Perl Intro for Bioinformatics

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Why Perl?

- Perl is a very, very concise language.
- Perl comes with extensive documentation.
 - See the *perl* and *perldoc* manpages for more.
- The Perl compiler and runtime system can provide an amazing amount of help to the user.

- In particular, see *perldoc diagnostics*.

- Perl is free, tried and true, and available for all major operating systems.
- Bioperl (http://www.bioperl.org/)

Basic Datatypes

- Perl has three main datatypes:
 - Scalars
 - Scalars hold single values, like integers, floating-point or real values, and strings.
 - Arrays
 - Arrays hold any number of scalars.
 - Hashes
 - Hashes associate keys with values. Both keys and values are scalar.

Scalar Variables

- All scalar variables begin with dollar-signs. my \$name = "Tim"; my (\$dx, \$dy) = (10, 20); my \$dist = sqrt (\$dx**2 + \$dy**2);
- Perl needs to be told to check for declarations.
 print \$x + 2; # Assumes \$x = 0, and prints 2 use strict; print \$x + 2; # Error, \$x undeclared!

Dynamic Typing

- Scalars can be integers, floats, or strings. my \$id = 123_456_789; print "ID = ", \$id; # prints ID = 123456789 \$id = "123 456 789"; print "ID = ", \$id; # prints ID = 123 456 789
- Perl will automatically convert, but be careful.
 \$next_id = \$id + 1; print "ID = ", \$next_id; # prints ID = 124

Strings

• Double quotes interpolate values.

my \$name = "Tim"; print "Hello, \$name!"; print 'Hello, \$name!'; # prints Hello, \$name!

prints Hello, Tim!

• You concatenate strings with '.' not '+'. print 'Hello, ' . \$name . '!'; # prints Hello, Tim! print '4' . '2'; # prints 42 print '4' + '2'; # prints 6 print 2 + 2.2 # prints 42

More Strings

- You can extract substrings from strings. my \$msg = "I am a fish!"; print substr(\$msg, 7, 4); # prints fish
- Or replace them with new substrings. substr(\$msg, 7, 4) = "hologram"; print \$msg; # prints I am a hologram!
- But you can't subscript strings like arrays.
 print \$word[0]; # Error: doesn't mean what you # think it means.

Other String Operators

- You can duplicate strings with 'x'.
 print "Fish" x 3; # prints FishFish
- Like numbers, strings can also be compared.
 # Numbers < <= == != >= > <=> # Strings lt le eq ne ge gt cmp
- Remember, Perl does automatic conversions.
 print '42' < '6' ? 'true' : 'false'; # prints false
 print 42 lt 6 ? 'true' : 'false'; # prints true

Arrays

- Arrays are like strings of scalar values. my @stuff = (3.14, 42, "hands of blue"); my (\$pi, \$ans, \$scary) = @stuff;
- Indexing returns a scalar, hence the '\$'.
 print "Two by two, \$stuff[2]!";
- You can also extract slices, replace slices, but you cannot nest arrays within other arrays.
 my @slice = @stuff[0..1]; # (3.14, 42)
 @slice[1..1] = (2, 3); # (3.14, 2, 3)
 my @more = (@slice, 5, 7); # (3.14, 2, 3, 5, 7)

Arrays and Strings

- Strings can be split, arrays can be joined.
 my @girls = split(" ", "Zoe Inara Kaylee River");
 print join(", ", @girls); # prints Zoe, Inara, Kaylee, ...
 print join(", ", sort @girls); # prints Inara, Kaylee, ...
- Array length and string length are different.
 print length(\$girls[1]); # prints 6
 my \$len = @girls;
 print \$len; # prints 4
 print @girls; # prints InaraKayleeRiverZoe
 print scalar @girls; # prints 4

Hashes

- Hashes are like Dictionaries in Java. my %ages = ('Tim' => 42, 'River' => 17, 'Summer' => 21); print \$ages { 'River' }; # prints 17
- Keys may not be kept in the order supplied.
 print join(", ", keys %ages); # prints River, Tim, ...
 print join(", ", values %ages); # prints 17, 42, 21
- Keys can be unquoted if they're barewords. delete \$ages {Tim}; print exists \$ages {Tim} ? 'true' : 'false'; # prints false

Advanced Datatypes

- In Perl, arrays and hashes can only hold scalars, but there is another kind of scalar: references.
- References are like pointers.

my @array = (1, 2, 3, 4); my \$aref = \@array;

• Dereferencing can get messy.

print @{\$aref}; # prints 1234
print \${\$aref}[0]; # prints 1
print \$\$aref[1]; # prints 2
print \$aref->[2]; # prints 3

More References

- There are easier ways to create references.
 my \$href = { Numbers => [1, 2, 3, 4] }; print @{\$href->{Numbers}}; prints 1234 # Without the @{}, Perl will only print the pointer value.
 print \$href->{Numbers}->[0]; prints 1; print \$href->{Numbers}[1]; prints 2; # Only the first -> is mandatory; without it, Perl would assume you were looking up 'Numbers' in %href.
- You can even make references to literals. my \$ten = \10; \$\$ten = 12; # Error: read-only value.

Data Structures

- Refs enable us to create complex data structures. my %people = (River => { age => 17, siblings => ['Simon'] }, Summer => { age => 21, gender => 'female' }
); print \$people { River } { age }; # prints 17 # Perl figures out that \$people{River} is a reference.
 print @ { \$people { River } { siblings } }; # prints Simon # Since we want the whole array, we have to use @{}
- A good tutorial on Perl references can be found in the *perlref* manpage.

Conditionals

- Perl has the traditional if-statement, but it also has 'unless'; note, the braces are not optional. my \$x = \$value <=> 42; if(\$x < 0) { print "Too small!"; } elsif(\$x > 0) { print "Too large!"; } else { print "Just right!"; } unless(@work) { print "Done."; }
- You can even suffix them for single statements no braces, and the parentheses are optional. die("Can't open \$file!") unless open(IN, "< \$file");

Loops

- It also has for-, while-, and also foreach-loops. for(my \$i = 0; \$i < @items; ++\$i) { print \$items[\$i]; } foreach my \$item (@items) { print \$item; } while(not \$done) { ...do something... } do { ...something... } until(\$done);
- Like if/unless, you can suffix each except for(;;).
 print "I am a fish!" while 1; # Infinite loop.
 print \$_ foreach @items; # Can't name the iterator.
- Perl uses next/last like C/Java's continue/break.
 foreach my \$x (@a) { next if \$x < 1; ... }

File I/O

• Arguments to 'open' resemble Unix sh redirects.

open IN, "< \$file"; # IN is the file handle. open OUT, ">> \$log"; # Append to \$log. open DAT, "+< \$db"; # Open for reading/writing. open LS, "ls -l |"; # We read the output of 'ls -l'.

- Reading and writing from streams is easy. my \$line = <IN>; print OUT "me: I just read \$line\n"; # No comma.
- Lines read may contain line-ends. chomp \$line;

Regular Expressions

- You can use R.E. to check the format of strings.
 print 'keyword' if \$token =~ m/^(if | then | else)\$/;
- But it is more interesting to extract data. my@numbers = (\$line =~ m/\d+/g); while(my \$line = <IN>) { next if \$line !~ m/([-.\w]+)@([-.\w]+)/; print "email: \$1\@\$2\n"; } my (\$dir, \$file) = \$path =~ m{(.+/)?([^/]+)};

Substitutions

- You can also use R.E. for search-and-replace. my \$msg = "She's wearing green."; \$msg =~ s/green/gold/; print \$msg; # prints She's wearing gold. my \$code = "if test then hiccup else wink"; \$msg =~ s/(if | then | else)/\U\1/g; print \$code; # prints IF test THEN hiccup ELSE wink
- There's also a similar utility for characters. my \$secret = "Don't tell anyone!"; \$secret = tr/a-z/k-za-j/; # Simple encryption. print \$secret; # prints Dyx'd dovv kxiyxo!

Subroutines

• Arguments are not formal, Perl puts them in (a). sub hypotenuse { my ($a, b) = @_;$ return sqrt (\$a**2 + \$b**2); } sub sum { my \$acc; acc += foreach $@_;$ return \$acc; }

```
print sum(1, 2, 3, 4); # prints 10
```

Reading FASTA files

```
open IN, "< $file" or die "Can't open $file: $!";
my $line = <IN>; # Read the descriptor (ignored).
my $seq;
while( $line = <IN> ) {
    last if $line =~ /^>/; # Stop if descriptor.
    $line = lc $line;
    $line = ~ s/[^a-z]//sg;
    $seq = $seq . $line;
}
return $seq;
```

Translate DNA into RNA

```
sub dna_to_rna {
    my ($dna) = @_;
    $dna = lc $dna;
    $dna = ~ s/[^acgt]//sg;
    my $rna = (reverse $dna) = ~ tr/acgt/ugca/;
    return $rna;
}
```

Translate RNA into Protein

```
my %codonMap; # ('gcu' => 'Ala', 'cgu' => 'Arg', ...)
```

```
sub rna_to_protein {
    my ($rna) = @_;
    my $protein;
    while( $rna =~ /(...)/g ) {
        $protein .= $codonMap{$1};
     }
    return $protein;
}
```

Initializing %codonMap

my %codonMap;

```
while( my $line = <DATA> ) {
    chomp $line;
    my @codons = split $line;
    my $residue = shift @codons;
    foreach my $c (@codons) {
        $codonMap{lc $c} = $residue;
    }
}
```

Other Points of Interest

- Modules
 - Build libraries of related subroutines which can be included with the *use* statement.
 - For an introduction see *man perlmod*.
- Object Oriented Programming
 - Perl supports OOP after a fashion. Robust, simple, and a bit off-putting at first, but you'll learn to like it.
 - For an introduction see *man perltoot*; if you're unfamiliar with OOP, start with *man perlboot*.
- Perl for Bioinformatics BioPerl
 - http://www.bioperl.org/