

Elements of Programming in Perl

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Functions and Perl Modules

Josep F. Abril

jabril@imim.es

Functions in Perl

& → &functions

```
sub reverse_seq() {  
    my $seq = shift;  
    $revseq = reverse $seq;  
    return $revseq;  
}
```

Arguments list: @_

```
&reverse_seq($seq)
```

Prototyping → `sub reverse_seq($);`

\ → References

```
$func = \&reverse_seq;  
$anon_func = sub { ...statements... };
```

&{\$func}(\$seq)
\$func->(\$seq)

Passing Arguments to Functions

By copy

```
...  
sub do_loop($@); # prototype  
...  
sub do_loop() { # implementation  
    my ($seq, @ary) = @_;  
    foreach my $q (@ary) {  
        print STDOUT "$seq : $q\n";  
    };  
} # do_loop  
...  
my $seq = "ATG";  
my @N = (1..10);  
my @M = ('A'..'Z');  
&do_loop($seq, @N, @M);  
...
```

By reference

```
...  
sub do_loop($$$); # prototype  
...  
sub do_loop() { # implementation  
    my ($seq, $ary1, $ary2) = @_;  
    foreach my $q (@$ary1) {  
        print STDOUT "A: $seq : $q\n";  
    };  
    foreach my $q (@$ary2) {  
        print STDOUT "B: $seq : $q\n";  
    };  
} # do_loop  
...  
my $seq = "ATG";  
my @N = (1..10);  
my @M = ('A'..'Z');  
&do_loop($seq, \@N, \@M);  
...
```

Working with Modules

```
sh$ perl -e 'print "BINC\n"'
sh$ export PERLSIBW="" # # or ## sh$ export PERLSOPT="-I.'
sh$ perl -e 'print "BINC\n"'
sh$ ./seq_analysis_modules.pl single_seq.fa

-----
SEQUENCE: dna_seq
--> STATS:
Length: 500bp
G+C: 61.20%

--> FORWARD Translations:
Phase 0 : TYSBLLGQNVHESGAWYRSEELFFFLFRSRSSFFCAARSELAAAGELPFIIFPSTVOVHTSASAGFNSGILLFW
GQELNFRPGRSDFPKI FPNVTR*HYATLQVEVLLGSDVDRAPAGLGRFFFLVFPVGVSPISASLKAPEAFT
Phase 1 : FGSRLTGTGSLQVWVNFARLLCLSELSPKCRGCPALHPGSLQADSCBQLPFLRMSILGNLQVYTAGCCBSC
GQTLSEIAGLGGQYLSQW*TLIGSQRBSGLAGWYHTLTERDAGLGGAFYFVFPVTV*VDEFFLALL*VQSGELL
Phase 2 : PGLI*PWEKGFH*IRCECSLAFAPAFAPFPFAAIVPLACTLRVCSRAVFNVPILLVTCQVY*DCVSRFPFVPLV
GGPFPSTRV*RT*DFRCERS*VQVCRDGG*FPGA*SQARSRGALPFPSSAQ*RRSIRGRFYGAS*GAVY

-----
--> REVERSE Translations:
Phase 0 : VVCSAGAFIEAL*NDGLPTLGGTCRGGKRAPSPAGALSSTLRPLSTVVFANIFEL*VFFGILGLLGLPVCFRMSF
PQNRGSRPFRSADGSLVFTVYRGLGIDRSPAAARISGAGCGDGRGCGGGGSGSSEAF*QDNRVTFQDNRDTR
Phase 1 : **APQLPL*IKCRNMF*RRHAELEGGSGRQPLBALCQLVAGPQPSRSTF*YRCSHLG*QV*TFKV*DFCGDP
PRTCTGQNLITQW*YGFV*RRGLGCTARLQPLVAVQDQTAAGGGAGKABASRHSILQCGQCGQDQDQ
Phase 2 : SRI*SNRCSGSRVWQVHT*YTCWQWCRSAPALCGVYHTVQAFVPLI*LE*SNVHTWESLQV*PACR*IKEMVFP
FQCGAVTC*RRCPNDLIGGGR*RLQAGCQL*CCSAGCEPGR*PGEGRER*SSANIT*PNV*VFPVRC*DCG

-----
```

Retrieving Command-line Options

```
#!/usr/bin/perl -s
# "s" is a rudimentary built-in switch parsing
defined($?) || die "EIOhelp";
print STDERR <<"EIOhelp";
# USAGE:
# ./script.pl --help=>file.out ...
EIOhelp
exit();
};
$verbose = defined($v) ? 1 : 0;
defined($outfile) || open(STDOUT, ">$outfile");

use Getopt::Std;
my ($opt_w, $opt_m, $opt_o) = (0,0,undef);
# declare vars when using strict
getopts("w:m:o:");
# sets $opt_w, $opt_m and $opt_o
$opt_w || $help();

my $args = (); # declare when using strict
getopts("w:m:o:", \%args);
# sets $arg{w}, $arg{m} and $arg{o}
defined($arg{H}) || $help();

sub help() {
    print STDERR <<"EIOhelp";
    # USAGE:
    # ./script.pl --help=>file.out ...
    # OPTIONS:
    # -w | --verbose
    #     Reporting program execution.
    # -o | --outfile "file.out"
    #     Setting output file (default STDOUT).
    # -h | --help
    #     Print this help.
    EIOhelp
    exit();
} # help

use Getopt::Long;
my ($verbose,$outfile) = (0,undef);
GetOptions(
    "v|verbose" => \$verbose,
    "o|outfile=" => \$outfile,
    "h|help" => \$help,
);
if (defined($outfile) {
    open(OFH, "> $outfile") || die("error...");
    $ofh = \*OFH;
} else {
    $ofh = \*STDOUT;
};
sub help() {
    print STDERR <<"EIOhelp";
    # USAGE:
    # ./script.pl (options) ...
    # OPTIONS:
    # -v | --verbose
    #     Reporting program execution.
    # -o | --outfile "file.out"
    #     Setting output file (default STDOUT).
    # -h | --help
    #     Print this help.
    EIOhelp
    exit();
} # help
```

CGI: Common Gateway Interface

```
# CGI script that creates a fill-out form
# and echoes back its values.

use CGI qw/:standard/;

print header,
start_html('A Simple Example'),
h1('A Simple Example'),
start_form,
"WHAT's your name? ",textfield('name'),p,
"WHAT's the combination?", p,
checkbox_group(-name=>'words',
-values=>['eemie','meenie','minie','noe'],
-defaults=>['eemie','minie']), p,
"WHAT's your favorite color? ",
popup_menu(-name=>'color',
-values=>['red','green','blue','chartreuse']),p,
submit,
end_form,
hr;

if (param()) {
    print "Your name is ",em(param('name')),p,
"The keywords are: ",em(join(", ",param('words'))),p,
"Your favorite color is ",em(param('color')),
hr;
}
exit();
```



DBI: DataBase Independent Interface

```
use DBI;
use strict;

$dbh = DBI->connect($dsn, $user, $password,
    { RaiseError => 1, AutoCommit => 0 });

$stmt = $dbh->prepare(
    "SELECT foo, bar FROM table WHERE baz=?");

$stmt->execute( $baz );

while ( @row = $stmt->fetchrow_array ) {
    print "@row\n";
};
```

BioPerl

<http://bioperl.org/>

```
use Bio::Seq;
use Bio::SeqIO;

$seqin = Bio::SeqIO->new(-file => "human.gb",
    -format => 'GenBank' );
$seqout = Bio::SeqIO->new(-file => ">forward.fa",
    -format => 'Fasta' );
$seqtwo = Bio::SeqIO->new(-file => ">reverse.fa",
    -format => 'Fasta' );

while ( my $seq = $seqin->next_seq() ) {
    $id = $seq->display_id();
    print STDERR "Working on: $id\n";
    $seqout->write_seq($seq);
    $id = "$id.rev";
    $rev = $seq->revcomp();
    $rev->display_id($id);
    $seqtwo->write_seq($rev);
};
```

Inline

Embedding code from other languages (Assembler, C, C++, Java, Python, Awk, ...).

Handles for us all the compilation steps required for those languages before binding the new functions into our script.

Easy access to the Advanced Programming Interface (API) to reuse perl internal implemented functions in C.

```
use Inline {Prism;};
use Inline C;

$filename = $ARGV[0];
die "Usage: perl vowels.pl filename\n" unless $filename;

$stat = join " ", <>; # skip input file
$fp = vowel_stats($stat); # call our function
$fp = sprintf("%03.1f", $fp * 100); # format for printing
print "The letters in $filename are $fp vowels.\n";

__END__
__C__

/* Find percentage of vowels to letters */
double vowel_ratio(char* str) {
    int letters = 0;
    int vowels = 0;
    int i = 0;
    char c;
    while (c = str[i++]) {
        /* normalize for case in ASCII; upper in EBCDIC */
        char a = normalize ? toLower : toUpper;
        char b = normalize ? toLower : toLower;
        char c = normalize ? toLower : toLower;
        char d = normalize ? toLower : toLower;
        char e = normalize ? toLower : toLower;
        char f = normalize ? toLower : toLower;

        while (c = str[i++]) {
            if (isVowel(c)) {
                letters++;
                if (isVowel(c)) {
                    vowels++;
                }
            }
        }
    }

    return (double)vowels / (double)letters;
}
```
