

Lab1 Part A

Exercise 1.1

Code

```
use warnings;
print "welcome to Perl\n";
print "TATA is a box\n";
print "\n===== \n";
#remove the first \n in first line
print "welcome to Perl";
print "TATA is a box\n";
print "\n===== \n";
#second \n also removed
print "welcome to Perl";
print "TATA is a box";
print "\n===== \n";
#first in but second removed
print "welcome to Perl\n";
print "TATA is a box";
print "\n===== \n";
#insert \t
print "welcome\t to Perl\n";
print "TATA is a box\n";
print "\n===== \n";
#insert \t at different location 转义了
print "welcome\t t\t to Pe\trl\n";
print "TA\tTA is a box\n";
print "\n===== \n";
#replace double quote to single quote 没有转义
print 'welcome\t t\t to Pe\trl\n';
print 'TA\tTA is a box\n';
print '\n===== \n';
#no quote
# print welcome\t t\t to Pe\trl\n;
# print TA\tTA is a box\n;
# print '\n===== \n';
# run fail

# my explanation
# \n is new line
# \t is insert tab
```

```
# '' not convert \n and \t which means disable escape character
# "" enable escape character
```

Exercise 1.2

Code

```
#Exercise 1.2
print "\n===== \n";
print("When in double quotes:\n");
print("Use \\ to output \\n");
print("Use \\t to output tabs");
print "\n===== \n";
```

Exercise 1.3

Code

```
#Exercise 1.3
$dna = 'acCtagGgCCTTAcga';
$dna =~ tr/atcg/ATCG/;
$dna =~ tr/ATCG/UAGC/;
print("$dna \n");
print "\n===== \n";
```

Exercise 1.4

Code

```
#Exercise 1.4
$silly = 'tttTTT';
$silly =~ tr/tT/Tt/;
print("$silly\n");
print "\n===== \n";
```

Exercise 1.5

Code

```
#Exercise 1.5
$string = 'VRNrIAEelrrFMVALILdIKrTPgNKPrIaemICDIDtYIVeA';
$result = '';
```

```

print("$string[1]");
for (my $i = 0; $i <= length($string); $i++) {
    $char = substr($string, $i, 1);
    $char_c = $char;
    #if is a uppercase
    if (($char =~ /[A-Z]/)) {
        $char_c =~ tr/A-Z/a-z/;
    }
    else {
        $char_c =~ tr/a-z/A-Z/;
    }
    $result = $result . $char_c;
}
print("$result \n");
print "\n===== \n";

```

Exercise 1.6

Code

```

#Exercise 1.6
my $string = 'SEETQMRLQLKRKLQRNRTSFTQEIEALEKEFERTHYPDVFARERL';
#enum info
my $A_s = 'ACGPSTWY';
my $E_s = 'RNDQEHK';
my $I_s = 'ILMFV';
# preparing data
my $result = '';
for (my $i = 0; $i <= length($string); $i++) {
    $char = substr($string, $i, 1);
    if ($A_s =~ m[$char]) {
        $result = $result . 'A';
    }
    if ($E_s =~ m[$char]) {
        $result = $result . 'E';
    }
    if ($I_s =~ m[$char]) {
        $result = $result . 'I';
    }
}
print("$result \n");
print "\n===== \n";

```

Exercise 1.7

Code

```
#Exercise 1.7
$string3 = $string2 = $string1 =
'albctttttaglkmelkgghtaatactxmljeixjcaelkjaxm';
$string1 =~ tr/atcg/x/c; #除了(c)atcg外替换成x
$string2 =~ tr/atcg//cd; #除了(c)atcg 不进行替换并且删除(d)
$string3 =~ tr/atcg/~ /cs; #除了(c)atcg外替换成~并且讲多个 ~压缩成一个~
(s)
print("$string1 \n $string2 \n $string3\n");
print "\n===== \n";
```

Exercise 1.8

Code

```
#program 1 complement DNA sequence
$stringDNA = lc('atCGaTCG');
$stringDNA =~ tr/atcg/tagc/;
#program 2 done a transcription
$stringRNA = $stringDNA;
$stringRNA =~ tr/atcg/uagc/;
print($stringRNA);
print "\n===== \n";
```

Exercise 1.9

Code

```
#Exercise 1.9
$stringDNA = lc('atCGaTCG');
$replcount = $stringDNA =~ tr/ag/ag/;
print($replcount);
```

Output Screenshot

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