781
## 1782
                            ciona_intestinalis_funcgen_60_2p
## 1783
                            ciona_intestinalis_funcgen_61_2q
## 1784
                            ciona_intestinalis_funcgen_62_2r
## 1785
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## 1786
                             ciona_intestinalis_funcgen_64_2
## 1787
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## 1791
                             ciona_intestinalis_funcgen_69_3
## 1792
                             ciona intestinalis funcgen 70 3
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## 1799
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                             ciona intestinalis funcgen 78 3
## 1800
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## 1802
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## 1803
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## 1805
                             ciona intestinalis funcgen 83 3
                             ciona_intestinalis_funcgen_84_3
## 1806
## 1807
                             ciona_intestinalis_funcgen_85_3
## 1808
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## 1809
                             ciona_intestinalis_funcgen_87_3
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## 1811
                             ciona intestinalis funcgen 89 3
## 1812
                             ciona intestinalis funcgen 90 3
## 1813
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## 1814
                             ciona intestinalis funcgen 92 3
## 1815
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## 1816
                             ciona intestinalis funcgen 94 3
## 1817
                             ciona intestinalis funcgen 95 3
## 1818
                             ciona intestinalis funcgen 96 3
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## 1820
                             ciona_intestinalis_funcgen_98_3
## 1821
                             ciona_intestinalis_funcgen_99_3
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## 1823
                     ciona_intestinalis_otherfeatures_49_2i
## 1824
                     ciona_intestinalis_otherfeatures_50_2j
## 1825
                     ciona_intestinalis_otherfeatures_51_2k
## 1826
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## 1827
                     ciona intestinalis otherfeatures 53 21
## 1828
                     ciona intestinalis otherfeatures 54 21
## 1829
                     ciona intestinalis otherfeatures 55 2m
## 1830
                     ciona_intestinalis_otherfeatures_56_2m
## 1831
                     ciona intestinalis otherfeatures 57 2m
## 1832
                     ciona_intestinalis_otherfeatures_58_2n
## 1833
                     ciona intestinalis otherfeatures 59 20
## 1834
                     ciona intestinalis otherfeatures 60 2p
                     ciona intestinalis otherfeatures 61 2q
## 1835
## 1836
                     ciona_intestinalis_otherfeatures_62_2r
## 1837
                      ciona_intestinalis_otherfeatures_63_2
                      ciona_intestinalis_otherfeatures_64_2
## 1838
## 1839
                      ciona intestinalis otherfeatures 65 2
## 1840
                      ciona_intestinalis_otherfeatures_66_3
## 1841
                      ciona_intestinalis_otherfeatures_67_3
                      ciona_intestinalis_otherfeatures_68_3
## 1842
## 1843
                      ciona_intestinalis_otherfeatures_69_3
                      ciona_intestinalis_otherfeatures_70_3
## 1844
## 1845
                      ciona_intestinalis_otherfeatures_71_3
## 1846
                      ciona intestinalis otherfeatures 72 3
```

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## 1847
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                      ciona_intestinalis_otherfeatures_74_3
## 1849
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## 1850
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## 1851
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## 1852
                      ciona intestinalis otherfeatures 78 3
## 1853
                      ciona intestinalis otherfeatures 79 3
                      ciona intestinalis otherfeatures 80 3
## 1854
## 1855
                      ciona intestinalis otherfeatures 81 3
## 1856
                      ciona_intestinalis_otherfeatures_82_3
## 1857
                      ciona_intestinalis_otherfeatures_83_3
                      ciona_intestinalis_otherfeatures_84_3
## 1858
## 1859
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## 1860
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## 1862
                       ciona_intestinalis_otherfeatures_88_3
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## 1867
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## 1868
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## 1869
                      ciona_intestinalis_otherfeatures_95_3
## 1870
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## 1871
                      ciona intestinalis otherfeatures 97 3
## 1872
                      ciona intestinalis otherfeatures 98 3
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## 1875
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## 1876
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## 1877
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## 1878
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## 1879
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## 1880
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## 1881
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## 1882
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## 1883
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## 1884
                                   ciona_savignyi_core_58_2i
## 1885
                                   ciona_savignyi_core_59_2j
## 1886
                                   ciona_savignyi_core_60_2k
## 1887
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## 1888
                                   ciona savignyi core 62 2m
## 1889
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## 1890
                                    ciona_savignyi_core_64_2
## 1891
                                    ciona_savignyi_core_65_2
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## 1893
                                    ciona_savignyi_core_67_2
## 1894
                                    ciona_savignyi_core_68_2
## 1895
                                    ciona_savignyi_core_69_2
## 1896
                                    ciona_savignyi_core_70_2
## 1897
                                    ciona_savignyi_core_71_2
## 1898
                                    ciona_savignyi_core_72_2
## 1899
                                    ciona_savignyi_core_73_2
## 1900
                                    ciona savignyi core 74 2
```

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## 1901
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                                    ciona_savignyi_core_78_2
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                                    ciona savignyi core 82 2
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                                    ciona_savignyi_core_83_2
## 1910
                                    ciona_savignyi_core_84_2
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                                    ciona_savignyi_core_85_2
## 1912
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## 1923
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## 1928
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                          ciona_savignyi_otherfeatures_51_2g
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## 1931
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## 1932
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## 1934
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## 1935
                          ciona savignyi otherfeatures 57 2h
                          ciona_savignyi_otherfeatures_58_2i
## 1936
## 1937
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## 1938
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                          ciona_savignyi_otherfeatures_75_2
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## 1954
                           ciona savignyi otherfeatures 76 2
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## 1964
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                          ciona_savignyi_otherfeatures_87_2
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## 1990
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## 2005
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## 2006
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## 2007
            colobus angolensis palliatus otherfeatures 98 1
## 2008
            colobus angolensis palliatus otherfeatures 99 1
```

```
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## 2014
                               cottoperca gobio rnaseq 99 31
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## 2016
                                 coturnix japonica core 97 2
## 2017
                                 coturnix japonica core 98 2
## 2018
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## 2019
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## 2020
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## 2021
                       coturnix_japonica_otherfeatures_98_2
## 2022
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## 2023
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## 2024
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## 2025
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## 2026
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## 2033
                     cricetulus griseus chok1gshd core 96 1
## 2034
                     cricetulus griseus chok1gshd core 97 1
## 2035
                     cricetulus_griseus_chok1gshd_core_98_1
## 2036
                      cricetulus_griseus_chok1gshd_core_99_1
## 2037
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## 2283	danio_rerio_funcgen_74_9
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## 2285	danio_rerio_funcgen_76_9
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	danio_rerio_vega_62_9b
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##	2464	dasypus_novemcinctus_core_51_1g
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##	2472	dasypus_novemcinctus_core_59_2c
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## 310 1 ## 3105	ensembl_production_83
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## 3222
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## 3239
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## 3291
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## 3292
                           equus caballus otherfeatures 67 2
## 3293
                           equus caballus otherfeatures 68 2
## 3294
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## 3295
                           equus_caballus_otherfeatures_70_2
## 3296
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## 3297
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## 3298
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## 3299
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## 3300
                           equus_caballus_otherfeatures_75_2
## 3301
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## 3302
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## 3303
                           equus_caballus_otherfeatures_78_2
## 3304
                           equus caballus otherfeatures 79 2
```

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## 3310
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## 3311
                           equus caballus otherfeatures 86 2
## 3312
                           equus caballus otherfeatures 87 2
## 3313
                           equus caballus otherfeatures 88 2
## 3314
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## 3315
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                           equus_caballus_otherfeatures_91_2
## 3316
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## 3318
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## 3319
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## 3327
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## 3329
                                  equus caballus rnaseq 99 3
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## 3334
                              equus_caballus_variation_61_2h
## 3335
                              equus_caballus_variation_62_2i
## 3336
                               equus_caballus_variation_63_2
## 3337
                               equus_caballus_variation_64_2
## 3338
                               equus_caballus_variation_65_2
## 3339
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## 3340
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## 3341
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## 3342
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## 3348
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## 3349
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## 3350
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## 3351
                               equus_caballus_variation_78_2
## 3352
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## 3353
                               equus_caballus_variation_80_2
## 3354
                               equus_caballus_variation_81_2
## 3355
                               equus_caballus_variation_82_2
## 3356
                               equus_caballus_variation_83_2
## 3357
                               equus_caballus_variation_84_2
## 3358
                               equus_caballus_variation_85_2
```

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## 3359
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## 3360
                               equus caballus variation 87 2
## 3361
                               equus caballus variation 88 2
## 3362
                               equus_caballus_variation_89_2
## 3363
                               equus caballus variation 90 2
## 3364
                               equus caballus variation 91 2
## 3365
                               equus caballus variation 92 2
## 3366
                               equus caballus variation 93 2
## 3367
                               equus caballus variation 94 2
## 3368
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## 3377
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## 3378
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## 3382
                              erinaceus europaeus core 57 1e
## 3383
                              erinaceus europaeus core 58 1f
## 3384
                              erinaceus europaeus core 59 1g
## 3385
                              erinaceus_europaeus_core_60_1h
## 3386
                              erinaceus_europaeus_core_61_1i
## 3387
                              erinaceus_europaeus_core_62_1j
## 3388
                               erinaceus_europaeus_core_63_1
## 3389
                               erinaceus_europaeus_core_64_1
## 3390
                               erinaceus_europaeus_core_65_1
## 3391
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## 3392
                               erinaceus_europaeus_core_67_1
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## 3394
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## 3395
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## 3396
                               erinaceus_europaeus_core_71_1
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## 3398
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## 3400
                               erinaceus europaeus core 75 1
## 3401
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## 3402
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## 3403
                               erinaceus_europaeus_core_78_1
## 3404
                               erinaceus_europaeus_core_79_1
## 3405
                               erinaceus_europaeus_core_80_1
## 3406
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## 3407
                               erinaceus_europaeus_core_82_1
## 3408
                               erinaceus_europaeus_core_83_1
## 3409
                               erinaceus_europaeus_core_84_1
## 3410
                               erinaceus_europaeus_core_85_1
## 3411
                               erinaceus_europaeus_core_86_1
## 3412
                               erinaceus europaeus core 87 1
```

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## 3414
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## 3418
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## 3419
                               erinaceus europaeus core 94 1
## 3420
                               erinaceus europaeus core 95 1
## 3421
                               erinaceus europaeus core 96 1
## 3422
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## 3435
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## 3436
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## 3437
                                       esox lucius core 98 3
## 3438
                                       esox lucius core 99 3
## 3439
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## 3445
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## 3446
                                     esox_lucius_rnaseq_95_3
## 3447
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## 3448
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## 3450
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## 3452
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## 3453
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## 3454
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## 3455
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## 3456
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## 3457
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## 3458
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## 3459
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## 3460
                                      felis_catus_core_57_1f
## 3461
                                      felis_catus_core_58_1g
## 3462
                                      felis_catus_core_59_1h
## 3463
                                      felis_catus_core_60_1i
## 3464
                                      felis_catus_core_61_1j
## 3465
                                      felis_catus_core_62_1k
## 3466
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```

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##	3475	felis_catus_core_72_62
##	3476	felis_catus_core_73_62
##	3477	felis_catus_core_74_62
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##	3490	felis_catus_core_87_62
##	3491	felis_catus_core_88_62
##	3492	felis_catus_core_89_62
##	3493	felis_catus_core_90_62
##	3494	felis_catus_core_91_8
##	3495	felis_catus_core_92_8
##	3496	felis_catus_core_93_9
##	3497	felis_catus_core_94_9
	3498	felis_catus_core_95_9
	3499	felis_catus_core_96_9
	3500	felis_catus_core_97_9
	3501	felis_catus_core_98_9
	3502	felis_catus_core_99_9
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	3504	felis_catus_funcgen_93_9
	3505	felis_catus_funcgen_94_9
	3506	felis_catus_funcgen_95_9
	3507	felis_catus_funcgen_96_9
	3508	felis_catus_funcgen_97_9
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##	3510	felis_catus_funcgen_99_9
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##	3512	<pre>felis_catus_otherfeatures_71_62</pre>
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##	3515	felis_catus_otherfeatures_74_62
	3516	felis_catus_otherfeatures_75_62
	3517	felis_catus_otherfeatures_76_62
	3518	felis_catus_otherfeatures_77_62
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	3520	felis_catus_otherfeatures_79_62
##	3020	Terra_carna_ornerrearnrea_19_02

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## 3521
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## 3526
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## 3527
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## 3528
                             felis catus otherfeatures 87 62
                             felis_catus_otherfeatures 88 62
## 3529
## 3530
                             felis_catus_otherfeatures_89_62
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## 3532
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## 3546
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## 3547
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## 3552
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## 3553
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## 3554
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## 3555
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## 3556
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## 3557
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## 3559
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## 3560
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## 3562
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                                     felis catus rnaseq 92 8
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## 3582
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                                 felis catus variation 73 62
## 3584
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## 3614
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## 3616
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                               ficedula albicollis core 80 1
## 3617
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## 3619
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## 3624
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## 3625
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## 3626
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## 3627
                               ficedula_albicollis_core_90_1
## 3628
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## 3636
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## 3649
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                             ficedula albicollis rnaseq 91 1
```

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## 3683
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## 3684
                             ficedula albicollis rnaseq 93 1
## 3685
                             ficedula albicollis rnaseq 94 1
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                             ficedula albicollis rnaseq 96 1
## 3688
                             ficedula albicollis rnaseq 97 1
## 3689
                             ficedula albicollis rnaseg 98 1
                             ficedula albicollis rnaseq 99 1
## 3690
## 3691
                                fukomys damarensis core 90 1
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## 3693
                                fukomys_damarensis_core_92_1
                                fukomys_damarensis_core_93_1
## 3694
## 3695
                                fukomys damarensis core 94 1
## 3696
                                fukomys_damarensis_core_95_1
## 3697
                                fukomys_damarensis_core_96_1
## 3698
                                fukomys_damarensis_core_97_1
## 3699
                                fukomys_damarensis_core_98_1
## 3700
                                fukomys damarensis core 99 1
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## 3702
                      fukomys damarensis otherfeatures 91 1
## 3703
                      fukomys_damarensis_otherfeatures_92_1
## 3704
                      fukomys damarensis otherfeatures 93 1
## 3705
                      fukomys_damarensis_otherfeatures_94_1
## 3706
                      fukomys damarensis otherfeatures 95 1
                      fukomys damarensis otherfeatures 96 1
## 3707
## 3708
                      fukomys damarensis otherfeatures 97 1
## 3709
                      fukomys_damarensis_otherfeatures_98_1
## 3710
                      fukomys_damarensis_otherfeatures_99_1
                              fukomys_damarensis_rnaseq_90_1
## 3711
## 3712
                              fukomys damarensis rnaseg 91 1
                              fukomys_damarensis_rnaseq_92_1
## 3713
## 3714
                              fukomys_damarensis_rnaseq_93_1
## 3715
                              fukomys_damarensis_rnaseq_94_1
## 3716
                              fukomys_damarensis_rnaseq_95_1
## 3717
                              fukomys damarensis rnaseg 96 1
## 3718
                              fukomys damarensis rnaseg 97 1
## 3719
                              fukomys damarensis rnaseg 98 1
## 3720
                              fukomys_damarensis_rnaseq_99_1
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                           fundulus_heteroclitus_core_95_302
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                           fundulus heteroclitus core 96 302
## 3724
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                           fundulus heteroclitus core 98 302
## 3725
## 3726
                           fundulus_heteroclitus_core_99_302
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                        fundulus_heteroclitus_funcgen_96_302
## 3728
                        fundulus_heteroclitus_funcgen_97_302
## 3729
                       fundulus heteroclitus funcgen 98 302
## 3730
                        fundulus heteroclitus funcgen 99 302
## 3731
                 fundulus heteroclitus otherfeatures 94 302
## 3732
                 fundulus_heteroclitus_otherfeatures_95_302
## 3733
                 fundulus_heteroclitus_otherfeatures_96_302
## 3734
                 fundulus heteroclitus otherfeatures 97 302
## 3735
                 fundulus heteroclitus otherfeatures 98 302
## 3736
                 fundulus heteroclitus otherfeatures 99 302
```

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## 3737
                         fundulus heteroclitus rnaseg 94 302
## 3738
                         fundulus heteroclitus rnaseq 95 302
## 3739
                         fundulus heteroclitus rnaseq 96 302
## 3740
                         fundulus_heteroclitus_rnaseq_97_302
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                         fundulus heteroclitus rnaseq 98 302
## 3742
                         fundulus heteroclitus rnaseq 99 302
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## 3746
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## 3747
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## 3749
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## 3750
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## 3752
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## 3753
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## 3754
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## 3755
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## 3756
                                      gadus_morhua_core_78_1
## 3757
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## 3758
                                      gadus morhua core 80 1
## 3759
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## 3761
                                      gadus_morhua_core_83_1
## 3762
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## 3772
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## 3773
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## 3774
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## 3775
                                      gadus_morhua_core_97_1
## 3776
                                      gadus_morhua_core_98_1
## 3777
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                                    gallus gallus core 48 2f
## 3779
                                    gallus gallus core 49 2g
## 3780
                                    gallus_gallus_core_50_2h
## 3781
                                    gallus_gallus_core_51_2i
## 3782
                                    gallus_gallus_core_52_2j
## 3783
                                    gallus_gallus_core_53_2k
## 3784
                                    gallus_gallus_core_54_21
## 3785
                                    gallus_gallus_core_55_2m
## 3786
                                    gallus_gallus_core_56_2m
## 3787
                                    gallus_gallus_core_57_2m
## 3788
                                    gallus_gallus_core_58_2n
## 3789
                                    gallus_gallus_core_59_2o
## 3790
                                    gallus_gallus_core_60_2p
```

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## 3792	gallus_gallus_core_62_2r
## 3793	gallus_gallus_core_63_2
## 3794	gallus_gallus_core_64_2
## 3795	gallus_gallus_core_65_2
## 3796	gallus_gallus_core_66_2
## 3797	gallus_gallus_core_67_2
## 3798	gallus_gallus_core_68_2
## 3799	gallus_gallus_core_69_2
## 3800	gallus_gallus_core_70_2
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## 3803	gallus_gallus_core_73_4
## 3804	gallus_gallus_core_74_4
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## 3807	gallus_gallus_core_77_4
## 3808	gallus_gallus_core_78_4
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## 3811	gallus_gallus_core_81_4
## 3812	gallus_gallus_core_82_4
## 3813	gallus_gallus_core_83_4
## 3814	gallus_gallus_core_84_4
## 3815	gallus_gallus_core_85_4
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## 3817	gallus_gallus_core_87_5
## 3818	gallus_gallus_core_88_5
## 3819	gallus_gallus_core_89_5
## 3820	
	gallus_gallus_core_90_5
## 3821	gallus_gallus_core_91_5
## 3822	gallus_gallus_core_92_5
## 3823	gallus_gallus_core_93_5
## 3824	gallus_gallus_core_94_5
## 3825	gallus_gallus_core_95_6
## 3826	gallus_gallus_core_96_6
## 3827	gallus_gallus_core_97_6
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## 3829	gallus_gallus_core_99_6
## 3830	gallus_gallus_funcgen_55_2m
## 3831	gallus_gallus_funcgen_56_2m
## 3832	gallus_gallus_funcgen_57_2m
## 3833	gallus_gallus_funcgen_58_2n
## 3834	gallus_gallus_funcgen_59_20
## 3835	gallus_gallus_funcgen_60_2p
## 3836	gallus_gallus_funcgen_61_2q
## 3837	gallus_gallus_funcgen_62_2r
## 3838	gallus_gallus_funcgen_63_2
## 3839	gallus_gallus_funcgen_64_2
## 3840 ## 3841	gallus_gallus_funcgen_65_2
## 3841 ## 2040	gallus_gallus_funcgen_66_2
## 3842	gallus_gallus_funcgen_67_2
## 3843	gallus_gallus_funcgen_68_2
## 3844	gallus_gallus_funcgen_69_2

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## 3845
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## 3846
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## 3847
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## 3848
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                                  gallus gallus funcgen 74 4
##
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## 3852
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                                  gallus gallus funcgen 78 4
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  3856
                                  gallus_gallus_funcgen_81_4
##
##
  3857
                                  gallus_gallus_funcgen_82_4
  3858
                                  gallus_gallus_funcgen_83_4
##
##
  3859
                                  gallus_gallus_funcgen_84_4
##
  3860
                                  gallus_gallus_funcgen_85_4
##
  3861
                                  gallus_gallus_funcgen_86_5
  3862
##
                                  gallus gallus funcgen 87 5
##
  3863
                                  gallus_gallus_funcgen_88_5
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                                  gallus gallus funcgen 89 5
##
  3865
                                  gallus_gallus_funcgen_90_5
  3866
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## 3867
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## 3869
                                  gallus gallus funcgen 94 5
  3870
                                  gallus gallus funcgen 95 6
## 3871
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  3872
                                  gallus_gallus_funcgen_97_6
  3873
##
                                  gallus_gallus_funcgen_98_6
  3874
                                  gallus gallus funcgen 99 6
## 3875
                           gallus_gallus_otherfeatures_48_2f
  3876
                           gallus_gallus_otherfeatures_49_2g
  3877
##
                           gallus_gallus_otherfeatures_50_2h
## 3878
                           gallus_gallus_otherfeatures_51_2i
  3879
                           gallus gallus otherfeatures 52 2j
##
  3880
                           gallus gallus otherfeatures 53 2k
## 3881
                           gallus gallus otherfeatures 54 21
## 3882
                           gallus_gallus_otherfeatures_55_2m
  3883
##
                           gallus gallus otherfeatures 56 2m
  3884
##
                           gallus_gallus_otherfeatures_57_2m
                          gallus_gallus_otherfeatures 58 2n
  3885
##
  3886
                           gallus gallus otherfeatures 59 20
   3887
                           gallus gallus otherfeatures 60 2p
  3888
                           gallus_gallus_otherfeatures_61_2q
##
  3889
                           gallus_gallus_otherfeatures_62_2r
## 3890
                            gallus_gallus_otherfeatures_63_2
  3891
                            gallus_gallus_otherfeatures_64_2
## 3892
                            gallus_gallus_otherfeatures_65_2
##
  3893
                            gallus_gallus_otherfeatures_66_2
  3894
##
                            gallus_gallus_otherfeatures_67_2
##
  3895
                            gallus_gallus_otherfeatures_68_2
## 3896
                            gallus_gallus_otherfeatures_69_2
## 3897
                            gallus_gallus_otherfeatures_70_2
## 3898
                            gallus gallus otherfeatures 71 4
```

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## 3899
                            gallus gallus otherfeatures 72 4
                            gallus_gallus_otherfeatures_73_4
## 3900
## 3901
                            gallus gallus otherfeatures 74 4
## 3902
                            gallus_gallus_otherfeatures_75_4
  3903
                            gallus gallus otherfeatures 76 4
##
  3904
                            gallus gallus otherfeatures 77 4
## 3905
                            gallus gallus otherfeatures 78 4
                            gallus_gallus_otherfeatures_79_4
## 3906
  3907
                            gallus gallus otherfeatures 80 4
## 3908
                            gallus_gallus_otherfeatures_81_4
  3909
                            gallus_gallus_otherfeatures_82_4
## 3910
                            gallus_gallus_otherfeatures_83_4
  3911
                            gallus_gallus_otherfeatures_84_4
## 3912
                            gallus_gallus_otherfeatures_85_4
## 3913
                            gallus_gallus_otherfeatures_86_5
## 3914
                            gallus_gallus_otherfeatures_87_5
## 3915
                            gallus_gallus_otherfeatures_88_5
## 3916
                            gallus gallus otherfeatures 89 5
## 3917
                            gallus_gallus_otherfeatures_90_5
## 3918
                            gallus gallus otherfeatures 91 5
                            gallus_gallus_otherfeatures_92_5
## 3919
## 3920
                            gallus gallus otherfeatures 93 5
## 3921
                            gallus_gallus_otherfeatures_94_5
## 3922
                            gallus gallus otherfeatures 95 6
## 3923
                            gallus gallus otherfeatures 96 6
## 3924
                            gallus_gallus_otherfeatures_97_6
## 3925
                            gallus_gallus_otherfeatures_98_6
  3926
                            gallus_gallus_otherfeatures_99_6
## 3927
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## 3928
                                   gallus_gallus_rnaseq_72_4
## 3929
                                   gallus_gallus_rnaseq_73_4
  3930
                                   gallus_gallus_rnaseq_74_4
## 3931
                                   gallus_gallus_rnaseq_75_4
## 3932
                                   gallus_gallus_rnaseq_76_4
## 3933
                                   gallus gallus rnaseq 77 4
  3934
                                   gallus_gallus_rnaseq_78_4
## 3935
                                   gallus gallus rnaseq 79 4
## 3936
                                   gallus_gallus_rnaseq_80_4
## 3937
                                   gallus_gallus_rnaseq_81_4
## 3938
                                   gallus_gallus_rnaseq_82_4
## 3939
                                   gallus gallus rnaseq 83 4
                                   gallus_gallus_rnaseq_84_4
## 3940
  3941
                                   gallus gallus rnaseq 85 4
## 3942
                                   gallus_gallus_rnaseq_86_5
## 3943
                                   gallus_gallus_rnaseq_87_5
## 3944
                                   gallus_gallus_rnaseq_88_5
## 3945
                                   gallus_gallus_rnaseq_89_5
## 3946
                                   gallus_gallus_rnaseq_90_5
## 3947
                                   gallus_gallus_rnaseq_91_5
## 3948
                                   gallus_gallus_rnaseq_92_5
## 3949
                                   gallus_gallus_rnaseq_93_5
## 3950
                                   gallus gallus rnaseq 94 5
## 3951
                                   gallus_gallus_rnaseq_95_6
## 3952
                                   gallus_gallus_rnaseq_96_6
```

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	3953	gallus_gallus_rnaseq_97_6
	3954	gallus_gallus_rnaseq_98_6
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##	3957	<pre>gallus_gallus_variation_49_2g</pre>
##	3958	gallus_gallus_variation_50_2h
	3959	gallus_gallus_variation_51_2i
	3960	gallus_gallus_variation_52_2j
	3961	gallus_gallus_variation_53_2k
	3962	gallus_gallus_variation_54_21
	3963	
		gallus_gallus_variation_55_2m
	3964	gallus_gallus_variation_56_2m
	3965	gallus_gallus_variation_57_2m
	3966	<pre>gallus_gallus_variation_58_2n</pre>
	3967	<pre>gallus_gallus_variation_59_2o</pre>
	3968	<pre>gallus_gallus_variation_60_2p</pre>
##	3969	<pre>gallus_gallus_variation_61_2q</pre>
##	3970	<pre>gallus_gallus_variation_62_2r</pre>
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##	3972	gallus_gallus_variation_64_2
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##	3974	gallus_gallus_variation_66_2
##	3975	gallus_gallus_variation_67_2
	3976	gallus_gallus_variation_68_2
	3977	gallus_gallus_variation_69_2
	3978	gallus_gallus_variation_70_2
	3979	gallus_gallus_variation_70_2
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	3981	-
		gallus_gallus_variation_73_4
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	3983	gallus_gallus_variation_75_4
	3984	gallus_gallus_variation_76_4
	3985	gallus_gallus_variation_77_4
	3986	gallus_gallus_variation_78_4
##	3987	gallus_gallus_variation_79_4
##	3988	gallus_gallus_variation_80_4
##	3989	gallus_gallus_variation_81_4
##	3990	gallus_gallus_variation_82_4
##	3991	gallus_gallus_variation_83_4
##	3992	gallus_gallus_variation_84_4
##	3993	gallus_gallus_variation_85_4
	3994	gallus_gallus_variation_86_5
	3995	gallus_gallus_variation_87_5
	3996	gallus_gallus_variation_88_5
	3997	gallus_gallus_variation_89_5
	3998	gallus_gallus_variation_90_5
	3999	
##	4000	gallus_gallus_variation_91_5
		gallus_gallus_variation_92_5
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## 4012
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## 4013
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## 4014
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## 4015
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## 4017
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## 4022
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## 4058
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## 4059
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## 4060
                            gasterosteus_aculeatus_core_88_1
```

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## 4062
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## 4073
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## 4074
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## 4467
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## 4468
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## 4492
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##	4708	homo_sapiens_vega_63_37

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## 4816
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## ## ## ## ## ##	5224 5225 5226 5227 5228 5229 5230 5231	macaca_mulatta_core_98_10 macaca_mulatta_core_99_10 macaca_mulatta_funcgen_55_101 macaca_mulatta_funcgen_56_101 macaca_mulatta_funcgen_57_101 macaca_mulatta_funcgen_58_10m macaca_mulatta_funcgen_59_10n macaca_mulatta_funcgen_60_100 macaca_mulatta_funcgen_61_10p
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## ## ## ## ## ## ##	5224 5225 5226 5227 5228 5229 5230 5231 5232 5233 5234	macaca_mulatta_core_98_10 macaca_mulatta_core_99_10 macaca_mulatta_funcgen_55_101 macaca_mulatta_funcgen_56_101 macaca_mulatta_funcgen_57_101 macaca_mulatta_funcgen_58_10m macaca_mulatta_funcgen_59_10n macaca_mulatta_funcgen_60_10o macaca_mulatta_funcgen_61_10p macaca_mulatta_funcgen_62_10q macaca_mulatta_funcgen_63_10 macaca_mulatta_funcgen_64_10
## ## ## ## ## ## ##	5224 5225 5226 5227 5228 5229 5230 5231 5232 5233 5234 5235	macaca_mulatta_core_98_10 macaca_mulatta_core_99_10 macaca_mulatta_funcgen_55_101 macaca_mulatta_funcgen_56_101 macaca_mulatta_funcgen_57_101 macaca_mulatta_funcgen_58_10m macaca_mulatta_funcgen_59_10n macaca_mulatta_funcgen_60_10o macaca_mulatta_funcgen_61_10p macaca_mulatta_funcgen_62_10q macaca_mulatta_funcgen_63_10 macaca_mulatta_funcgen_64_10 macaca_mulatta_funcgen_65_10
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## ## ## ## ## ## ## ## ##	5224 5225 5226 5227 5228 5229 5230 5231 5232 5233 5234 5235 5236 5237 5238 5239	macaca_mulatta_core_98_10 macaca_mulatta_core_99_10 macaca_mulatta_funcgen_55_101 macaca_mulatta_funcgen_56_101 macaca_mulatta_funcgen_57_101 macaca_mulatta_funcgen_58_10m macaca_mulatta_funcgen_59_10n macaca_mulatta_funcgen_60_10o macaca_mulatta_funcgen_61_10p macaca_mulatta_funcgen_62_10q macaca_mulatta_funcgen_62_10q macaca_mulatta_funcgen_63_10 macaca_mulatta_funcgen_64_10 macaca_mulatta_funcgen_65_10 macaca_mulatta_funcgen_66_10 macaca_mulatta_funcgen_66_10 macaca_mulatta_funcgen_68_10 macaca_mulatta_funcgen_68_10 macaca_mulatta_funcgen_69_10
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######################################	5224 5225 5226 5227 5228 5229 5230 5231 5232 5233 5234 5235 5236 5237 5238 5239 5240 5241 5242 5243	macaca_mulatta_core_98_10 macaca_mulatta_core_99_10 macaca_mulatta_funcgen_55_101 macaca_mulatta_funcgen_56_101 macaca_mulatta_funcgen_57_101 macaca_mulatta_funcgen_58_10m macaca_mulatta_funcgen_59_10n macaca_mulatta_funcgen_60_10o macaca_mulatta_funcgen_61_10p macaca_mulatta_funcgen_62_10q macaca_mulatta_funcgen_62_10q macaca_mulatta_funcgen_63_10 macaca_mulatta_funcgen_64_10 macaca_mulatta_funcgen_65_10 macaca_mulatta_funcgen_66_10 macaca_mulatta_funcgen_66_10 macaca_mulatta_funcgen_68_10 macaca_mulatta_funcgen_69_10 macaca_mulatta_funcgen_70_10 macaca_mulatta_funcgen_70_10 macaca_mulatta_funcgen_71_10 macaca_mulatta_funcgen_72_10 macaca_mulatta_funcgen_73_10
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##	6628	mus_musculus_vega_49_37b
##	6629	mus_musculus_vega_50_37c
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	6632	mus_musculus_vega_53_37f
	6633	mus_musculus_vega_54_37g
	6634	mus_musculus_vega_55_37h
	6635	mus_musculus_vega_56_37i
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##	6636 6637	mus_musculus_vega_57_37j mus_musculus_vega_58_37k
##	6638	mus_musculus_vega_50_37k mus_musculus_vega_59_371
##	6639	mus_musculus_vega_59_371 mus_musculus_vega_60_37m
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##	6644	mus_musculus_vega_65_37
##	6645	mus_musculus_vega_66_37
##	6646	mus_musculus_vega_67_37
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##	6648	mus_musculus_vega_69_38
##	6649	mus_musculus_vega_70_38
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##	6651	mus_musculus_vega_72_38
444	6652	mus_musculus_vega_73_38
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## 7373
                          ornithorhynchus_anatinus_core_82_1
## 7374
                          ornithorhynchus anatinus core 83 1
## 7375
                          ornithorhynchus_anatinus_core_84_1
## 7376
                          ornithorhynchus anatinus core 85 1
                          ornithorhynchus_anatinus_core_86_1
## 7377
## 7378
                          ornithorhynchus anatinus core 87 1
                          ornithorhynchus anatinus core 88 1
## 7379
                          ornithorhynchus_anatinus_core_89_1
## 7380
## 7381
                          ornithorhynchus_anatinus_core_90_1
                          ornithorhynchus_anatinus_core_91_1
## 7382
## 7383
                          ornithorhynchus_anatinus_core_92_1
## 7384
                          ornithorhynchus_anatinus_core_93_1
## 7385
                          ornithorhynchus_anatinus_core_94_1
## 7386
                          ornithorhynchus_anatinus_core_95_1
## 7387
                          ornithorhynchus_anatinus_core_96_1
## 7388
                          ornithorhynchus_anatinus_core_97_1
## 7389
                          ornithorhynchus anatinus core 98 1
## 7390
                          ornithorhynchus anatinus core 99 1
## 7391
                     ornithorhynchus anatinus funcgen 55 1k
## 7392
                     ornithorhynchus_anatinus_funcgen_56_1k
## 7393
                     ornithorhynchus anatinus funcgen 57 1k
## 7394
                     ornithorhynchus_anatinus_funcgen_58_11
## 7395
                     ornithorhynchus anatinus funcgen 59 1m
## 7396
                     ornithorhynchus anatinus funcgen 60 1n
                     ornithorhynchus anatinus funcgen 61 10
## 7397
## 7398
                     ornithorhynchus_anatinus_funcgen_62_1p
## 7399
                      ornithorhynchus_anatinus_funcgen_63_1
## 7400
                      ornithorhynchus_anatinus_funcgen_64_1
## 7401
                      ornithorhynchus anatinus funcgen 65 1
## 7402
                       ornithorhynchus_anatinus_funcgen_66_1
## 7403
                      ornithorhynchus_anatinus_funcgen_67_1
                       ornithorhynchus_anatinus_funcgen_68_1
## 7404
## 7405
                       ornithorhynchus_anatinus_funcgen_69_1
## 7406
                      ornithorhynchus_anatinus_funcgen_70_1
## 7407
                      ornithorhynchus anatinus funcgen 71 1
## 7408
                      ornithorhynchus anatinus funcgen 72 1
```

```
ornithorhynchus anatinus funcgen 73 1
## 7409
## 7410
                      ornithorhynchus anatinus funcgen 74 1
                      ornithorhynchus anatinus funcgen 75 1
## 7411
## 7412
                      ornithorhynchus anatinus funcgen 76 1
## 7413
                      ornithorhynchus anatinus funcgen 77 1
## 7414
                      ornithorhynchus anatinus funcgen 78 1
## 7415
                      ornithorhynchus anatinus funcgen 79 1
                      ornithorhynchus anatinus funcgen 80 1
## 7416
## 7417
                      ornithorhynchus anatinus funcgen 81 1
## 7418
                      ornithorhynchus_anatinus_funcgen_82_1
                      ornithorhynchus_anatinus_funcgen_83_1
## 7419
## 7420
                      ornithorhynchus anatinus funcgen 84 1
## 7421
                      ornithorhynchus anatinus funcgen 85 1
## 7422
                      ornithorhynchus_anatinus_funcgen_86_1
## 7423
                      ornithorhynchus_anatinus_funcgen_87_1
## 7424
                      ornithorhynchus_anatinus_funcgen_88_1
## 7425
                      ornithorhynchus_anatinus_funcgen_89_1
## 7426
                      ornithorhynchus anatinus funcgen 90 1
## 7427
                      ornithorhynchus anatinus funcgen 91 1
## 7428
                      ornithorhynchus anatinus funcgen 92 1
## 7429
                      ornithorhynchus_anatinus_funcgen_93_1
## 7430
                      ornithorhynchus anatinus funcgen 94 1
## 7431
                      ornithorhynchus_anatinus_funcgen_95_1
## 7432
                      ornithorhynchus anatinus funcgen 96 1
## 7433
                      ornithorhynchus anatinus funcgen 97 1
## 7434
                      ornithorhynchus anatinus funcgen 98 1
## 7435
                      ornithorhynchus_anatinus_funcgen_99_1
## 7436
               ornithorhynchus_anatinus_otherfeatures_48_1e
               ornithorhynchus_anatinus_otherfeatures_49_1f
## 7437
## 7438
               ornithorhynchus anatinus otherfeatures 50 1g
               ornithorhynchus anatinus otherfeatures 51 1h
## 7439
## 7440
               ornithorhynchus anatinus otherfeatures 52 1i
## 7441
               ornithorhynchus_anatinus_otherfeatures 53 1;
## 7442
               ornithorhynchus_anatinus_otherfeatures_54_1j
## 7443
               ornithorhynchus anatinus otherfeatures 55 1k
## 7444
               ornithorhynchus anatinus otherfeatures 56 1k
## 7445
               ornithorhynchus anatinus otherfeatures 57 1k
## 7446
               ornithorhynchus anatinus otherfeatures 58 11
## 7447
               ornithorhynchus anatinus otherfeatures 59 1m
## 7448
               ornithorhynchus_anatinus_otherfeatures_60_1n
## 7449
               ornithorhynchus anatinus otherfeatures 61 10
               ornithorhynchus anatinus otherfeatures 62 1p
## 7450
## 7451
                ornithorhynchus anatinus otherfeatures 63 1
## 7452
                ornithorhynchus_anatinus_otherfeatures_64_1
## 7453
                ornithorhynchus_anatinus_otherfeatures_65_1
## 7454
                ornithorhynchus_anatinus_otherfeatures_66_1
                ornithorhynchus anatinus otherfeatures 67 1
## 7455
## 7456
                ornithorhynchus anatinus otherfeatures 68 1
## 7457
                ornithorhynchus anatinus otherfeatures 69 1
## 7458
                ornithorhynchus_anatinus_otherfeatures_70_1
## 7459
                ornithorhynchus_anatinus_otherfeatures_71_1
## 7460
                ornithorhynchus_anatinus_otherfeatures_72_1
## 7461
                ornithorhynchus anatinus otherfeatures 73 1
                ornithorhynchus anatinus otherfeatures 74 1
## 7462
```

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ornithorhynchus anatinus otherfeatures 75 1
## 7463
## 7464
                ornithorhynchus anatinus otherfeatures 76 1
                ornithorhynchus anatinus otherfeatures 77 1
## 7465
## 7466
                ornithorhynchus_anatinus_otherfeatures_78_1
## 7467
                ornithorhynchus anatinus otherfeatures 79 1
## 7468
                ornithorhynchus anatinus otherfeatures 80 1
## 7469
                ornithorhynchus anatinus otherfeatures 81 1
                ornithorhynchus anatinus otherfeatures 82 1
## 7470
## 7471
                ornithorhynchus anatinus otherfeatures 83 1
## 7472
                ornithorhynchus_anatinus_otherfeatures_84_1
## 7473
                ornithorhynchus_anatinus_otherfeatures_85_1
## 7474
                ornithorhynchus_anatinus_otherfeatures_86_1
## 7475
                ornithorhynchus anatinus otherfeatures 87 1
## 7476
                ornithorhynchus_anatinus_otherfeatures_88_1
## 7477
                ornithorhynchus_anatinus_otherfeatures_89_1
## 7478
                ornithorhynchus_anatinus_otherfeatures_90_1
## 7479
                ornithorhynchus_anatinus_otherfeatures_91_1
## 7480
                ornithorhynchus anatinus otherfeatures 92 1
## 7481
                ornithorhynchus anatinus otherfeatures 93 1
## 7482
                ornithorhynchus anatinus otherfeatures 94 1
## 7483
                ornithorhynchus_anatinus_otherfeatures_95_1
## 7484
                ornithorhynchus anatinus otherfeatures 96 1
## 7485
                ornithorhynchus_anatinus_otherfeatures_97_1
## 7486
                ornithorhynchus anatinus otherfeatures 98 1
## 7487
                ornithorhynchus anatinus otherfeatures 99 1
## 7488
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## 7489
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                       ornithorhynchus anatinus rnaseg 71 1
## 7490
## 7491
                       ornithorhynchus_anatinus_rnaseq_72_1
## 7492
                       ornithorhynchus anatinus rnaseg 73 1
## 7493
                       ornithorhynchus_anatinus_rnaseq_74_1
## 7494
                       ornithorhynchus anatinus rnaseg 75 1
## 7495
                       ornithorhynchus_anatinus_rnaseq_76_1
## 7496
                       ornithorhynchus_anatinus_rnaseq_77_1
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                       ornithorhynchus anatinus rnaseg 78 1
## 7498
                       ornithorhynchus anatinus rnaseq 79 1
## 7499
                       ornithorhynchus anatinus rnaseg 80 1
## 7500
                       ornithorhynchus anatinus rnaseg 81 1
## 7501
                       ornithorhynchus anatinus rnaseg 82 1
## 7502
                       ornithorhynchus_anatinus_rnaseq_83_1
## 7503
                       ornithorhynchus anatinus rnaseq 84 1
                       ornithorhynchus anatinus rnaseq 85 1
## 7504
                       ornithorhynchus anatinus rnaseg 86 1
## 7505
## 7506
                       ornithorhynchus_anatinus_rnaseq_87_1
## 7507
                       ornithorhynchus_anatinus_rnaseq_88_1
## 7508
                       ornithorhynchus_anatinus_rnaseq_89_1
## 7509
                       ornithorhynchus anatinus rnaseg 90 1
## 7510
                       ornithorhynchus_anatinus_rnaseq_91_1
## 7511
                       ornithorhynchus_anatinus_rnaseq_92_1
                       ornithorhynchus_anatinus_rnaseq_93_1
## 7512
## 7513
                       ornithorhynchus_anatinus_rnaseq_94_1
## 7514
                       ornithorhynchus_anatinus_rnaseq_95_1
## 7515
                       ornithorhynchus anatinus rnaseq 96 1
## 7516
                       ornithorhynchus anatinus rnaseg 97 1
```

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## 7517
                       ornithorhynchus anatinus rnaseg 98 1
## 7518
                       ornithorhynchus anatinus rnaseg 99 1
## 7519
                   ornithorhynchus anatinus variation 48 1e
## 7520
                   ornithorhynchus_anatinus_variation_49_1f
## 7521
                   ornithorhynchus anatinus variation 50 1g
## 7522
                   ornithorhynchus anatinus variation 51 1h
## 7523
                   ornithorhynchus anatinus variation 52 1i
## 7524
                   ornithorhynchus anatinus variation 53 1j
## 7525
                   ornithorhynchus anatinus variation 54 1j
## 7526
                   ornithorhynchus_anatinus_variation_55_1k
## 7527
                   ornithorhynchus_anatinus_variation_56_1k
                   ornithorhynchus_anatinus_variation_57_1k
## 7528
## 7529
                   ornithorhynchus anatinus variation 58 11
## 7530
                   ornithorhynchus_anatinus_variation_59_1m
## 7531
                   ornithorhynchus_anatinus_variation_60_1n
## 7532
                   ornithorhynchus_anatinus_variation_61_1o
## 7533
                   ornithorhynchus_anatinus_variation_62_1p
## 7534
                    ornithorhynchus anatinus variation 63 1
## 7535
                    ornithorhynchus_anatinus_variation_64_1
## 7536
                    ornithorhynchus anatinus variation 65 1
## 7537
                    ornithorhynchus_anatinus_variation_66_1
## 7538
                    ornithorhynchus anatinus variation 67 1
                    ornithorhynchus_anatinus_variation_68_1
## 7539
## 7540
                    ornithorhynchus anatinus variation 69 1
## 7541
                    ornithorhynchus anatinus variation 70 1
## 7542
                    ornithorhynchus anatinus variation 71 1
## 7543
                    ornithorhynchus_anatinus_variation_72_1
                    ornithorhynchus_anatinus_variation_73_1
## 7544
## 7545
                    ornithorhynchus_anatinus_variation_74_1
## 7546
                    ornithorhynchus anatinus variation 75 1
## 7547
                    ornithorhynchus_anatinus_variation_76_1
## 7548
                    ornithorhynchus_anatinus_variation_77_1
## 7549
                    ornithorhynchus_anatinus_variation_78_1
## 7550
                    ornithorhynchus_anatinus_variation_79_1
## 7551
                    ornithorhynchus anatinus variation 80 1
## 7552
                    ornithorhynchus anatinus variation 81 1
## 7553
                    ornithorhynchus anatinus variation 82 1
## 7554
                    ornithorhynchus_anatinus_variation_83_1
## 7555
                    ornithorhynchus anatinus variation 84 1
## 7556
                    ornithorhynchus_anatinus_variation_85_1
## 7557
                    ornithorhynchus anatinus variation 86 1
## 7558
                    ornithorhynchus anatinus variation 87 1
                    ornithorhynchus anatinus variation 88 1
## 7559
## 7560
                    ornithorhynchus_anatinus_variation_89_1
## 7561
                    ornithorhynchus_anatinus_variation_90_1
## 7562
                    ornithorhynchus_anatinus_variation_91_1
## 7563
                    ornithorhynchus anatinus variation 92 1
## 7564
                    ornithorhynchus_anatinus_variation_93_1
## 7565
                    ornithorhynchus_anatinus_variation_94_1
                    ornithorhynchus_anatinus_variation_95_1
## 7566
## 7567
                    ornithorhynchus_anatinus_variation_96_1
## 7568
                    ornithorhynchus_anatinus_variation_97_1
## 7569
                    ornithorhynchus_anatinus_variation_98_1
                    ornithorhynchus anatinus variation 99 1
## 7570
```

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## 7571
                            oryctolagus cuniculus core 48 1e
## 7572
                            oryctolagus_cuniculus_core_49_1f
## 7573
                            oryctolagus cuniculus core 50 1g
## 7574
                           oryctolagus_cuniculus_core_51_1g
##
  7575
                            oryctolagus cuniculus core 52 1h
## 7576
                            oryctolagus cuniculus core 53 1h
## 7577
                            orvctolagus cuniculus core 54 1h
## 7578
                            oryctolagus cuniculus core 55 1h
  7579
                           oryctolagus cuniculus core 56 1h
## 7580
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## 7581
                           oryctolagus_cuniculus_core_58_2a
## 7582
                            oryctolagus_cuniculus_core_59_2b
##
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## 7584
                            oryctolagus_cuniculus_core_61_3a
## 7585
                            oryctolagus_cuniculus_core_62_3b
## 7586
                             oryctolagus_cuniculus_core_63_3
## 7587
                             oryctolagus_cuniculus_core_64_3
## 7588
                             oryctolagus cuniculus core 65 3
## 7589
                             oryctolagus_cuniculus_core_66_3
## 7590
                             oryctolagus cuniculus core 67 3
## 7591
                             oryctolagus_cuniculus_core_68_3
## 7592
                             oryctolagus cuniculus core 69 3
## 7593
                             oryctolagus_cuniculus_core_70_3
## 7594
                             oryctolagus cuniculus core 71 3
## 7595
                             oryctolagus cuniculus core 72 3
## 7596
                             oryctolagus cuniculus core 73 2
## 7597
                             oryctolagus_cuniculus_core_74_2
                             oryctolagus_cuniculus_core_75_2
##
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## 7599
                             oryctolagus_cuniculus_core_76_2
## 7600
                             oryctolagus cuniculus core 77 2
## 7601
                             oryctolagus_cuniculus_core_78_2
## 7602
                             oryctolagus_cuniculus_core_79_2
## 7603
                             oryctolagus_cuniculus_core_80_2
## 7604
                             oryctolagus_cuniculus_core_81_2
##
  7605
                             oryctolagus cuniculus core 82 2
                             oryctolagus_cuniculus_core_83_2
  7606
##
## 7607
                             oryctolagus cuniculus core 84 2
## 7608
                             oryctolagus_cuniculus_core_85_2
## 7609
                             oryctolagus cuniculus core 86 2
## 7610
                             oryctolagus_cuniculus_core_87_2
## 7611
                             oryctolagus cuniculus core 88 2
## 7612
                             oryctolagus cuniculus core 89 2
                             oryctolagus cuniculus core 90 2
##
  7613
## 7614
                             oryctolagus_cuniculus_core_91_2
## 7615
                             oryctolagus_cuniculus_core_92_2
## 7616
                             oryctolagus_cuniculus_core_93_2
## 7617
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## 7618
                             oryctolagus_cuniculus_core_95_2
## 7619
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                             oryctolagus_cuniculus_core_98 2
## 7621
## 7622
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## 7623
## 7624
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## 7625
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## 7627
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  7628
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                         oryctolagus cuniculus funcgen 76 2
  7630
                         oryctolagus cuniculus funcgen 77 2
##
## 7631
                         orvctolagus cuniculus funcgen 78 2
## 7632
                         oryctolagus cuniculus funcgen 79 2
                         oryctolagus_cuniculus funcgen 80 2
  7633
## 7634
                         oryctolagus_cuniculus_funcgen_81_2
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                         oryctolagus_cuniculus_funcgen_82_2
                         oryctolagus_cuniculus_funcgen_83_2
## 7636
##
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                         oryctolagus cuniculus funcgen 84 2
## 7638
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## 7639
                         oryctolagus_cuniculus_funcgen_86_2
## 7640
                         oryctolagus_cuniculus_funcgen_87_2
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                         oryctolagus cuniculus funcgen 90 2
## 7644
                         oryctolagus cuniculus funcgen 91 2
##
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  7646
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                         oryctolagus_cuniculus_funcgen_94_2
## 7647
## 7648
                         orvctolagus cuniculus funcgen 95 2
## 7649
                         oryctolagus cuniculus funcgen 96 2
## 7650
                         orvctolagus cuniculus funcgen 97 2
## 7651
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  7652
                         oryctolagus_cuniculus_funcgen_99_2
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## 7653
## 7654
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## 7655
                  oryctolagus_cuniculus_otherfeatures_59_2b
## 7656
                   oryctolagus_cuniculus_otherfeatures_60_3
## 7657
                  oryctolagus_cuniculus_otherfeatures_61 3a
## 7658
                  oryctolagus_cuniculus_otherfeatures_62_3b
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                   oryctolagus cuniculus otherfeatures 63 3
  7660
                   oryctolagus cuniculus otherfeatures 64 3
##
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                   oryctolagus cuniculus otherfeatures 65 3
## 7662
                   oryctolagus_cuniculus_otherfeatures_66_3
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                   oryctolagus cuniculus otherfeatures 67 3
## 7664
                   oryctolagus_cuniculus_otherfeatures_68_3
## 7665
                   oryctolagus cuniculus otherfeatures 69 3
## 7666
                   oryctolagus cuniculus otherfeatures 70 3
                   orvctolagus cuniculus otherfeatures 71 3
##
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## 7668
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## 7670
                   oryctolagus_cuniculus_otherfeatures_74_2
## 7671
                   oryctolagus cuniculus otherfeatures 75 2
## 7672
                   oryctolagus_cuniculus_otherfeatures_76_2
## 7673
                   oryctolagus_cuniculus_otherfeatures_77_2
## 7674
                   oryctolagus_cuniculus_otherfeatures_78_2
## 7675
                   oryctolagus_cuniculus_otherfeatures_79_2
## 7676
                   oryctolagus_cuniculus_otherfeatures_80_2
## 7677
                   oryctolagus_cuniculus_otherfeatures_81_2
## 7678
                   oryctolagus cuniculus otherfeatures 82 2
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## 7679
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                   oryctolagus cuniculus otherfeatures 85 2
## 7682
                   oryctolagus_cuniculus_otherfeatures_86_2
##
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## 7684
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## 7685
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## 7686
                   oryctolagus cuniculus otherfeatures 90 2
  7687
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                   oryctolagus_cuniculus_otherfeatures_93_2
## 7690
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##
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## 7694
                   oryctolagus_cuniculus_otherfeatures_98_2
## 7695
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                           oryctolagus_cuniculus_rnaseq_74_2
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## 7698
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## 7699
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## 7701
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## 7704
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## 7705
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##
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## 7708
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## 7709
                           oryctolagus_cuniculus_rnaseq_86_2
## 7710
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## 7711
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## 7715
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                           oryctolagus_cuniculus_rnaseq_99_2
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## 7723
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## 7724
                                  oryzias_latipes_core_49_1e
  7725
                                  oryzias_latipes_core_50_1f
## 7726
                                  oryzias_latipes_core_51_1g
## 7727
                                  oryzias_latipes_core_52_1h
## 7728
                                  oryzias_latipes_core_53_1i
                                  oryzias_latipes_core_54_1i
## 7729
## 7730
                                  oryzias_latipes_core_55_1i
## 7731
                                  oryzias_latipes_core_56_1i
## 7732
                                  oryzias latipes core 57 1i
```

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## 7733
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## 7734
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## 7735
                                  oryzias_latipes_core_60_11
## 7736
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##
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                                  oryzias_latipes_core_62_1n
## 7738
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## 7739
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## 7740
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##
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## 7746
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## 7747
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## 7748
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##
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## 7750
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## 7757
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## 7763
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## 7772
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## 7782
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## 7784
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## 7785
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## 7786
                             oryzias_latipes_hni_rnaseq_99_1
```

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## 7788
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## 7790
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                              oryzias_latipes_hsok_core_98_1
## 7792
                              oryzias latipes hsok core 99 1
## 7793
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## 7794
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## 7801
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## 7804
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  7805
##
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##
  7806
                         oryzias latipes otherfeatures 55 1i
##
  7807
                         oryzias_latipes_otherfeatures_56_1i
## 7808
                         oryzias_latipes_otherfeatures_57_1i
## 7809
                         oryzias_latipes_otherfeatures_58_1j
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                         oryzias latipes otherfeatures 60 11
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## 7838
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## 7840
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```

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                               otolemur garnettii core 61 1i
## 7883
                               otolemur garnettii core 62 1j
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## 7894
                                otolemur garnettii core 73 3
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                                otolemur garnettii core 79 3
## 7901
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## 7903
                                otolemur garnettii core 82 3
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                                otolemur_garnettii_core_85_3
## 7906
## 7907
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## 7920
                                otolemur garnettii core 99 3
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## 7923
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## 7925
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## 7932
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                       otolemur_garnettii_otherfeatures_78_3
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## 7936
                       otolemur garnettii otherfeatures 80 3
## 7937
                       otolemur garnettii otherfeatures 81 3
## 7938
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                       otolemur_garnettii_otherfeatures_85_3
## 7942
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## 7943
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## 7944
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## 7945
                       otolemur_garnettii_otherfeatures_89_3
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## 7948
                       otolemur garnettii otherfeatures 92 3
```

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## 7949
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## 7954
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## 7955
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## 7957
                                        ovis aries core 75 31
## 7958
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## 7959
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## 7961
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## 7962
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## 7963
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## 7964
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## 7965
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## 7967
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## 7968
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## 7969
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## 7970
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## 7973
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## 7974
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## 7999
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## 8001
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## 8002
                              ovis aries otherfeatures 94 31
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## 8009
                                      ovis aries rnaseq 75 31
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## 8011
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## 8012
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## 8013
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## 8014
                                      ovis_aries_rnaseq_80_31
## 8015
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## 8016
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## 8017
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## 8019
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## 8021
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## 8023
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## 8051
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## 8053
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## 8056
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## 8109
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	9232	rattus_norvegicus_core_92_6
	9233	rattus_norvegicus_core_93_6
	9234	rattus_norvegicus_core_94_6
	9235	rattus_norvegicus_core_95_6
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	9237	rattus_norvegicus_core_97_6
	9238	rattus_norvegicus_core_98_6
	9239	rattus_norvegicus_core_99_6
	9240	rattus_norvegicus_funcgen_55_34w
	9241	rattus_norvegicus_funcgen_56_34x
	9242	rattus_norvegicus_funcgen_57_34y
	9243	rattus_norvegicus_funcgen_58_34z
##	9244	rattus_norvegicus_funcgen_59_34a

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## 9246
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## 9247
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## 9280
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## 9286
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                     rattus_norvegicus_otherfeatures 50 34t
## 9287
## 9288
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## 9289
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## 9290
                     rattus_norvegicus_otherfeatures_53_34u
## 9291
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## 9293
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## 9294
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## 9295
                     rattus_norvegicus_otherfeatures_58_34z
## 9296
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## 9298
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## 9313
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## 9314
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## 9352
                               rattus norvegicus rnaseq 95 6
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## 9355
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## 9358
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## 9361
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## 9363
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                         rattus norvegicus variation 56 34x
## 9366
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## 9367
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## 9370
                         rattus norvegicus variation 61 34c
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                            rattus_norvegicus_variation_74_5
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## 9390
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                            rattus_norvegicus_variation_94 6
## 9403
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## 9406
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## 9414
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## 9415
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## 9426
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                               rhinopithecus_bieti_core_95_1
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                               rhinopithecus bieti core 96 1
                               rhinopithecus_bieti_core_97_1
## 9429
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## 9431
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## 9434
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                            rhinopithecus_bieti_funcgen_94_1
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                     rhinopithecus_bieti_otherfeatures_94_1
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## 9448
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## 9455
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## 9458
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## 9460
                           rhinopithecus roxellana core 92 1
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## 9467
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## 9469
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## 9485
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## 9496
                         saccharomyces cerevisiae core 49 1h
## 9497
                         saccharomyces cerevisiae core 50 1i
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## 9499
                         saccharomyces cerevisiae core 52 1i
## 9500
                         saccharomyces_cerevisiae_core_53_1i
## 9501
                         saccharomyces cerevisiae core 54 1i
## 9502
                         saccharomyces cerevisiae core 55 1j
                         saccharomyces cerevisiae core 56 1i
## 9503
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                         saccharomyces_cerevisiae_core_57_1j
## 9505
                         saccharomyces_cerevisiae_core_58_1j
## 9506
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## 9507
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## 9508
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## 9509
                         saccharomyces_cerevisiae_core_62_2d
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## 9511
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## 9513
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## 9514
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## 9534
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## 9539
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## 9550
                     saccharomyces cerevisiae funcgen 58 1j
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## 9552
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## 9553
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## 9554
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## 9556
                      saccharomyces cerevisiae funcgen 64 3
## 9557
                      saccharomyces cerevisiae funcgen 65 4
## 9558
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## 9568
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## 9571
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## 9575
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## 9588
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                      saccharomyces_cerevisiae_funcgen_99_4
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## 9593
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## 9594
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## 9595
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               saccharomyces cerevisiae otherfeatures 52 1i
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## 9598
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## 9600
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## 9601
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## 9604
               saccharomyces cerevisiae otherfeatures 60 2b
## 9605
               saccharomyces cerevisiae otherfeatures 61 2c
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## 10421
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## 10428
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## 10429
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## 10430
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## 10432
                        taeniopygia_guttata_variation_61_1g
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                         taeniopygia_guttata_variation_86_1
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## 10474
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## 10479
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## 10482
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## 10483
                                takifugu_rubripes_core_60_4n
## 10484
                                takifugu_rubripes_core_61_4o
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## 10486
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## 10529
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## 10535
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## 10536
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## 10540
                       takifugu rubripes otherfeatures 65 4
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## 10543
                       takifugu rubripes otherfeatures 68 4
## 10544
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## 10549
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## 10573
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## 10581
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## 10584
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## 10585
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## 10586
                                 tarsius_syrichta_core_55_1c
## 10587
                                 tarsius_syrichta_core_56_1c
## 10588
                                 tarsius_syrichta_core_57_1c
## 10589
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## 10590
                                 tarsius_syrichta_core_59_1e
## 10591
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## 10592
                                 tarsius_syrichta_core_61_1g
## 10593
                                 tarsius_syrichta_core_62_1h
## 10594
                                  tarsius syrichta core 63 1
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                                  tarsius_syrichta core 70 1
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## 10624
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## 10636
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## 10685
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                      tetraodon nigroviridis variation 67 8
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## 10765
                      tetraodon nigroviridis variation 88 8
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## 11156
                             xiphophorus maculatus core 72 1
## 11157
                            xiphophorus_maculatus_core_73_1
## 11158
                            xiphophorus maculatus core 74 1
## 11159
                             xiphophorus maculatus core 75 1
## 11160
                            xiphophorus maculatus core 76 1
## 11161
                             xiphophorus_maculatus_core_77_1
## 11162
                             xiphophorus_maculatus_core_78_1
## 11163
                             xiphophorus_maculatus_core_79_1
## 11164
                             xiphophorus_maculatus_core_80_1
## 11165
                             xiphophorus_maculatus_core_81_1
## 11166
                             xiphophorus_maculatus_core_82_1
## 11167
                             xiphophorus_maculatus_core_83_1
## 11168
                             xiphophorus_maculatus_core_84_1
## 11169
                             xiphophorus maculatus core 85 1
## 11170
                            xiphophorus_maculatus_core_86_1
## 11171
                             xiphophorus maculatus core 87 1
## 11172
                            xiphophorus_maculatus_core_88_1
## 11173
                             xiphophorus maculatus core 89 1
## 11174
                             xiphophorus_maculatus_core_90_1
## 11175
                            xiphophorus maculatus core 91 1
## 11176
                             xiphophorus maculatus core 92 1
## 11177
                             xiphophorus maculatus core 93 1
## 11178
                             xiphophorus_maculatus_core_94_5
## 11179
                             xiphophorus_maculatus_core_95_5
## 11180
                             xiphophorus_maculatus_core_96_5
## 11181
                             xiphophorus_maculatus_core_97_5
## 11182
                             xiphophorus_maculatus_core_98_5
## 11183
                            xiphophorus_maculatus_core_99_5
## 11184
                   xiphophorus_maculatus_otherfeatures_69_1
## 11185
                   xiphophorus_maculatus_otherfeatures_70_1
## 11186
                   xiphophorus_maculatus_otherfeatures_71_1
## 11187
                   xiphophorus_maculatus_otherfeatures_72_1
## 11188
                   xiphophorus maculatus otherfeatures 73 1
```

```
## 11189
                   xiphophorus maculatus otherfeatures 74 1
## 11190
                   xiphophorus maculatus otherfeatures 75 1
                   xiphophorus maculatus otherfeatures 76 1
## 11191
## 11192
                   xiphophorus_maculatus_otherfeatures_77_1
## 11193
                   xiphophorus maculatus otherfeatures 78 1
## 11194
                   xiphophorus maculatus otherfeatures 79 1
## 11195
                   xiphophorus maculatus otherfeatures 80 1
## 11196
                   xiphophorus maculatus otherfeatures 81 1
## 11197
                   xiphophorus maculatus otherfeatures 82 1
## 11198
                   xiphophorus_maculatus_otherfeatures_83_1
## 11199
                   xiphophorus_maculatus_otherfeatures_84_1
## 11200
                   xiphophorus_maculatus_otherfeatures_85_1
## 11201
                   xiphophorus maculatus otherfeatures 86 1
## 11202
                   xiphophorus_maculatus_otherfeatures_87_1
## 11203
                   xiphophorus_maculatus_otherfeatures_88_1
## 11204
                   xiphophorus_maculatus_otherfeatures_89_1
## 11205
                   xiphophorus_maculatus_otherfeatures_90_1
## 11206
                   xiphophorus maculatus otherfeatures 91 1
## 11207
                   xiphophorus maculatus otherfeatures 92 1
## 11208
                   xiphophorus maculatus otherfeatures 93 1
## 11209
                   xiphophorus_maculatus_otherfeatures_94_5
## 11210
                   xiphophorus maculatus otherfeatures 95 5
## 11211
                   xiphophorus_maculatus_otherfeatures_96_5
## 11212
                   xiphophorus maculatus otherfeatures 97 5
## 11213
                   xiphophorus maculatus otherfeatures 98 5
## 11214
                   xiphophorus maculatus otherfeatures 99 5
## 11215
                          xiphophorus_maculatus_rnaseq_69_1
## 11216
                          xiphophorus_maculatus_rnaseq_70_1
## 11217
                          xiphophorus_maculatus_rnaseq_71_1
## 11218
                          xiphophorus maculatus rnaseg 72 1
## 11219
                          xiphophorus_maculatus_rnaseq_73_1
## 11220
                          xiphophorus_maculatus_rnaseq_74_1
## 11221
                          xiphophorus_maculatus_rnaseq_75_1
## 11222
                          xiphophorus_maculatus_rnaseq_76_1
## 11223
                          xiphophorus maculatus rnaseg 77 1
## 11224
                          xiphophorus maculatus rnaseq 78 1
## 11225
                          xiphophorus maculatus rnaseq 79 1
## 11226
                          xiphophorus_maculatus_rnaseq_80_1
## 11227
                          xiphophorus maculatus rnaseq 81 1
## 11228
                          xiphophorus_maculatus_rnaseq_82_1
## 11229
                          xiphophorus maculatus rnaseq 83 1
## 11230
                          xiphophorus maculatus rnaseq 84 1
## 11231
                          xiphophorus maculatus rnaseq 85 1
## 11232
                          xiphophorus_maculatus_rnaseq_86_1
## 11233
                          xiphophorus_maculatus_rnaseq_87_1
## 11234
                          xiphophorus_maculatus_rnaseq_88_1
## 11235
                          xiphophorus maculatus rnaseg 89 1
## 11236
                          xiphophorus_maculatus_rnaseq_90_1
## 11237
                          xiphophorus_maculatus_rnaseq_91_1
## 11238
                          xiphophorus_maculatus_rnaseq_92_1
## 11239
                          xiphophorus_maculatus_rnaseq_93_1
## 11240
                          xiphophorus_maculatus_rnaseq_94_5
## 11241
                          xiphophorus maculatus rnaseq 95 5
## 11242
                          xiphophorus maculatus rnaseq 96 5
```

```
## 11243
                          xiphophorus maculatus rnaseg 97 5
## 11244
                          xiphophorus_maculatus_rnaseq_98_5
## 11245
                          xiphophorus maculatus rnaseq 99 5
## 11246
                         zonotrichia_albicollis_core_96_101
## 11247
                         zonotrichia_albicollis_core_97_101
## 11248
                         zonotrichia albicollis core 98 101
                         zonotrichia albicollis core 99 101
## 11249
## 11250
                zonotrichia_albicollis_otherfeatures_96_101
## 11251
                zonotrichia_albicollis_otherfeatures_97_101
## 11252
                zonotrichia_albicollis_otherfeatures_98_101
## 11253
                zonotrichia_albicollis_otherfeatures_99_101
## 11254
                       zonotrichia_albicollis_rnaseq_96_101
## 11255
                       zonotrichia_albicollis_rnaseq_97_101
## 11256
                       zonotrichia_albicollis_rnaseq_98_101
## 11257
                       zonotrichia_albicollis_rnaseq_99_101
```

Podemos ahora trabajar para encontrar las tablas que contengan el string "compara"

```
dbGetQuery(ensembl_con, "SHOW DATABASES") %>%
  # filter the table for cases where the the string "compara" is present
  filter(str_detect(Database, "compara")) %>%
  tail()
```

```
## Database
## 47 ensembl_compara_94
## 48 ensembl_compara_95
## 49 ensembl_compara_96
## 50 ensembl_compara_97
## 51 ensembl_compara_98
## 52 ensembl_compara_99
```

Nos quedamos con la mas reciente:

Para ver las tablas que hay en la base de datos que hemos elegido:

```
library(dbplyr)
```

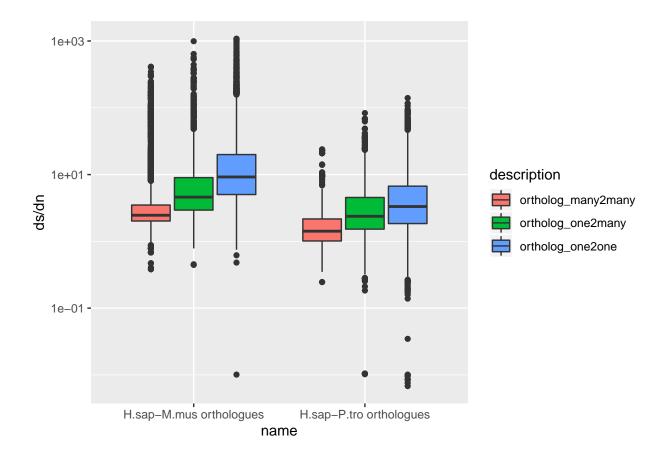
```
## Warning: package 'dbplyr' was built under R version 3.6.2
##
## Attaching package: 'dbplyr'
## The following objects are masked from 'package:dplyr':
##
## ident, sql
src_dbi(compara_con)
```

```
## src: mysql [anonymous@ensembldb.ensembl.org:NA/ensembl_compara_99]
## tbls: CAFE_gene_family, CAFE_species_gene, conservation_score,
## constrained_element, dnafrag, dnafrag_region, exon_boundaries,
## external_db, family, family_member, gene_align, gene_align_member,
```

```
##
     gene_member, gene_member_hom_stats, gene_member_qc, gene_tree_node,
##
     gene_tree_node_attr, gene_tree_node_tag, gene_tree_object_store,
     gene_tree_root, gene_tree_root_attr, gene_tree_root_tag, genome_db,
##
##
     genomic_align, genomic_align_block, genomic_align_tree, hmm_annot,
##
     hmm_curated_annot, hmm_profile, homology, homology_member,
##
    mapping session, member xref, meta, method link,
##
    method link species set, method link species set attr,
     method_link_species_set_tag, ncbi_taxa_name, ncbi_taxa_node,
##
##
     other_member_sequence, peptide_align_feature, seq_member,
##
     seq_member_projection, seq_member_projection_stable_id, sequence,
##
     species_set, species_set_header, species_set_tag, species_tree_node,
##
     species_tree_node_attr, species_tree_node_tag, species_tree_root,
     stable_id_history, synteny_region
Accedemos a la tabla "homology
compara_homology <- tbl(compara_con, "homology")</pre>
head(compara_homology)
## # Source:
               lazy query [?? x 15]
## # Database: mysql [anonymous@ensembldb.ensembl.org:NA/ensembl_compara_99]
    homology_id method_link_spe~ description is_tree_complia~
##
##
           <int>
                           <int> <chr>
                                                         <int> <dbl> <dbl>
## 1
                           132551 other_para~
                                                             1 1.02 16.6
              1
## 2
              2
                          132551 other para~
                                                             1 1.07
                                                                      7.82
## 3
               3
                           132551 other_para~
                                                             1 1.07 66.7
## 4
               4
                           132551 other para~
                                                             1 1.20 70.1
## 5
               5
                           132551 other para~
                                                             1 1.04 71.9
               6
                                                             1 0.914 77.6
                           132551 other para~
## # ... with 9 more variables: n <dbl>, s <dbl>, lnl <dbl>,
       species_tree_node_id <int>, gene_tree_node_id <int>,
## #
       gene_tree_root_id <int>, goc_score <int>, wga_coverage <dbl>,
       is_high_confidence <int>
compara_linksp <- tbl(compara_con, "method_link_species_set")</pre>
head(compara linksp)
              lazy query [?? x 8]
## # Source:
## # Database: mysql [anonymous@ensembldb.ensembl.org:NA/ensembl_compara_99]
    method_link_spe~ method_link_id species_set_id name    source url
##
                               <int>
                <int>
                                              <int> <chr> <chr> <chr>
                                              34144 G.ac~ ensem~ ""
## 1
                  555
                                  16
                  569
                                              34350 C.in~ ensem~ ""
## 2
                                  16
                  587
                                              34701 M.mu~ ensem~ ""
## 3
                                  16
## 4
                  592
                                  16
                                              34706 M.mu~ ensem~ ""
                  594
                                              34708 C.fa~ ensem~ ""
## 5
                                  16
                  596
                                  16
                                              34710 A.ca~ ensem~ ""
## # ... with 2 more variables: first release <int>, last release <int>
sapiens_ortho <- compara_linksp %>%
 filter(name == "H.sap-P.tro orthologues" | name == "H.sap-M.mus orthologues")
sapiens_ortho
## # Source:
               lazy query [?? x 8]
## # Database: mysql [anonymous@ensembldb.ensembl.org:NA/ensembl_compara_99]
    method_link_spe~ method_link_id species_set_id name source url
```

```
##
                <int>
                                <int>
                                               <int> <chr> <chr> <chr>
## 1
                50976
                                 201
                                               35673 H.sa~ ensem~ ""
               103890
                                               39100 H.sa~ ensem~ ""
## 2
                                 201
## # ... with 2 more variables: first_release <int>, last_release <int>
compara_result <- sapiens_ortho %>%
  left_join(compara_homology, by = "method_link_species_set_id") %>%
  collect()
head(compara_result)
## # A tibble: 6 x 22
     method_link_spe~ method_link_id species_set_id name source url
                               <int>
##
                <int>
                                               <int> <chr> <chr> <chr>
## 1
                50976
                                  201
                                               35673 H.sa~ ensem~ ""
## 2
                                  201
                                               35673 H.sa~ ensem~ ""
                50976
## 3
                50976
                                  201
                                               35673 H.sa~ ensem~ ""
## 4
                50976
                                  201
                                               35673 H.sa~ ensem~ ""
## 5
                                               35673 H.sa~ ensem~ ""
                50976
                                  201
                                               35673 H.sa~ ensem~ ""
## 6
                50976
                                  201
## # ... with 16 more variables: first_release <int>, last_release <int>,
## #
       homology_id <int>, description <chr>, is_tree_compliant <int>,
## #
       dn <dbl>, ds <dbl>, n <dbl>, s <dbl>, lnl <dbl>,
## #
       species_tree_node_id <int>, gene_tree_node_id <int>,
       gene_tree_root_id <int>, goc_score <int>, wga_coverage <dbl>,
## #
       is_high_confidence <int>
compara_result %>%
  count(name, description)
## # A tibble: 6 x 3
    name
##
                             description
                                                     n
     <chr>
                             <chr>
                                                 <int>
## 1 H.sap-M.mus orthologues ortholog_many2many
                                                  5773
## 2 H.sap-M.mus orthologues ortholog_one2many
                                                  3450
## 3 H.sap-M.mus orthologues ortholog_one2one
                                                 16534
## 4 H.sap-P.tro orthologues ortholog_many2many
                                                   556
## 5 H.sap-P.tro orthologues ortholog_one2many
                                                  3214
## 6 H.sap-P.tro orthologues ortholog_one2one
                                                 19410
compara_result %>%
  ggplot(aes(name, ds/dn, fill = description)) +
  geom_boxplot() + scale_y_log10()
## Warning: Transformation introduced infinite values in continuous y-axis
```

Warning: Removed 10524 rows containing non-finite values (stat boxplot).



Datos estructurados: APIs y paquetes dedicados

La base de datos del Worl Bank contiene una API desde la que podemos acceder a los World Data Indicators (http://datatopics.worldbank.org/world-development-indicators/). El paquete de R "WDI" baja los datos que le pidamos de esa base de datos, parsea el XML correspondiente y formatea el objeto resultante. Ademas contiene funciones para hacer busquedas en las bases de datos:

```
library(WDI)

WDIsearch(string = "life.*expectancy", field = "name", cache = NULL)

## indicator
```

```
[1,] "UIS.SLE.12"
##
##
    [2,] "UIS.SLE.1.M"
    [3,] "UIS.SLE.1.GPI"
##
    [4,] "UIS.SLE.1.F"
##
    [5,] "UIS.SLE.1"
##
    [6,] "UIS.SLE.O.M"
##
##
    [7,] "UIS.SLE.O.GPI"
##
    [8,] "UIS.SLE.O.F"
    [9,] "UIS.SLE.O"
##
##
   [10,] "UIS.SLEN.12.T"
  [11,] "UIS.SLEN.12.M"
  [12,] "UIS.SLEN.12.GPI"
## [13,] "UIS.SLEN.12.F"
## [14,] "UIS.SLE.56.M"
## [15,] "UIS.SLE.56.GPI"
```

```
## [16,] "UIS.SLE.56.F"
## [17,] "UIS.SLE.56"
## [18,] "UIS.SLE.4.M"
## [19,] "UIS.SLE.4.GPI"
## [20,] "UIS.SLE.4.F"
## [21,] "UIS.SLE.4"
## [22.] "UIS.SLE.23.M"
## [23,] "UIS.SLE.23.GPI"
## [24,] "UIS.SLE.23.F"
## [25,] "UIS.SLE.23"
## [26,] "UIS.SLE.1t6.GPI"
## [27,] "UIS.SLE.123.M"
## [28,] "UIS.SLE.123.GPI"
## [29,] "UIS.SLE.123.F"
## [30,] "UIS.SLE.123"
## [31,] "UIS.SLE.12.M"
## [32,] "UIS.SLE.12.F"
## [33,] "SP.DYN.LE60.MA.IN"
## [34,] "SP.DYN.LE60.FE.IN"
## [35,] "SP.DYN.LEOO.MA.IN"
## [36,] "SP.DYN.LEOO.IN"
## [37,] "SP.DYN.LEOO.FE.IN"
## [38,] "SE.SCH.LIFE.MA"
## [39,] "SE.SCH.LIFE.FE"
## [40,] "SE.SCH.LIFE"
         name
##
  [1,] "School life expectancy, primary and lower secondary, both sexes (years)"
   [2,] "School life expectancy, primary, male (years)"
## [3,] "School life expectancy, primary, gender parity index (GPI)"
## [4,] "School life expectancy, primary, female (years)"
## [5,] "School life expectancy, primary, both sexes (years)"
## [6,] "School life expectancy, pre-primary, male (years)"
## [7,] "School life expectancy, pre-primary, gender parity index (GPI)"
## [8,] "School life expectancy, pre-primary, female (years)"
   [9,] "School life expectancy, pre-primary, both sexes (years)"
## [10,] "School life expectancy, primary and lower secondary (excluding repetition), both sexes (years
## [11,] "School life expectancy, primary and lower secondary (excluding repetition), male (years)"
## [12,] "School life expectancy, primary and lower secondary (excluding repetition), gender parity ind
## [13,] "School life expectancy, primary and lower secondary (excluding repetition), female (years)"
## [14,] "School life expectancy, tertiary, male (years)"
## [15,] "School life expectancy, tertiary, gender parity index (GPI)"
## [16,] "School life expectancy, tertiary, female (years)"
## [17,] "School life expectancy, tertiary, both sexes (years)"
## [18,] "School life expectancy, post-secondary non-tertiary, male (years)"
## [19,] "School life expectancy, post-secondary non-tertiary, gender parity index (GPI)"
## [20,] "School life expectancy, post-secondary non-tertiary, female (years)"
## [21,] "School life expectancy, post-secondary non-tertiary, both sexes (years)"
## [22,] "School life expectancy, secondary, male (years)"
## [23,] "School life expectancy, secondary, gender parity index (GPI)"
## [24,] "School life expectancy, secondary, female (years)"
## [25,] "School life expectancy, secondary, both sexes (years)"
## [26,] "School life expectancy, primary to tertiary, gender parity index (GPI)"
## [27,] "School life expectancy, primary and secondary, male (years)"
## [28,] "School life expectancy, primary and secondary, gender parity index (GPI)"
```

```
## [29,] "School life expectancy, primary and secondary, female (years)"
## [30,] "School life expectancy, primary and secondary, both sexes (years)"
## [31,] "School life expectancy, primary and lower secondary, male (years)"
## [32,] "School life expectancy, primary and lower secondary, female (years)"
## [33,] "Life expectancy at age 60, male (years)"
## [34,] "Life expectancy at age 60, female (years)"
## [35,] "Life expectancy at birth, male (years)"
## [36,] "Life expectancy at birth, total (years)"
## [37,] "Life expectancy at birth, female (years)"
## [38,] "School life expectancy, primary to tertiary, male (years)"
## [39,] "School life expectancy, primary to tertiary, female (years)"
## [40,] "School life expectancy, primary to tertiary, both sexes (years)"
df.le = WDI(country = "all", indicator = c("SP.DYN.LE00.IN"), start = 1900,
            end = 2012)
head(df.le)
##
     iso2c
              country SP.DYN.LEOO.IN year
## 1
        1A Arab World
                            70.70383 2012
## 2
        1A Arab World
                            70.52877 2011
## 3
                            70.34992 2010
        1A Arab World
## 4
        1A Arab World
                            70.15756 2009
## 5
        1A Arab World
                            69.94217 2008
                            69.70155 2007
        1A Arab World
is.data.frame(df.le)
## [1] TRUE
levels(factor(df.le$country))
##
     [1] "Afghanistan"
##
     [2] "Albania"
##
     [3] "Algeria"
##
     [4] "American Samoa"
##
     [5] "Andorra"
##
     [6] "Angola"
##
     [7] "Antigua and Barbuda"
##
     [8] "Arab World"
##
     [9] "Argentina"
##
    [10] "Armenia"
##
   [11] "Aruba"
   [12] "Australia"
   [13] "Austria"
##
    [14] "Azerbaijan"
##
   [15] "Bahamas, The"
   [16] "Bahrain"
   [17] "Bangladesh"
##
   [18] "Barbados"
   [19] "Belarus"
##
##
   [20] "Belgium"
##
   [21] "Belize"
##
   [22] "Benin"
   [23] "Bermuda"
##
##
  [24] "Bhutan"
## [25] "Bolivia"
```

```
[26] "Bosnia and Herzegovina"
##
    [27] "Botswana"
##
    [28] "Brazil"
    [29] "British Virgin Islands"
##
##
    [30] "Brunei Darussalam"
    [31] "Bulgaria"
##
    [32] "Burkina Faso"
##
    [33] "Burundi"
##
##
    [34] "Cabo Verde"
    [35] "Cambodia"
##
    [36] "Cameroon"
    [37] "Canada"
##
    [38] "Caribbean small states"
##
##
    [39] "Cayman Islands"
##
    [40] "Central African Republic"
##
    [41] "Central Europe and the Baltics"
##
    [42] "Chad"
##
    [43] "Channel Islands"
##
   [44] "Chile"
    [45] "China"
##
##
    [46] "Colombia"
    [47] "Comoros"
##
    [48] "Congo, Dem. Rep."
##
    [49] "Congo, Rep."
##
    [50] "Costa Rica"
##
    [51] "Cote d'Ivoire"
##
    [52] "Croatia"
    [53] "Cuba"
##
##
    [54] "Curacao"
##
    [55] "Cyprus"
##
    [56] "Czech Republic"
##
    [57] "Denmark"
##
    [58] "Djibouti"
    [59] "Dominica"
##
##
    [60] "Dominican Republic"
    [61] "Early-demographic dividend"
##
##
    [62] "East Asia & Pacific"
##
    [63] "East Asia & Pacific (excluding high income)"
##
    [64] "East Asia & Pacific (IDA & IBRD countries)"
    [65] "Ecuador"
##
    [66] "Egypt, Arab Rep."
##
    [67] "El Salvador"
    [68] "Equatorial Guinea"
##
    [69] "Eritrea"
##
    [70] "Estonia"
##
    [71] "Eswatini"
##
    [72] "Ethiopia"
##
##
    [73] "Euro area"
    [74] "Europe & Central Asia"
    [75] "Europe & Central Asia (excluding high income)"
##
    [76] "Europe & Central Asia (IDA & IBRD countries)"
##
##
   [77] "European Union"
##
   [78] "Faroe Islands"
##
   [79] "Fiji"
```

```
## [80] "Finland"
##
  [81] "Fragile and conflict affected situations"
  [82] "France"
  [83] "French Polynesia"
##
   [84] "Gabon"
##
  [85] "Gambia, The"
  [86] "Georgia"
   [87] "Germany"
##
   [88] "Ghana"
##
  [89] "Gibraltar"
  [90] "Greece"
  [91] "Greenland"
##
  [92] "Grenada"
## [93] "Guam"
## [94] "Guatemala"
##
   [95] "Guinea"
## [96] "Guinea-Bissau"
## [97] "Guyana"
## [98] "Haiti"
## [99] "Heavily indebted poor countries (HIPC)"
## [100] "High income"
## [101] "Honduras"
## [102] "Hong Kong SAR, China"
## [103] "Hungary"
## [104] "IBRD only"
## [105] "Iceland"
## [106] "IDA & IBRD total"
## [107] "IDA blend"
## [108] "IDA only"
## [109] "IDA total"
## [110] "India"
## [111] "Indonesia"
## [112] "Iran, Islamic Rep."
## [113] "Iraq"
## [114] "Ireland"
## [115] "Isle of Man"
## [116] "Israel"
## [117] "Italy"
## [118] "Jamaica"
## [119] "Japan"
## [120] "Jordan"
## [121] "Kazakhstan"
## [122] "Kenya"
## [123] "Kiribati"
## [124] "Korea, Dem. Peopleâ\200\231s Rep."
## [125] "Korea, Rep."
## [126] "Kosovo"
## [127] "Kuwait"
## [128] "Kyrgyz Republic"
## [129] "Lao PDR"
## [130] "Late-demographic dividend"
## [131] "Latin America & Caribbean"
## [132] "Latin America & Caribbean (excluding high income)"
## [133] "Latin America & the Caribbean (IDA & IBRD countries)"
```

```
## [134] "Latvia"
## [135] "Least developed countries: UN classification"
## [136] "Lebanon"
## [137] "Lesotho"
## [138] "Liberia"
## [139] "Libya"
## [140] "Liechtenstein"
## [141] "Lithuania"
## [142] "Low & middle income"
## [143] "Low income"
## [144] "Lower middle income"
## [145] "Luxembourg"
## [146] "Macao SAR, China"
## [147] "Madagascar"
## [148] "Malawi"
## [149] "Malaysia"
## [150] "Maldives"
## [151] "Mali"
## [152] "Malta"
## [153] "Marshall Islands"
## [154] "Mauritania"
## [155] "Mauritius"
## [156] "Mexico"
## [157] "Micronesia, Fed. Sts."
## [158] "Middle East & North Africa"
## [159] "Middle East & North Africa (excluding high income)"
## [160] "Middle East & North Africa (IDA & IBRD countries)"
## [161] "Middle income"
## [162] "Moldova"
## [163] "Monaco"
## [164] "Mongolia"
## [165] "Montenegro"
## [166] "Morocco"
## [167] "Mozambique"
## [168] "Myanmar"
## [169] "Namibia"
## [170] "Nauru"
## [171] "Nepal"
## [172] "Netherlands"
## [173] "New Caledonia"
## [174] "New Zealand"
## [175] "Nicaragua"
## [176] "Niger"
## [177] "Nigeria"
## [178] "North America"
## [179] "North Macedonia"
## [180] "Northern Mariana Islands"
## [181] "Norway"
## [182] "Not classified"
## [183] "OECD members"
## [184] "Oman"
## [185] "Other small states"
## [186] "Pacific island small states"
```

[187] "Pakistan"

```
## [188] "Palau"
## [189] "Panama"
## [190] "Papua New Guinea"
## [191] "Paraguay"
## [192] "Peru"
## [193] "Philippines"
## [194] "Poland"
## [195] "Portugal"
## [196] "Post-demographic dividend"
## [197] "Pre-demographic dividend"
## [198] "Puerto Rico"
## [199] "Qatar"
## [200] "Romania"
## [201] "Russian Federation"
## [202] "Rwanda"
## [203] "Samoa"
## [204] "San Marino"
## [205] "Sao Tome and Principe"
## [206] "Saudi Arabia"
## [207] "Senegal"
## [208] "Serbia"
## [209] "Seychelles"
## [210] "Sierra Leone"
## [211] "Singapore"
## [212] "Sint Maarten (Dutch part)"
## [213] "Slovak Republic"
## [214] "Slovenia"
## [215] "Small states"
## [216] "Solomon Islands"
## [217] "Somalia"
## [218] "South Africa"
## [219] "South Asia"
## [220] "South Asia (IDA & IBRD)"
## [221] "South Sudan"
## [222] "Spain"
## [223] "Sri Lanka"
## [224] "St. Kitts and Nevis"
## [225] "St. Lucia"
## [226] "St. Martin (French part)"
## [227] "St. Vincent and the Grenadines"
## [228] "Sub-Saharan Africa"
## [229] "Sub-Saharan Africa (excluding high income)"
## [230] "Sub-Saharan Africa (IDA & IBRD countries)"
## [231] "Sudan"
## [232] "Suriname"
## [233] "Sweden"
```

[240] "Togo" ## [241] "Tonga"

[234] "Switzerland"

[236] "Tajikistan" ## [237] "Tanzania" ## [238] "Thailand" ## [239] "Timor-Leste"

[235] "Syrian Arab Republic"

```
## [242] "Trinidad and Tobago"
## [243] "Tunisia"
## [244] "Turkey"
## [245] "Turkmenistan"
## [246] "Turks and Caicos Islands"
## [247] "Tuvalu"
## [248] "Uganda"
## [249] "Ukraine"
## [250] "United Arab Emirates"
## [251] "United Kingdom"
## [252] "United States"
## [253] "Upper middle income"
## [254] "Uruguay"
## [255] "Uzbekistan"
## [256] "Vanuatu"
## [257] "Venezuela, RB"
## [258] "Vietnam"
## [259] "Virgin Islands (U.S.)"
## [260] "West Bank and Gaza"
## [261] "World"
## [262] "Yemen, Rep."
## [263] "Zambia"
## [264] "Zimbabwe"
levels(factor(df.le$year))
   [1] "1960" "1961" "1962" "1963" "1964" "1965" "1966" "1967" "1968" "1969"
## [11] "1970" "1971" "1972" "1973" "1974" "1975" "1976" "1977" "1978" "1979"
## [21] "1980" "1981" "1982" "1983" "1984" "1985" "1986" "1987" "1988" "1989"
## [31] "1990" "1991" "1992" "1993" "1994" "1995" "1996" "1997" "1998" "1999"
## [41] "2000" "2001" "2002" "2003" "2004" "2005" "2006" "2007" "2008" "2009"
## [51] "2010" "2011" "2012"
```

Datos no estructurados:

Datos de un link de internet

El paquete dslab está en github, por lo que podriamos descargarnos los datos del fichero "murders.csv" directamente de alli.

```
url <- "https://raw.githubusercontent.com/rafalab/dslabs/master/inst/extdata/murders.csv"</pre>
```

Y, aún mejor, read_csv puede leerlo directamente

```
dat <- read_csv(url)</pre>
```

Podemos no solo leer los datos en R sino bajarnos el fichero a una unidad de disco o de red utilizando R:

```
download.file(url, "murders.csv")
```

Web scratching

Queremos comparar las estadisticas de asesinato en EEUU con las estadisticas de europa, pero el dato de eurostat no esta desglosado y ese número no es comparable. En wikipedia buscamos las estadisticas de muertes no casuales a nivel mundial (CTRL+U para ver el código de la web)

```
url="https://en.wikipedia.org/wiki/List_of_countries_by_intentional_homicide_rate"
h <- read_html(url)</pre>
```

```
class(h)
## [1] "xml_document" "xml_node"
## {html_document}
## <html class="client-nojs" lang="en" dir="ltr">
## [1] <head>\n<meta http-equiv="Content-Type" content="text/html; charset= ...</pre>
## [2] <body class="mediawiki ltr sitedir-ltr mw-hide-empty-elt ns-0 ns-sub ...
Hay una seccion del codigo html que comienza con
. Ahi estan los datos de la tabla que queremos leer en R.
tab <- h %>% html nodes("table")
tab <- tab[[4]] %>% html_table
head(tab)
##
     Country (or dependent territory, subnational area, etc.) Region
## 1
                                                      Burundi Africa
## 2
                                                      Comoros Africa
## 3
                                                     Djibouti Africa
## 4
                                                      Eritrea Africa
## 5
                                                     Ethiopia Africa
## 6
                                                        Kenya Africa
##
          Subregion Rate Count Yearlisted
                                                 Source
## 1 Eastern Africa 6.02
                           635
                                      2016
                                                CTS/SDG
## 2 Eastern Africa 7.70
                                     2015 WHO Estimate
                            60
## 3 Eastern Africa 6.48
                            60
                                     2015 WHO Estimate
## 4 Eastern Africa 8.04
                           390
                                      2015 WHO Estimate
## 5 Eastern Africa 7.56 7,552
                                      2015 WHO Estimate
## 6 Eastern Africa 5.00 2,466
                                      2017
                                                    CTS
class(tab)
## [1] "data.frame"
tab <- tab %>%
  select(starts_with("Country"),
         Region,Count,Rate,starts_with("Year")) %>%
  setNames(c("country", "continent", "total", "murder_rate", "year"))
head(tab)
      country continent total murder_rate year
## 1 Burundi
              Africa 635
                                     6.02 2016
## 2 Comoros
              Africa
                          60
                                     7.70 2015
## 3 Djibouti
              Africa
                           60
                                     6.48 2015
                 Africa 390
## 4 Eritrea
                                     8.04 2015
## 5 Ethiopia
                 Africa 7,552
                                     7.56 2015
## 6
       Kenya
                 Africa 2,466
                                     5.00 2017
###Import from JSON
library(jsonlite)
```

Attaching package: 'jsonlite'

```
## The following object is masked from 'package:purrr':
##
##
citi_bike <- fromJSON("http://citibikenyc.com/stations/json")</pre>
citi_bike$executionTime
## [1] "2020-01-24 05:34:46 AM"
head(citi_bike$stationBeanList)
##
                            stationName availableDocks totalDocks latitude
       id
## 1
      304
                  Broadway & Battery Pl
                                                     18
                                                                 33 40.70463
## 2 3255
                        8 Ave & W 31 St
                                                                19 40.75059
                                                     16
                       W 52 St & 11 Ave
                                                                55 40.76727
## 3
       72
                                                     33
## 4
       79
               Franklin St & W Broadway
                                                     32
                                                                33 40.71912
                                                      7
                                                                27 40.71117
## 5
       82
                 St James Pl & Pearl St
       83 Atlantic Ave & Fort Greene Pl
                                                     25
## 6
                                                                62 40.68383
##
     longitude statusValue statusKey availableBikes
## 1 -74.01362 In Service
                                   1
## 2 -73.99468 In Service
                                   1
                                                   3
## 3 -73.99393 In Service
                                    1
                                                  21
## 4 -74.00667 In Service
                                    1
                                                   1
## 5 -74.00017 In Service
                                                  20
## 6 -73.97632 In Service
                                                  37
                                    1
                        stAddress1 stAddress2 city postalCode location
## 1
             Broadway & Battery Pl
## 2
                   8 Ave & W 31 St
## 3
                  W 52 St & 11 Ave
## 4
          Franklin St & W Broadway
## 5
            St James Pl & Pearl St
## 6 Atlantic Ave & Fort Greene Pl
##
     altitude testStation lastCommunicationTime landMark
                    FALSE 2020-01-24 05:32:48 AM
## 1
## 2
                    FALSE 2020-01-24 05:34:16 AM
## 3
                    FALSE 2020-01-24 05:33:40 AM
## 4
                    FALSE 2020-01-24 05:32:51 AM
## 5
                    FALSE 2020-01-24 05:32:43 AM
## 6
                    FALSE 2020-01-24 05:32:08 AM
```