

epos: Estimating Population Size from the Site Frequency Spectrum

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June 18, 2018

1 Introduction

Epos estimates historical population sizes based on a site frequency spectrum. The program implements theory developed by Michael Lynch (Arizona State University) and Peter Pfaffelhuber (Freiburg University), which will be described in a forthcoming paper.

2 Prerequisites

Epos depends on two libraries, the Gnu Scientific Library (`lgsl`), and the Basic Linear Algebra Subprograms (`lblas`).

3 Getting Started

epos 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.

- Change into the `epos` directory

```
cd epos
```

and list its contents

```
ls
```

- Generate `epos`

```
make
```

- List its options

```
./epos -h
```

- Test program

```
./epos data/kap144i.dat
```

4 Change Log

- Version 0.1 (Oct. 25, 2017)
 - First running version based on the GSL.
- Version 0.2 (Oct 26, 2017)
 - Used `LAPACKEdgesv` in `getPopSizes`; makes no difference compared to the previous version.
- Version 0.3 (Oct 26, 2017)
 - Used `LAPACKEdgesvx` in `getPopSizes`; makes no difference compared to previous version.
- Version 0.4 (Oct 27, 2017)
 - Allow construction of SFS based on `-t` switch.
 - Allow switching between the GSL algorithm (`-g`) and the default LAPACK algorithm.
- Version 0.5 (November 2, 2017)
 - Allow the straight usage of the trapezoid matrix for solving the system (`-T`).
 - Print out coefficient matrix (`-p`).
- Version 0.6 (November 10, 2017)
 - Use `long double` in `getPopSizesTri2`; did not help.
- Version 0.7 (November 10, 2017)
 - Implement Peter Pfaffelhuber's formula; working only partially.
- Version 0.8 (November 10, 2017)
 - Peter's equation (6) is working.
- Version 0.9 (November 10, 2017)
 - Peter's equation in arbitrary precision in MPFR library. Numerical stability achieved with 329 bits per number.
- Version 0.10 (November 30, 2017)
 - Implemented equation (3) from Peter's memo dated Nov. 13. Not working.
- Version 0.11 (December 1, 2017)
 - Implemented revised equation (3) from Peter's memo dated Nov. 30. Not working.
- Version 0.12 (December 15, 2017)
 - Implemented Estimator 2.1 from Peter's memo dated Dec. 2, 2017. Code is running, but the results look odd.
- Version 0.13 (December 16, 2017)
 - Fixed the implementation of Estimator 2.1; results OK now.
- Version 0.14 (December 18, 2017)
 - Implemented optimization strategy for folded SFS; working.
 - Implemented optimization strategy for unfolded/even SFS; working.
 - Implemented optimization strategy for folded/odd SFS; not working yet.

- Version 0.15 (December 18, 2017)
 - Fixed error in search for optimal number of steps.
- Version 0.16 (December 19, 2017)
 - Fixed error in left-hand side of folded/odd equation; working.
 - Changed search strategy.
 - Changed default value of $-d$ from 10^{-6} to 10^{-3} .
 - Simplified user interface.
 - Included set of test cases (`test.sh`).
- Version 0.17 (December 20, 2017)
 - Fixed searching routine.
 - Changed default value of $-d$ from 10^{-3} to 10^{-2} .
 - Included addition of the λ -factor; seems to make no difference.
- Version 0.18 (December 20, 2017)
 - Fixed the λ -factor; computation is now much stabilized.
- Version 0.19 (January 11, 2018)
 - Included the λ -factor in the computation of Ψ . Computations now applicable to real data.
 - Changed the default-value of λ from 10^{-7} to 10^{-5} .
- Version 0.20 (January 11, 2018)
 - Consider zero-class mutations in computation, if present.
 - Changed default λ to 2×10^{-5} to get all data sets to run.
- Version 0.21 (January 13, 2018)
 - Reverted output of levels, going from the present into the past.
 - Default output is now as a function of times instead as a function of levels.
 - Included “step-wise” option for plotting times and levels.
 - Fixed time computation.
 - Included error message for negative population sizes.
 - Removed memory leaks and other subtle bugs using `valgrind`.
- Version 0.22 (January 16, 2018)
 - Fixed important bug in function `foldedEpsi`, where variable `b` was computed as a function of `sfs->f[n/2]` instead of, now `sfs->f[n/2-1]`.
- Version 0.23 (January 16, 2018)
 - Search for optimal λ .
 - Allow arbitrary level as first entry in level list.
 - Output λ , Ψ , and the levels added to make it easier to follow the program.
 - Reduced program to folded/even case.
- Version 0.23 (January 17, 2018)
 - Catch GSL-exceptions.

- Changed output format
- Version 0.24 (January 18, 2018)
 - Not quite sure what changed.
- Version 0.25 (January 18, 2018)
 - Fixed bug in computation of the λ -term in `foldedEpsi`.
- Version 0.26 (January 19, 2018)
 - Set $\lambda = 0$ and add levels until negative population sizes appear. This is fast and appears to be effective.
- Version 0.27 (January 24, 2018)
 - Output number of polymorphic and monomorphic sites surveyed.
- Version 0.28 (???)
- Version 0.29 (January 28, 2018)
 - Fixed missing resetting of times during iteration over files.
 - Removed superfluous option for step-wise output (`-s`).
 - Output name of input file.
 - Removed inclusion of `mpfr.h` from `epos.c`.
 - Removed search for the initial level to add; by definition this must be 2, i. e. one population size for the entire coalescent.
- Version 0.30 (January 31, 2018)
 - Fixed computation of the mutation rate. Previously I multiplied the per site mutation rate with the number of monomorphic positions. Now it is mutated by the number of all positions.
 - Added bootstrapping.
- Version 0.31 (February 1, 2018)
 - Computation of mutation rate was correct in previous version, after all, so reverted to that.
- Version 0.32 (February 7, 2018)
 - Introduced the `-m` switch.
- Version 0.33 (February 7, 2018)
 - Fixed δ computation in function `delta` in `util.c`. This reduces the computation for $m = 1$ to Watterson's estimator, as expected.
- Version 0.34 (February 9, 2018)
 - Reintroduced unfolded spectrum (`-u`) and compared to the equations in Peter's memo of December 19, 2017.
 - Reintroduced λ and set it by default to 10^{-7} .
 - Reintroduced δ and set it by default to 0.0.
 - Fixed sample size computation at the end of `sfs.c`.
- Version 0.35 (February 9, 2018)
 - Introduced estimation of $\lambda = 1./\mu/\text{args->f}$.

- Version 0.36 (February 14, 2018)
 - Changed setting of λ to $\lambda = \mu \times \text{args} \rightarrow f$. By default $\text{args} \rightarrow f = 1$.
- Version 0.37 (February 16, 2018)
 - Fixed sample size computation for folded/even in function `getSfs` in `sfs.c`.
- Version 0.38 (February 16, 2018)
 - Included working `-m` switch.
- Version 0.39 (February 19, 2018)
 - Fixed passing of λ in iterated runs.
 - Fixed numerical underflow when multiplying with λ in `foldedEpsi` in `foldedE.c`.
 - Fixed numerical underflow when multiplying with λ in `psi` in `unfolded.c`.
 - Included check for positive Ψ in both cases.
 - Expanded verbose output.
- Version 0.40 (February 21, 2018)
 - Fixed error in `foldedEpsi` in `foldedE.c`
 - Removed `if (m > 1)` from `getCoeffMat` in `foldedE.c`.
 - Switched `n/2` in `foldedEpsi` to `n/2..`
 - Ensured that `sfs->u` is always set to a value in `getSfs`.
 - Replace `u = args->u` by `u = sfs->u` in `getCoeffMat` in `unfolded.c`.
 - Re-implemented `foldedEpsi` in `foldedE.c`
- Version 0.41 (February 22, 2018)
 - Changed `4.*u*(n/2.)` to `4.*u/(n/2.)` in `foldedEpsi` in `foldedE.c`. This was a bug in the computation of Ψ for the folded/even case.
- Version 0.42 (February 22, 2018)
 - Removed line `prevMinPsi = DBL_MAX` in `foldedE` in `foldedE.c`.
- Version 0.43 (February 22, 2018)
 - Added diagnostic output in case negative population sizes are found.
- Version 0.44 (February 22, 2018)
 - Ψ now also reported if only one level is included.
- Version 0.45 (February 23, 2018)
 - Included the `-n` option to allow negative population sizes.
- Version 0.46 (February 23, 2018)
 - Fixed `if (change > args->d)`-phrase in `unfolded` and `foldedE`. This lacked re-computation of the population sizes with the best new level added, and assignment of Ψ .
- Version 0.47 (February 26, 2018)
 - Reorganized code to remove duplication. The searching for best population sizes is now done in only one place, `getPopSizes` in `popSizes.c`.

- Added printing of intermediate population sizes if the `-V` option is used.
- Version 0.48 (March 2, 2018)
 - Multi-threaded version.
- Version 0.49 (March 6, 2018)
 - Reverted to single-threaded behavior by removing `-t` from the options list and setting it to 1 in the background. This avoids the occasional race-conditions observed with the multi-threaded version.
- Version 0.50 (March 14, 2018)
 - Find number of levels through cross-validation (`-c`).
- Version 0.51 (March 14, 2018)
 - Find lambda through cross-validation (`-L`).
- Version 0.52 (March 17, 2018)
 - Include reporting of negative population sizes in verbose output (`-V`).
 - Changed `prevMinPsi = prevMinPsi`; in `getPopSizes` to `prevMinPsi = currMinPsi`;
- Version 0.53 (April 7, 2018)
 - Always allow negative population sizes.
 - Cross-validation by default.
 - $\lambda = 0$ by default.
 - If negative population sizes are found, the program searches for optimal λ by going through $\lambda = 0..\mu$. This is slow and would need to be optimized in future versions.
 - Added Scripts for extracting quantiles from `epos` output.
- Version 0.54 (April 11, 2018)
 - Fixed array out-of-bounds error in `shuffleArr` in `sfs.c`.
- Version 0.55 (April 12, 2018)
 - “Unfolded” mode not working; so I removed that option for now.
- Version 0.56 (May 31, 2018)
 - Fixed Error in documentation.
- June 13, 2018
 - Posted `epos` on `github`.