

Lecture 2:

Programming in Perl: Introduction 1

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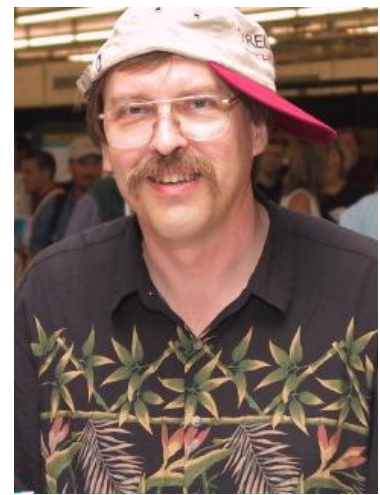
Guest lecturer
Umeå Plant Science Centre
Computational Life Science Cluster (CLiC)

This lecture

- Introduction to Perl 1
 - basic expressions
 - scalars
 - arrays
 - loops
 - conditions
 - file handling

What is Perl ?

- Perl was created by Larry Wall
- Perl = Practical Extraction and Report Language
- Perl is an Open Source project
- Perl is a cross-platform programming language



Why Perl

- Perl is a very popular programming language
- Perl allows a rapid development cycle
- Perl has strong text manipulation capabilities
- Perl can easily call other programs

- Existing Perl modules exists for nearly everything
 - <http://www.bioperl.org>
 - <http://www.cpan.org/> (Comprehensive Perl Archive Network)

ActivePerl

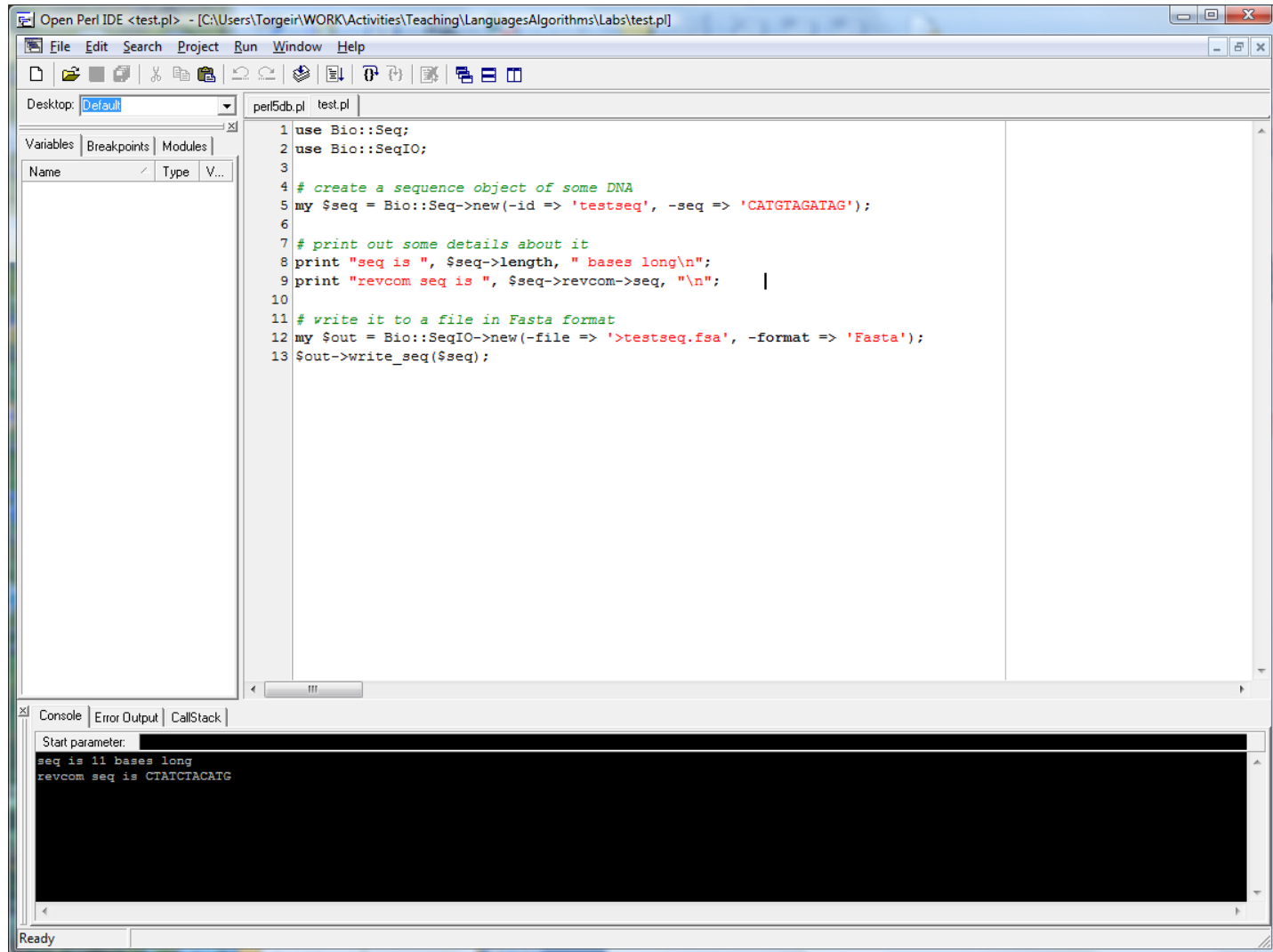
The screenshot shows the Perl Package Manager interface. The search bar contains 'bioperl'. The main list displays various packages with columns for Package Name, Area, Installed, Available, and Abstract. The 'BioPerl' package is highlighted in blue. Below the list, the 'Details' tab is active, showing the following information for BioPerl:

BioPerl
Bioinformatics Toolkit
Version: 1.5.9_4
Released: 2009-1-21
Author: BioPerl Team <bioperl-l@bioperl.org>
CPAN: http://search.cpan.org/dist/BioPerl-1.5.9_4/

At the bottom of the window, a status bar indicates: 10781 packages, 11 listed | 88 installed, 0 to install, 0 to remove | Install Area: **site**

Package Name	Area	Installed	Available	Abstract
Bundle-BioPerl-Run			1.5.2_100	Bundle of pre-requisites for bioperl-run
Bundle-BioPerl-N...			1.5.2_100	Bundle of pre-requisites for bioperl-network
Bundle-BioPerl-Db			1.5.2_100	Bundle of pre-requisites for bioperl-db
Bundle-BioPerl-Core			1.5.2_100	Bundle of pre-requisites for bioperl
Bundle-BioPerl-Core			1.5.9_4	Bundle of pre-requisites for BioPerl
Bundle-BioPerl			2.1.8	A bundle to install external CPAN modules used by BioPerl 1.5.2
bioperl-run			1.5.2_100	bioperl-run - wrapper toolkit
bioperl-network			1.5.2_100	bioperl-network - package for biological networks
bioperl-db			1.5.2_100	bioperl-db - package for biological databases
bioperl			1.5.2_100	Bioinformatics Toolkit
BioPerl	site	1.5.9_4	1.5.9_4	Bioinformatics Toolkit
Bio-mGen			1.03	a fast and simple gene loading, helping automate BioPerl processes.

Open Perl IDE



Our first Perl program

```
use strict;  
use warnings;
```

```
print "Hello world!\n";
```

Hello world!

“use strict” makes it harder to write bad software

”use warnings” makes Perl complain at a huge variety of things that are almost always sources of bugs in your programs

”\n” prints a new line

Perl scalars

- Perl variables that hold single values are called *Scalars*.
- Scalars hold values of many different **types** such as *strings*, *characters*, *floats*, *integers*, and *references*
- Scalars are written with a leading \$, like: \$sum
- Scalars, as all variables, are declared with **my**, like **my \$sum**
- Perl is not a typed language: scalars can be strings, numbers, etc.

- You can reassign values of different types to a scalar:

```
my $b = 42; $b = "forty-two"; print "$b\n";
```

```
forty-two
```

- Perl will convert between strings and numbers for you:

```
my $a = "42" + 8; print "$a\n";
```

```
50
```

```
my $a = "Perl" + 8; print "$a\n";
```

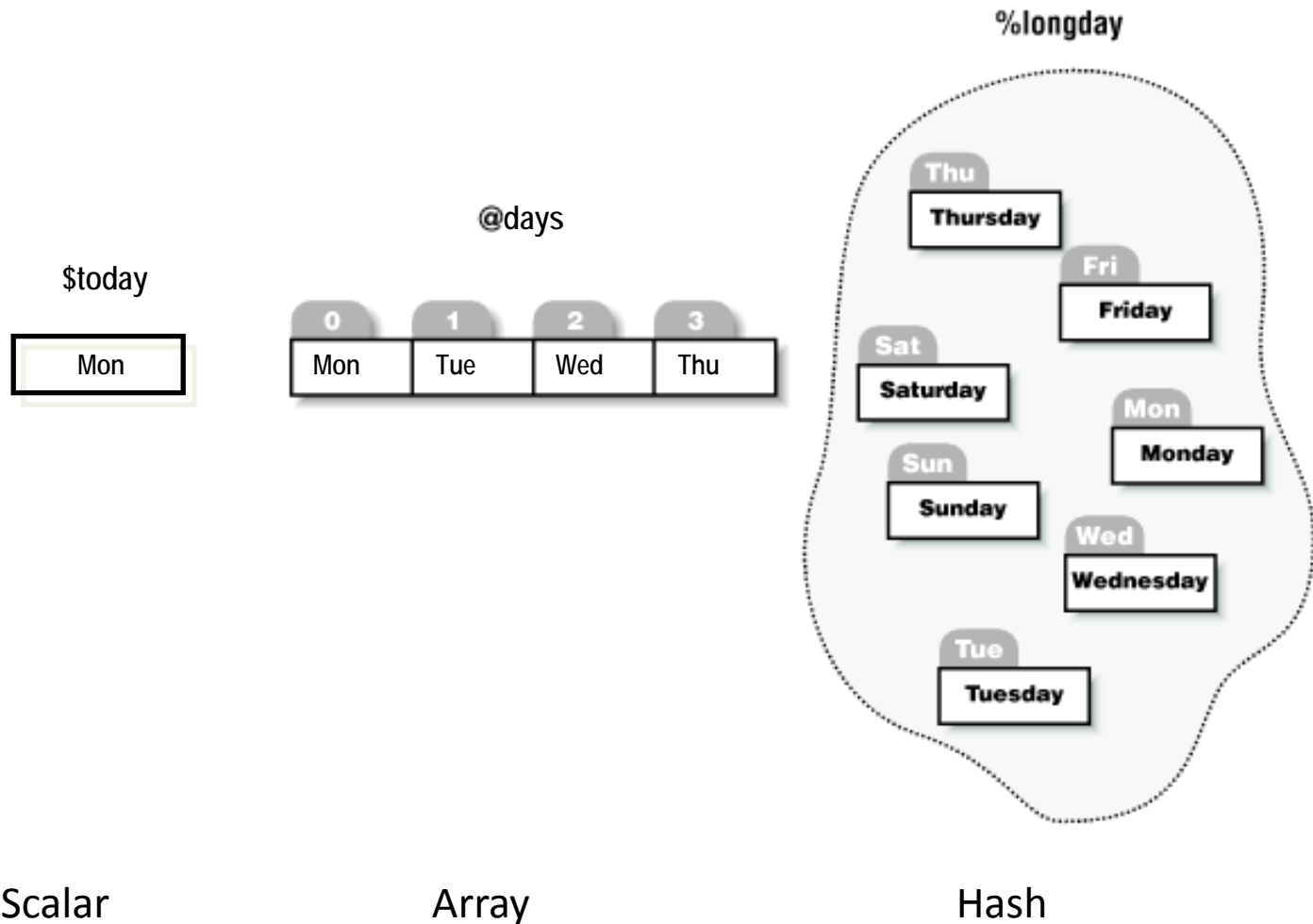
```
Argument "Perl" isn't numeric in addition (+) at test.pl line 4.
```

```
8
```


Perl scalars: some numerical operators

- `$v = 1+4;` `# addition`
- `$v = 5-4;` `# subtraction`
- `$v = 3*4;` `# multiplication`
- `$v = 7/8;` `# division`
- `$v = 2**8;` `# power`
- `$v = sqrt(4);` `# square root`
- `$i++;` `# $i = $i + 1;`
- `$i--;` `# $i = $i - 1;`
- `$i+= 5;` `# $i = $i + 5;`
- `$i/=5;` `# $i = $i / 5;`
- Everything after ”#” is not executed

The three fundamental datatypes in Perl



- The *sigills* `$`, `@`, `%` must always be used.
- You can use different datatypes with the same name in the same program.

Perl Arrays

- *Arrays* hold multiple ordered values.

- Arrays are written with a leading `@`, like: `@shopping_list`

- Arrays can be initialized by lists.

```
my @s = ("milk","eggs","butter"); print "@s\n";
```

milk eggs butter

- Arrays are indexed by integer. The first scalar in an array has index `0` and no matter its size, the last scalar has index `-1`:

```
my @s = ("milk","eggs","butter"); print "$s[0] - $s[-1]\n";
```

milk - butter

- The sizes of arrays are not declared; they grow and shrink as necessary.

```
my @s = ("milk","eggs","butter"); $s[4] = "beer"; print "@s\n";
```

Use of uninitialized value in join or string at test.pl line 4.

milk eggs butter beer

Perl Arrays

- Arrays can be iterated over in `foreach` loops. You don't need to know their size:

```
my @s = ("milk","eggs","butter");  
foreach (@s) {  
    print "$_\n";  
}
```

milk

eggs

butter

`$_` is known as the "default input and pattern matching variable".

This is all equivalent:

```
my @s = ("milk","eggs","butter");
foreach (@s) {
    print;
    print "\n";
}
```

```
my @s = ("milk","eggs","butter");
foreach my $item (@s) {
    print "$item\n";
}
```

```
my @s = ("milk","eggs","butter");
foreach (@s) {
    print "$_\n";
}
```

Perl Arrays

An array in scalar context evaluates to its size. You can easily get the index of the last item in an array.

```
my @s = ("milk","eggs","butter");  
my $length = @s;  
print "$length\n";  
3
```

```
my @s = ("milk","eggs","butter");  
my $last_index = $#s;  
print "$last_index\n";  
2
```

```
my @s = ("milk","eggs","butter");  
print "$s[$#s]\n";  
butter
```

Perl Arrays

Special commands add or remove items to the front or back of arrays.

`push` and `pop` add to the back, making a stack.

```
my @s = ("milk","eggs","butter");  
push @s, "beer";  
print "@s\n";  
milk eggs butter beer
```

```
my @s = ("milk","eggs","butter");  
pop @s;  
print "@s\n";  
milk eggs
```

```
my @s = ("milk","eggs","butter");  
my $last_item = pop @s;  
print "$last_item\n";  
butter
```

Perl arrays

grow or shrink as needed

"fred"	"wilma"
--------	---------

@data

```
my @data = ("fred","wilma");
```


Perl arrays

grow or shrink as needed

"fred"	"wilma"	42
--------	---------	----

@data

```
my @data = ("fred","wilma");  
push @data, 42;
```

Perl arrays

grow or shrink as needed

"fred"	"wilma"	42	undef	undef	"dino"
--------	---------	----	-------	-------	--------

@data

```
my @data = ("fred","wilma");  
push @data, 42;  
$data[5] = "dino";
```

undef

- The value of all uninitialized scalars (and scalar elements of arrays and hashes) has the special scalar value `undef`.
- `undef` evaluates as `0` when used as a number and `""` when used as a string, which is why you most often don't have to initialize variables explicitly before you use them.

```
my $a; $a++; print "$a\n";
```

```
1
```

```
my @a = (1,2);
```

```
$a[3] = 23; print "@a\n";
```

Use of uninitialized value in join or string at test.pl line 4.

```
1 2 23
```

- Even after a scalar has been assigned, you can undefine them using the `undef` operator.

```
$a = undef;
```

```
undef @a;
```

Arrays and lists in assignments

"fred"	"wilma"	42	undef	undef	"dino"
--------	---------	----	-------	-------	--------

@data

You can initialize or set arrays or lists by arrays or lists:

```
my ($man,$wmn) = ($data[0],$data[1]); print "$man $wmn\n";
```

```
fred wilma
```

```
my ($man,$wmn) = @data; print "$man $wmn\n";
```

```
fred wilma
```

```
@data = ("barney", "bambam"); print "@data\n";
```

```
barney bambam
```

```
my @mydata = @data; print "@data | @mydata\n";
```

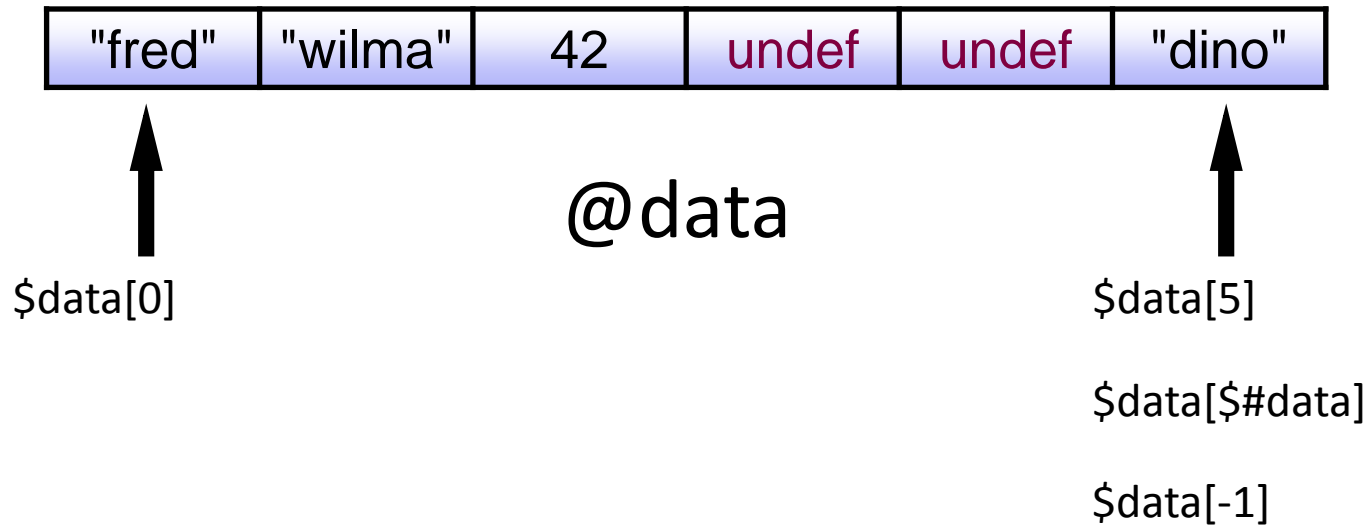
```
barney bambam | barney bambam
```

You can swap elements without a temporary:

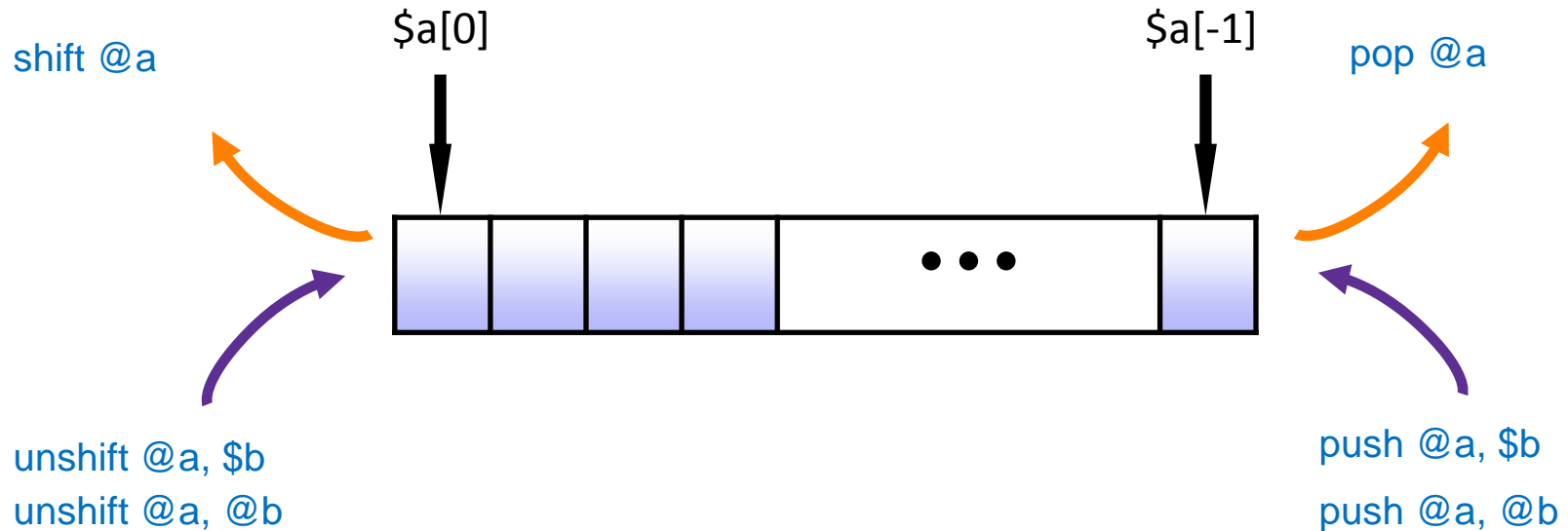
```
($data[1],$data[0]) = ($data[0],$data[1]); print "$data[0] $data[1]\n";
```

```
bambam barney
```

Array indexing



Adding elements to array ends



Loops: Iterating over Arrays

When we need the index:

```
for ($i = 0; $i < @data ; $i++) { # c-style
    print "$data[$i]\n";
}
```

When we only need the element:

```
foreach (@data) { # perl-style
    print "$_\n";
}
```

conditions

- `if – else` statements are used to test whether an expression is true or false

```
if ($a < 0) {  
    print "$a is a negative number\n";  
} elsif ($a == 0) {  
    print "$a is zero\n";  
} else {  
    print "$a is a positive number\n";  
}
```

- Use the function `defined` to test if a scalar has the value `undef`

```
if (defined $a) {  
    $a++;  
}
```

equivalent to

```
$a++ if defined $a;
```


The rules of truth in Perl

- Only Scalars can be True or False
- `undef` is False
- `""` is False
- `0` is False
- `0.0` is False
- `"0"` is False
- Everything else is True (including `"0.0"` !)

Logical expression

- `$a == $b` # compare numbers, true if \$a equal to \$b
- `$a != $b` # compare numbers, true if \$a is not equal to \$b
- `$a eq $b` # compare strings, true if \$a is equal to \$b
- `$a ne $b` # compare strings, true if \$a is not equal to \$b
- `!$a` # boolean, true if \$a is 0, false if \$a is 1

Controlling loops: next and last

next skip to the next iteration

```
my @a = (1,2,5,6,7,0);
```

```
my @filtered;
```

```
foreach (@a) {  
    next if $_ < 5;  
    push @filtered, $_;  
}
```

```
print "@filtered\n";
```

```
5 6 7
```

last ends the loop

```
my @a = (1,2,5,6,7,0);
```

```
my $found_zero = 0;
```

```
foreach (@a) {  
    if ($_ == 0) {  
        $found_zero = 1;  
        last;  
    }  
}
```

```
print "$found_zero \n";
```

```
1
```

Sorting arrays

- Use the built in function `sort`

- The results may surprise you!

```
my @words = ("c","b","a","B");
```

```
@words = sort @words;
```

```
print "@words\n";
```

```
B a b c
```

```
my @numbers = (10,3,1,2,100);
```

```
@numbers = sort @numbers;
```

```
print "@numbers\n";
```

```
1 10 100 2 3
```

sort

- `sort` uses a default sorting operator `cmp` that sorts "ASCIIbetically", with capital letters ranking over lower-case letters, and then numbers.

`sort @words;`

is equivalent to:

`sort {$a cmp $b} @words;`

- `cmp` is a function that returns three values:
 - -1 if `$a le $b`
 - 0 if `$a eq $b`
 - +1 if `$a ge $b`
- where `le`, `eq`, and `ge` are string comparison operators.
- `$a` and `$b` are special scalars that only have meaning inside the subroutine block argument of `sort`. They are aliases to the members of the list being sorted.

sort {\$a <=> \$b} @numbers

- `<=>` (the "spaceship operator") is the numerical equivalent to the `cmp` operator:
 - -1 if `$a < $b`
 - 0 if `$a == $b`
 - +1 if `$a > $b`

- You can provide your own named or anonymous comparison subroutine to `sort`:

```
my @numbers = (10,3,1,2,100);
```

```
@numbers = sort {$a <=> $b} @numbers;
```

```
print "@numbers\n";
```

```
1 2 3 10 100
```

```
@numbers = sort {$b <=> $a} @numbers;
```

```
print "@numbers\n";
```

```
100 10 3 2 1
```

Syntax summary: scalars

- Declare: `my $age;`
- Set: `$age = 29; $age = "twenty-nine";`
- Access: `print "$age\n";` `twenty-nine`

Syntax summary: arrays

- Declare: `my @children;`
- Set all: `@children = ("Troy","Anea");`
- Set element: `$children[0] = "Troy Alexander";`
- Access all: `print "@children\n";` `Troy Alexander Anea`
- Access element: `print "$children[1]\n";` `Anea`

Syntax summary: loops

```
foreach my $child (@children) {  
    print "$child\n";  
}
```

Troy Alexander

Anea

```
for (my $i = 0; $i < @children; $i++) {  
    print "$i: $children[$i]\n";  
}
```

0: Troy Alexander

1: Anea

Syntax summary: conditions

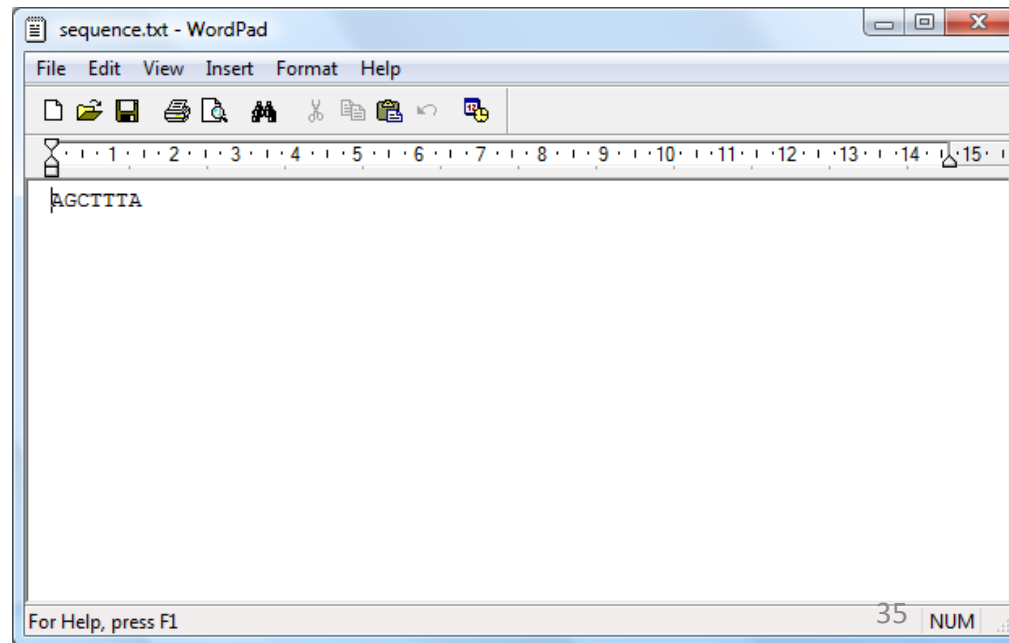
```
foreach my $child (@children) {  
    if (length($child) > 4) {  
        print "$child\n";  
    }  
}
```

Troy Alexander

Reading and writing to files

- `open(A, ">sequence.txt")` – creates a new file and opens it for writing
- `open(A, ">>sequence.txt")` – opens an existing file for writing
- `open(A, "sequence.txt")` – opens an already existing file for reading

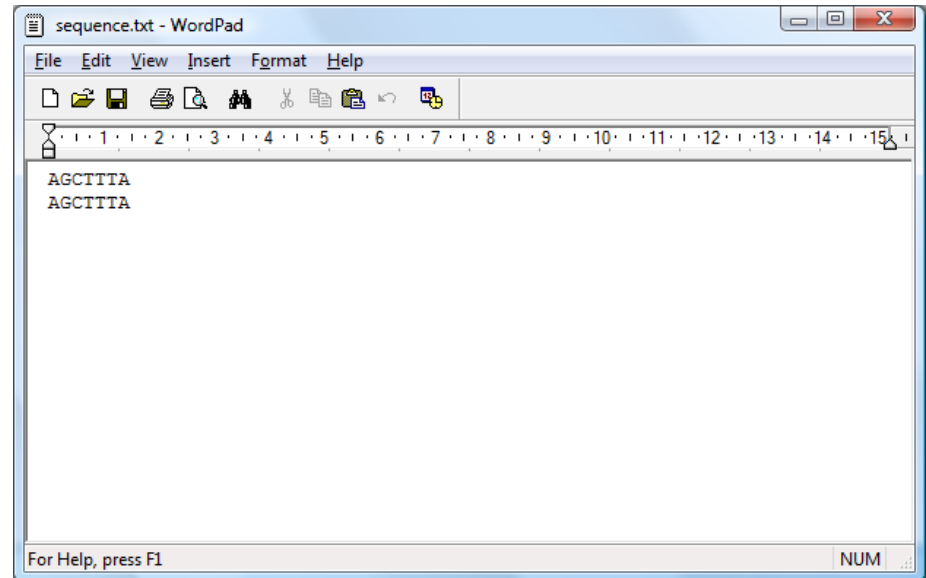
```
open(A, ">sequence.txt");  
print A "AGCTTTA\n";  
close(A);
```



Reading and writing to files

```
open(A , ">>sequence.txt");  
print A "AGCTTTA\n";  
close(A);
```

```
open(A , "sequence.txt");  
my $line1 = readline *A;  
my $line2 = readline *A;  
close(A);  
print "$line1 | $line2\n";  
AGCTTTA  
| AGCTTTA
```



Reading files

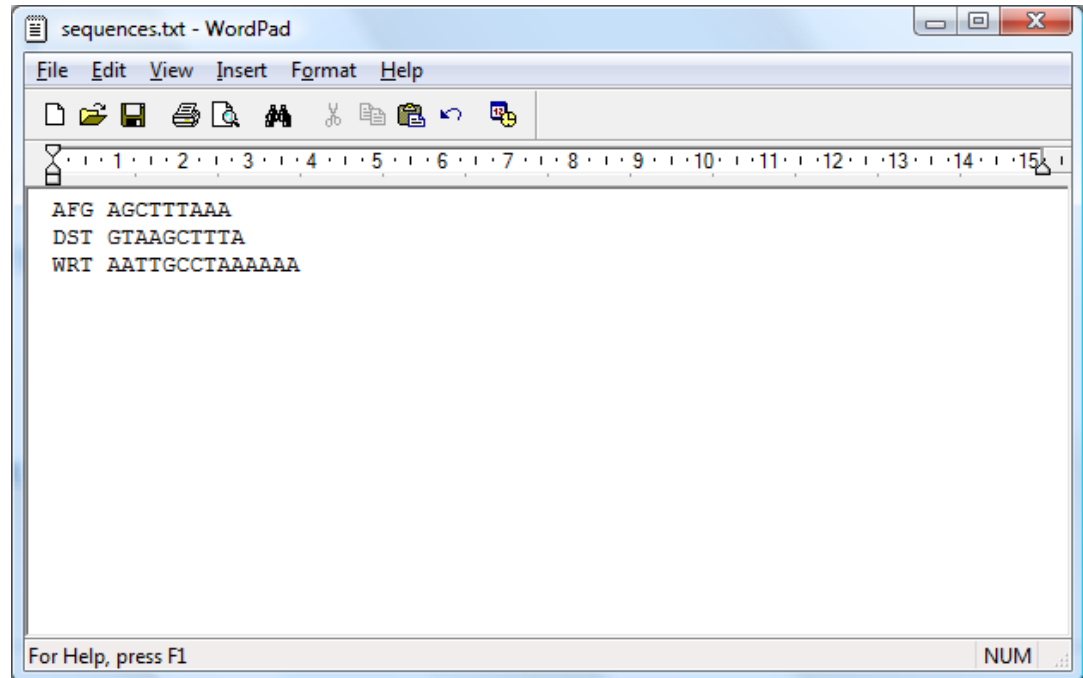
```
my @seqs;  
open(A, "sequence.txt");  
while (<A>) {  
    chomp;  
    push @seqs, $_;  
}  
close(A);  
print "@seqs\n";  
AGCTTTA AGCTTTA
```

`chomp` removes "\n" from the end of the line if it exists

Splitting strings: split

- You can split a string on any substrings that match a regular-expression with:
 - `@array = split /PATTERN/, $string;`
 - `split /\s/, "do the twist";` # gives ("do","the","twist")
 - `split //, "dice me";` # gives ("d","i","c","e"," ","m","e");
- Extremely useful when parsing files:

```
my @genes;  
open(A, "sequences.txt");  
while (<A>) {  
    chomp;  
    my ($gene) = split /\s/;  
    push @genes, $gene;  
}  
close(A);  
print "@genes\n";  
AFG DST WRT
```



Extracting fragments: substr

```
my $string = "AC Milan";  
my $fragment = substr $string, 3;  
print "$fragment\n";
```

Milan

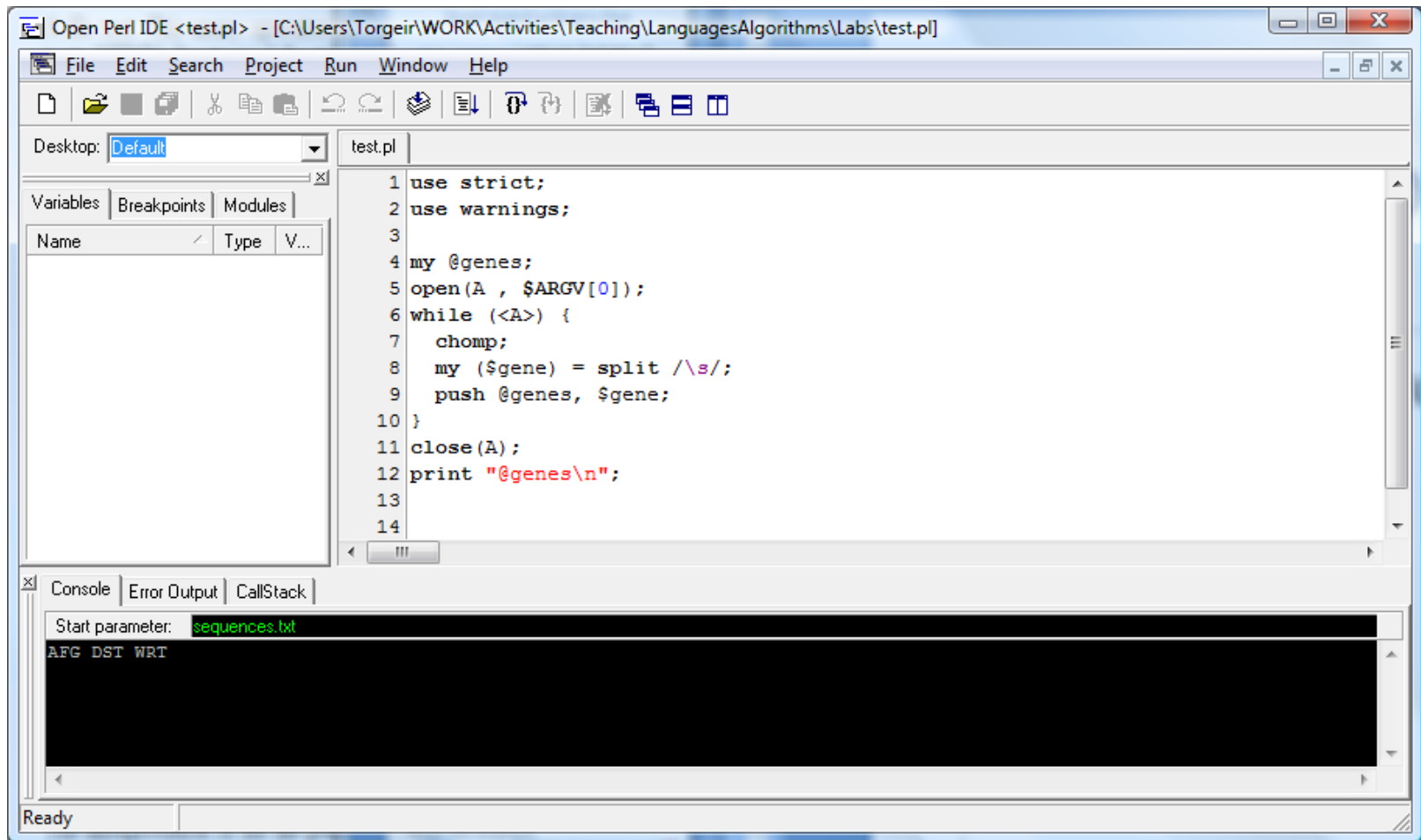
```
my $string = "F.C. Internazionale";  
my $fragment = substr $string, 5, 5;  
print "$fragment\n";
```

Inter

```
my $string = "F.C. Internazionale";  
my $fragment = substr $string, -7, 4;  
print "$fragment\n";
```

zion

@ARGV: command-line arguments



Acknowledgements

- Several slides were taken or re-worked from David Ardell and Yannick Pouliot.