inspectPro, v. 0.2: DESCRIPTION

Bernhard Haubold

Max-Planck-Institute for Evolutionary Biology, Plön, Germany

November 14, 2012

1 Introduction

2 Getting Started

inspectPro 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 inspectPro_XXX.tgz
```

where XXX indicates the version.

• Change into the newly created directory

```
cd InspectPro_XXX
and list its contents
```

ls

• Generate inspectPro

make

• List its options

./inspectPro -h

3 Listing

The following listing documents the driver program for inspectPro.

```
/***** inspectPro.c ******************
* Description:
    * Author: Bernhard Haubold, haubold@evolbio.mpg.de
    * Date: Fri Oct 19 09:41:32 2012
    *******************

#include <stdio.h>
#include <stdlib.h>
#include <assert.h>
#include <string.h>
#include "interface.h"
```

```
#include "profileTree.h"
  #include "eprintf.h"
  void writeProfiles(Profile *pro, int n) {
16
    printf("#ID\tCount\tA\tC\tG\tT\n");
    for(i=0;i<n;i++)
      printf("%d\t%d\t%d\t%d\t%d\t%d\n",i,pro[i].n,
              pro[i].profile[0],
              pro[i].profile[1],
              pro[i].profile[2],
              pro[i].profile[3]);
  }
void writeLengths(int *len, int n) {
    int i;
    printf("#ID\tLength\n");
    for (i=0; i<n; i++)</pre>
      printf("%d\t%d\n",i,len[i]);
  void writePositions(FILE *fp) {
    int numRead;
   Position *pos;
    printf("#Pos\tPro\n");
    pos = (Position *)emalloc(sizeof(Position));
    numRead = fread(pos, sizeof(Position), 1, fp);
    assert(numRead == 1);
    while(!feof(fp)){
      printf("%d\t%d\n",pos->pos,pos->pro);
      numRead = fread(pos, sizeof(Position), 1, fp);
    free (pos);
  void scanFile(FILE *fp, Args *args) {
    char *tag;
    int numRead, *lengths;
    Profile *profiles;
    int n;
    tag = (char *)emalloc(3*sizeof(char));
    numRead = fread(tag, sizeof(char), 3, fp);
    assert(numRead == 3);
    while(!feof(fp)){
      if(strcmp(tag, "sum") == 0) {
        numRead = fread(&n, sizeof(int), 1, fp);
        assert(numRead == 1);
        profiles = (Profile *)emalloc(n*sizeof(Profile));
        numRead = fread(profiles, sizeof(Profile), n, fp);
        assert(numRead == n);
```

```
writeProfiles(profiles, n);
         free (profiles);
66
       }else if(strcmp(tag, "con") == 0) {
         numRead = fread(&n, sizeof(int), 1, fp);
         assert(numRead == 1);
         lengths = (int *)emalloc(n*sizeof(int));
         numRead = fread(lengths, sizeof(int), n, fp);
         assert(numRead == n);
         writeLengths(lengths, n);
         free(lengths);
       }else if(strcmp(tag, "pos") == 0) {
         writePositions(fp);
       }else
         assert(0);
       numRead = fread(tag, sizeof(char), 3, fp);
     }
     free(tag);
   int main(int argc, char *argv[]){
     int i;
     char *version;
     Args *args;
     FILE *fp;
     version = "0.2";
     setprogname2("inspectPro");
     args = getArgs(argc, argv);
     if(args->v)
       printSplash(version);
     if(args->h || args->e)
       printUsage(version);
     if(args->numInputFiles == 0){
       fp = stdin;
       scanFile(fp, args);
     }else{
       for(i=0;i<args->numInputFiles;i++) {
101
         fp = efopen(args->inputFiles[i], "rb");
         scanFile(fp, args);
         fclose(fp);
       }
     free (args);
     free(progname());
     return 0;
   }
```

4 Change Log

- Version 0.1 (October 22, 2012)
 - First version that runs.
- Version 0.2 (October 24, 2012)

- When printing positions the last position was printed twice; fixed.
- Filled in interface.
- Version 0.3 (November 14, 2012)
 - Implement inspection of * .lik file; works only with mlRho v. $\geq 1.25.$