1.0.1.8 – Introduction to Perl





Introduction to Perl Session 5

while loop

- I/O

printf and sprintf

G E N O

a 🖻



Hash Recap

hash variables are prefixed by %

```
%fruits = ( red=>"apple", green=>"grape", yellow=>"lemon" );
for $color (keys %fruits) {
    print qq($fruits{$color} is a $color fruit);
}
for $fruit (values %fruits) {
    print qq(For lunch, I brought a $fruit);
}
print "There is a blue fruit" if exists $fruits{blue};
$fruits{purple} = "eggplant";
print "Liar, $fruits{purple} is not a fruit" if exists $fruits{purple};
```

- hashes are used when you need to associate two values together (key and value)

- you want to keep a count (or other statistic) associated with a key
- store information centrally that can be recalled by using a string \$options{margin}



Sort Recap

- sorting is performed on a list
 - an optional CODE block specifies how the sort is to be done

```
@nums = (1..100);
@names = qw(spot fuzz mug flop mac chew wagger);
# default sort is ascending asciibetic
for $name (sort @names) { ... }
# descending asciibetic
for $name (sort {$b cmp $a} @names) { ... }
# ascending numerical
for $num (sort {$a <=> $b} @num) { ... }
# descending numerical
for $num (sort {$b <=> $a} @num) { ... }
# complex - sort by length of name, ascending
for $name (sort {length($a) <=> length($b)} @names) { ... }
```



Shorthand and *crement Operators

 several operations are so frequently performed, that most languages have shorthand versions

• man perlop

- do not confuse the match operator ~= here



while Loop

- we've already seen the for loop
 - iterates over items in a list

```
for $num ( @nums ) { ... }
for $key ( keys %hash ) { ... }
for $value ( sort {$b cmp $a} values %hash ) { ... }
for $word ( split(" ",$string) ) { ... }
for $char ( split("",$string) ) { ... }
```

- we now turn to the while loop, which is an iterated if

- while iterates as long as a specified condition is true





while Loop

-while can be used in the same way as for, but this practise is not encouraged

- I include it here because it nicely illustrates how while works

```
$count = 0;
while ( $count < 5 ) {
    print qq($count);
    # alter the variable used in condition
    $count++;
}
0
1
2
3
4
```

```
# this achieves the same result
for $count (0..4) {
   print qq($count);
}
0
1
2
3
4
```



while Loop

 while loop is used in cases when the appearance of the end-condition cannot be easily predicted

```
(\$sum, \$count) = (0, 0);
while (\$sum < 5) {
  # alter the variable used in condition
  $sum += rand();
  $count++;
  print qq(Sum of $count numbers is $sum);
Sum of 1 numbers is 0.520913321059197
Sum of 2 numbers is 0.644604011438787
Sum of 3 numbers is 1.22890125820413
Sum of 4 numbers is 2.09695924632251
Sum of 5 numbers is 2.73787972517312
Sum of 6 numbers is 3.21515524713323
Sum of 7 numbers is 4.19210437964648
Sum of 8 numbers is 4.62139639491215
Sum of 9 numbers is 5.02754724724218
next check of $sum < 5 fails and while loop is not repeated</pre>
```



Loop Control

- you can skip to the next loop iteration without executing the rest of the block using next
 - next if CONDITION;
- you can force termination of the loop using last
 - last if CONDITION;

```
($sum,$count) = (0,0);
while ( $sum < 5 ) {
    $x = rand();
    # skip numbers smaller than 0.5
    next if $x < 0.5;
    $sum += $x;
    $count++;
    print qq(Sum of $count numbers from [0.5,1\) is $sum);
}
Sum of 1 numbers from [0.5,1) is 0.830018592532724
Sum of 2 numbers from [0.5,1) is 1.60385857429355
Sum of 3 numbers from [0.5,1) is 2.58709897752851
...
```



```
while with shift or pop
```

you can destructively iterate across an array

- by using shift or pop, iteratively whittle an array until there are no elements left

```
@nums = (1..5);
# shift is inside while loop
while ( @nums ) {
   $num = shift @nums;
   print qq(Found $num at front of @nums);
}
Found 1 in front of 2 3 4 5
Found 2 in front of 3 4 5
Found 2 in front of 4 5
Found 3 in front of 4 5
Found 4 in front of 5
Found 5 in front of
```

```
@nums = (1..5);
# shift/assignment in condition
while ( $num = shift @nums ) {
    print qq(Found $num at front of @nums);
  }
Found 1 in front of 2 3 4 5
Found 2 in front of 3 4 5
Found 2 in front of 3 4 5
Found 3 in front of 4 5
Found 4 in front of 5
Found 5 in front of
```



challenge

- what does this code print?

```
@nums = (0..5);
while ( $num = shift @nums ) {
    print qq(Found $num at front of @nums);
}
```



challenge - answer

- the code prints nothing
 - the first shifted element is 0, which is FALSE and the while block immediately ends
- be careful not to jump out of while when encountering zero
 - zero is a perfectly reasonable value (in a file, in an array)
 - be conscious of the potential need for define in the condition
 - are you testing for truth (O is not true) or definition (O is defined)?

```
@nums = (0..5);
while ( defined($num = shift @nums) ) {
    print qq(Found $num at front of @nums);
}
```



I/O – Reading from a File

- reading from a file is easy just like most other things in Perl
- programs which use files typically follow these steps
 - open a file and create a filehandle (special variable type)
 - fetch one line at a time from the file, typically in a while loop
 - you parse each line and construct a data structure that holds this information
 - close the file when EOF (end of file) is encountered
 - the EOF evaluates to FALSE, which nicely terminates any while loop
 - process data
 - optionally, write data out to another file
- we will see how to
 - read from a file
 - write to a file
 - format strings with printf/sprintf



Reading from a File

reading from a file is done in three steps

```
# 1. open the file, creating a filehandle (conventionally capitalized)
open(FH,$file);
# 2. fetch next line from the file (via filehandle) using line operator <>
while( $line = <FH> ) {
    ...
}
# 3. close the filehandle (not strictly necessary, but good housekeeping)
close(FH);
```

- capitalize your filehandles (convention)
 - FILE, FH, HANDLE but not file, fh, or handle
 - you can have multiple handles
 - eventually you will use modules (IO:::File) that abstract raw filehandles
 - I lied, there is another variable type



Reading from a File

>	cat	file1.txt	

- 0 1 2 3

- Δ

```
open(FH,"file1.txt");
while( $line = <FH> ) {
  print $line;
close(FH);
0
1
2
3
4
```

- the <> operator (slurp operator) returns the next line includes the trailing newline
- the test within the while() loop when used with <> is implicitly a defined test
 - a file with a "O" and trailing newline "O\n" will evaluate to true because of the trailing newline
 - a file with a trailing "0" without a newline will evaluate to false, which is inconvenient and thus Perl applies defined
 - loop ends when <> returns EOF after the last line

• <> actually returns the next record in the file

- default record terminator is "\n", thus you get a line at a time
- you can change the record terminator and modify the behaviour of <> (here be dragons)

chomping lines

```
> cat file2.txt
a b c
      f
d
  e
  1
         2
0
3 4
5
open(FH,"file2.txt");
while($line = <FH>) {
 # remove trailing newline in $line
 chomp $line;
 # split line at whitespace (i.e. into words)
 @words = split(" ",$line);
 # concatenate words together using ":"
  print join(":",@words);
close(FH);
a:b:c
d:e:f
0:1:2
3:4
5
```

 chomp safely removes the trailing newline in a string
 it does nothing if a newline is not present

a 🖷

@list=split(" ",\$line) tokenizes the line into words
at whitespace (spaces or tabs)

03

IDIOM

- PP

GENOM



File Analysis Script

- we will create a script that performs the following
 - reads from a file
 - reports the number of words on each line
 - reports the total number of lines and words in the file
 - reports the average number of words per line
 - returns the 5 most common words
 - returns the 5 longest words



Step 1 – parsing the file

```
# keep count of each word in a hash
%words = ();
# keep number of words per line in an array
@wordcount = ();
open(FH, "sherlock.txt");
while($line = <FH>) {
  chomp $line;
  # split line at a run (one or more) non-word characters : \W is the opposite of \w
  @words = split(/\W+/,$line);
  # iterate through all words in the line
  wordcount = 0;
  for $word (@words) {
    # accept only words which have a letter character (e.g. no numbers)
   if($word =~ /[a-z]/i) { # /i is for case-insensitive match
       # increment count for this word
       $words{$word}++;
       $wordcount++;
    }
  }
  # add number of passed words in this line to the array
  push @wordcount, $wordcount;
close(FH);
```



Step 2 – reporting word statistics

```
$wordcount total = 0;
for $i (0..@wordcount-1) {
 # maintain count of all words seen
 $wordcount total += $wordcount[$i];
 # report on words on this line
 print qq(line $i had $wordcount[$i] words);
# report on word count statistics
print qq(saw ),scalar(@wordcount),qq( lines in file);
print qq(saw $wordcount total words in file);
print qq(average words/line ),$wordcount total/@wordcount;
# create sorted word lists - by frequency and length
@words common = sort { $words{$b} <=> $words{$a} } keys %words;
@words length = sort { length($b) <=> length($a) } keys %words;
for $i (0..4) {
  print qq(common word $i $words common[$i]);
for $i (0..4) {
  print qq(longest word $i $words length[$i]);
```

line 0 had 9 words line 1 had 0 words line 2 had 0 words

line 12649 had 0 words line 12650 had 0 words line 12651 had 0 words

saw 12652 lines in file saw 105999 words in file average words/line 8.3780429971546

common word 0 the common word 1 I common word 2 and common word 3 to common word 4 of longest word 0 disproportionately longest word 1 indistinguishable longest word 2 conventionalities longest word 3 scissorsgrinder longest word 4 inconsequential



Reading FASTA files

- FASTA file is a simple sequence format
 - first line starts with ">" and contains a header
 - first word in header is referred to as the ID
 - sequence follows, usually 50-80 bp per line

>gi|4878025|gb|U80929.2|CVU80929 Cloning vector pBACe3.6, complete sequence GATCCGCGGAATTCGAGCTCACGCGTACTGATGCATGATCCGGGTTTAAACCCAGTACTCTAGATCCTCT AGAGTCGACCTGCAGGCATGCAAGCTTGGCGTAATCATGGTCATAGCTGTTTCCTGTGTGAAATTGTTAT CCGCTCACAATTCCACACAACATACGAGCCGGAAGCATAAAGTGTAAAGCCTGGGGTGCCTAATGAGTGA GCTAACTCACATTAATTGCGTTGCGCTCACTGCCCGCTTTCCAGTCGGGAAACCTGTCGTGCCAGCTGCA TTAATGAATCGGCCAACGCGCGGGGAGAGGCGGTTTGCGTATTGGGCGCTCTTCCGCTTCCTCGCTCACT

. . .

GTCATGCCCCGCGCCCACCGGAAGGAGCTGACTGGGTTGAAGGCTCTCAAGGGCATCGGTCGAGCTTGAC ATTGTAGGACTATATTGCTCTAATAAATTTGCGGCCGCTAATACGACTCACTATAGGGAGAG

- there are modules that help you process FASTA files
- Iet's write a script to read a FASTA file and produce statistics

1.0.1.8 - Introduction to Perl



Step 1 – reading FASTA file

```
%bp = ();
# open human chr22 assembly
open(FH,"/home/martink/work/ucsc/hg18/fasta/chr22.fa");
while($line = <FH>) {
    chomp $line;
    # skip header
    next if $line =~ /^>/;
    # we're in the sequence
    # store count for each unique bp
    for $bp (split("",$line)) {
        $bp{$bp}++;
    }
}
```

- the ^ in regexps is an anchor which matches start of line

· /^hello/ matches hello at the start of a line

- the \$ in regexps is an anchor which matches end of line

- /goodbye\$/ matches goodbye at the end of a line

- challenge – what does /^\$/ match?



Step 2 – processing base pair types

<pre>%bpstats = (); for \$bp (keys %bp) { # print count for each bp type seen print qq(\$bp \$bp{\$bp}); # keep independent count of different bp types if (\$bp =~ /[atcg]/) { \$bpstats{repeat} += \$bp{\$bp}; } elsif (\$bp =~ /n/i) { \$bpstats{padding} += \$bp{\$bp}; } elsif (\$bp =~ /n/i) { \$bpstats{padding} += \$bp{\$bp}; } elsif (\$bp =~ /[GC]/) { \$bpstats{gc} += \$bp{\$bp}; } elsif (\$bp =~ /[AT]/) { \$bpstats{at} += \$bp{\$bp}; } else { theretats{up}}</pre>	a 4510978 A 4555927 c 3824338 C 4516771 t 4480960 T 4544042 n 17 N 14789904 g 3813956 G 4517817
<pre>\$bpstats{unk} += \$bp{\$bp}; } # return count of bps, by category for \$statistic (sort {\$bpstats{\$b} <=> \$bpstats{\$a}} keys %bpstats) { print qq(\$statistic \$bpstats{\$statistic}); }</pre>	repeat 16630232 padding 14789921 at 9099969 gc 9034588



Writing to a File

- the easiest way to write to a file is to redirect the output of your script to a file

- anything printed to STDOUT will be sent to a file
- anything printed to STDERR will be sent to the screen

% my_script.pl > file.txt

• to redirect both STDOUT and STDERR to a file,

% my_script.pl &> file.txt

to redirect to different files,

% my_script.pl > file.txt 2> file.err.txt



- sometimes you need to write to a file from your script
- open the file with open() but prefix filename with > or >>
 - •>file create and overwrite if necessary

• >>file append

- pass the filehandle as the first argument to print

•print FH \$num

```
open(FH,">file.txt");
for $num (@nums) {
    print FH $num,"\n";
}
close(FH);
```



- no comma between filehandle and arguments to print

• print FH, \$num1, num2;

G E



Creating a Random FASTA file

- let's create a random 100,000 bp FASTA file

```
@bp = qw(a t g c A T G C);
$sequence = "";
for (1..100000) {
   # $array[rand(@array)] idiom - randomly samples array
    $sequence .= $bp[rand(@bp)];
}
open(FH,">randseq.fa");
print FH ">random sequence\n";
# 4-argument substr returns 70 characters and replaces them with empty string
while( $line = substr($sequence,0,70,"") ) {
    print FH $line,"\n";
}
close(FH);
>random sequence
CCCGagttcAtGCGTCcTcATAAtgTTaGAGTcGAAtTTTgCctTaatTAGcagAcatcGTgAttaTcGg
aatctCAgagCCTCttcgcGtttTggTaTcgGcAgTcGaAaCcGCTagacatTgGaActgCcacagtAtt
. . .
cGcaACctCCacaAcTGgGtGgGTacagtCATGaTgCtAGtTgttTCCaTaGaGcagAcAcCttCGcCaa
TtTCgGTtTTCGACTCCCAccCgTagAAtAACGtCaCTgT
```



Formatted Output

- it is often desirable to prettify output for better readability
 - pad strings to fixed number of characters
 - specify number of decimals in a string
- printf is used to output a prettified string
- sprintf is used to generate a prettified string (which may be printed)

```
printf FORMAT_STRING,LIST;
```

```
$formatted_string = sprintf FORMAT_STRING,LIST;
```

the FORMAT_STRING specifies how the elements in the LIST are to be presented
 contains special entries like %s, %d, %f used to format LIST elements

printf

 $@x = qw(0 \ 1 \ 1.0 \ 1.6234);$ # truncated, not rounded printf "%d %d %d \n",@x; 0 1 1 1 # default 6 decimals printf "%f %f %f %f \n",@x; 0.000000 1.000000 1.000000 1.623400 # each field length is 10 printf "%10d %10d %10f %10f\n",@x; 0 1 1.000000 1.623400 # fix decimal places for floats printf "%10d %10d %10.3f %10.3f\n",@x; 0 1 1.000 1.623 # left justify first two fields printf "%-10d %-10d %10.3f %10.3f \n",@x; 0 1 1.000 1.623 # and now zero padding printf "%-010d %010d %010.3f %010.3f\n",@x; 0 000000001 000001.000 000001.623

- %d integer output
- %f float output
- %s string output
- %Nx field length N (%3d)
- %.Dx D decimal digits, where applicable (%.2f)

a 🖷

GEN O

- %N.Dx field length N with D decimal digits, where applicable (%5.2f)
- %<mark>0</mark>x 0-pad (%05.2f)

"%-x - left justify (%-05.2f)



sprintf

- let's create a script that produces the following
 - a random number
 - a 5-decimal truncated version of it and its square
 - a digit map (number of times each digit 0-9 seen in the number)

```
open(FH,">data.txt");
for $i (1..100) {
 x = rand();
 @digits = ();
 for $char (split("",$x)) {
    # count the number of times digit $char is seen
    # e.g. increment $digits[5] everytime 5 is seen
     $digits[$char]++ if $char =~ /\d/;
 }
 $i, $x, $x, $x**2, @digits);
 print FH $line,"\n";
close(FH);
line
     1 rand
              0.929494368378073 trunc 0.92949 trunc^2 0.86396 digitmap 2013201223
line
              0.903590672183782 trunc 0.90359 trunc^2 0.81648 digitmap 3122011222
     2 rand
. . .
```



Example - Filtering Files

- let's take all end sequence alignments of human clones and report the average clone sizes for different groups of clones, defined by regular expressions
 - one file contains the data (coordinates)
 - another file contains the filters (regexps) used to process the data
- coordinates defined in a file like this

```
M2131014 CTD-2131014 3 80809618 80926601
M2131015 CTD-2131015 6 121610096 121675696
...
```

- clone groups defined by file containing regular expressions

#comment		
A01\$		
10\$		
^N		
^D		
^.0001		



Step 1 – Reading coordinates and categories

```
%clonesize = ();
open(FH,"/home/martink/work/ucsc/hg17/bes/bacend.parsed.txt");
while($line = <FH>) {
    chomp $line;
   # M2131015 CTD-2131015 6 121610096 121675696
    # assigning to undef effectively skips the field
    ($clone,undef,$chr,$start,$end) = split(" ",$line);
    $size = $end - $start + 1;
   # keep track of size of each clone
    $clonesize{$clone} = $size;
}
close(FH);
# make an array of the regular expressions
(arx = ();
open(RX,"rx.txt");
while($line = <RX>) {
    chomp $line;
   # skip past comment lines
    next if $line =~ /^\s*#/;
    push @rx, $line;
}
close(RX);
```

6

IDIOM

G E N O M E SCIENCES C E N O M E

Step 2 – Processing clones

```
\%sum = ();
%count = ();
# check each clone whose size we know
for $clone (keys %clonesize) {
 # iterate through each regular expression
 for $rx (@rx) {
   # if the clone matches then keep track of total size/count for this category
   if($clone =~ /$rx/) {
     $sum{$rx} += $clonesize{$clone};
     $count{$rx} ++;
     # last is a flow-control key word which terminates the innermost enclosing loop
     last;
for $rx (sort keys %sum) {
 printf("group %10s num %6d avgsize %8.1f\n",
          $rx, $count{$rx}, $sum{$rx}/$count{$rx});
}
            10$ num 8502 avgsize 148126.2
group
          A01$ num 389 avgsize 140548.0
group
         ^.0001 num
                    195 avgsize 163235.3
group
             ^D num 39519 avgsize 138941.2
group
             ^N num 99252 avgsize 170593.5
group
```

1.0.1.8 – Introduction to Perl

1.0.8.1.5

Introduction to Perl Session 5

- you now know
 - while loop
 - reading from a file
 - writing to a file
 - printf/sprintf
- next time
 - subroutines
 - introduction to special variables
 - \$_ and friends



1.0.1.8.5 - Introduction to Perl - I/O

GENOM

a 🖭