

testMeans, v. 0.2:

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1 Introduction

TestMeans implements the t-test.

2 Getting Started

TestMeans was written in C on a computer running Linux and should work on any standard UNIX system. However, please contact me at haubold@evolbio.mpg.de if you have any problems with the program.

- Unpack the program

```
tar -xvzf testMeans_XXX.tgz
```

where XXX indicates the version.

- Change into the newly created directory

```
cd TestMeans_XXX
```

and list its contents

```
ls
```

- Generate testMeans

```
make
```

- List its options

```
./testMeans -h
```

- Inspect the example data, which consists of paired measurements, which you can think of as control

```
cat exampleControl.txt
Cwc22 11.961 12.401 11.661 11.96 10.454 11.584 11.175 11.343
Yars2 9.347 9.341 9.29 9.441 9.602 9.892 10.058 9.99
```

and treatment

```
cat exampleTreatment.txt
Cwc22 8.479 8.523 8.793 8.726 9.677 8.728 8.383 11.086
Yars2 9.924 9.953 9.568 9.79 10.381 10.156 10.045 10.079
```

where for each gene (first column) eight measures were made (columns 2–9). `TestMeans` can deal with an arbitrary number of measurements as long as the files being compared have the same number of columns.

- Test program

```
./testMeans exampleControl.txt exampleTreatment.txt
Cwc22 1.157e+01 9.049e+00 2.807e-05
Yars2 9.620e+00 9.987e+00 2.166e-02
```

where the four columns in each row indicate

1. Sample name (gene in this case)
2. Average of measurements for sample 1
3. Average of measurements for sample 2
4. P -value, that is, the error probability when rejecting the null hypothesis that both samples were drawn from the same population.

3 Listing

The following listing documents the driver program for `testMeans`.

```
1  /***** testMeans.c *****/
* Description: Compare two means.
* Author: Bernhard Haubold, haubold@evolbio.mpg.de
* Date: Thu Dec 15 10:44:31 2016
*****/
6  #include <stdio.h>
#include <stdlib.h>
#include <string.h>
#include "interface.h"
#include <gsl/gsl_statistics_double.h>
11 #include "eprintf.h"
#include "tab.h"
#include "tTest.h"
#include "mcTest.h"
#include "gsl_rng.h"
16 #define MAX_ID_LEN 256

void test(FILE *fp1, FILE *fp2, Args *args) {
    char *line, *idl, *id2;
21    double *s1, *s2;
    int n1, n2, f, l, i;
    Result *res;
    gsl_rng *ran;

26    s1 = s2 = NULL;
    n1 = n2 = 0;
    l = 0;
    id1 = (char *)emalloc(MAX_ID_LEN);
    id2 = (char *)emalloc(MAX_ID_LEN);
31    res = (Result *)emalloc(sizeof(Result));
    while((line = tabGetLine(fp1)) != NULL) {
        l++;
    }
}
```

```

f = tabNfield() - 1;
36   if(f > n1)
      s1 = (double *)erealloc(s1, f * sizeof(double));
n1 = f;
id1 = strncpy(id1,tabField(0),MAX_ID_LEN);
for(i=1; i<=n1; i++)
  s1[i-1] = atof(tabField(i));
41   line = tabGetLine(fp2);
if(!line){
  printf("ERROR:_unpaired_line_#%d\n",l);
  exit(-1);
}
46   id2 = strncpy(id2,tabField(0),MAX_ID_LEN);
if(strcmp(id1,id2)){
  printf("ERROR:_%s_on_line_%d_paired_with_%s.\n", id1, l, id2);
  exit(-1);
}
51   f = tabNfield() - 1;
if(f > n2)
  s2 = (double *)erealloc(s2, f * sizeof(double));
n2 = f;
for(i=1; i<=n2; i++)
  s2[i-1] = atof(tabField(i));
56   if(args->t == 'm'){
    ran = ini_gsl_rng(args);
    res = mcTest(res, ran, args->i, s1, n1, s2, n2);
    free_gsl_rng(ran, args);
}else if(args->t == 's')
  res = student(res, s1, n1, s2, n2);
else if(args->t == 'w')
  res = welch(res, s1, n1, s2, n2);
printf("%s\t%.3e\t%.3e\t%.3e\n", id1, res->m1, res->m2, res->p);
61 }
free(id1);
free(id2);
}

71 int main(int argc, char *argv[]){
  int i, j;
  char *version;
  Args *args;
76   FILE *fp1, *fp2;

  version = "0.2";
  setprogname2("testMeans");
  args = getArgs(argc, argv);
  if(args->v)
    printSplash(version);
  if(args->h || args->e)
    printUsage(version);
  for(i=0;i<args->numInputFiles;i++){
    for(j=i+1;j<args->numInputFiles;j++){
86      fp1 = efopen(args->inputFiles[i],"r");

```

```

fp2 = efopen(args->inputFiles[j], "r");
test(fp1, fp2, args);
fclose(fp1);
fclose(fp2);
}
}
free(args);
free(progname());
96   return 0;
}

```

4 Change Log

- Version 0.1 (March 3, 2017)
 - First running version
- Version 0.2 (May 12, 2017)
 - Explain data format in help output (-h).