gfp_gaussian_process

Compile

cd src; make

Run

cd bin

./gfp_gaussian <csv file created by moma>

The columns that are taken from the csv file can be set by modifying csv_config.txt, although defaults are set.

The parameter space is defined in parameter_bound.txt.

Notes gfp_gaussian_process

Libraries

- Minimization: nlopt
 - o can be installed via cmake
 - can be statically compiled easily
- Linear algebra: Eigen
 - o available via modules

TODO: Likelihood calculation

- ✓ mean and covariance matrix at cell division, log_likelihood, posterior
- ✓ set initial guess for mean and covariance matrix, (not really tested yet)
- prior distribution for the next cell state: calculation of mean and covariance matrix

- get a simulated data set as reference and test everything
- ☐ decide on output

Structure

```
CSVconfig config("csv_config.txt");
Parameter_set params("parameter_bounds.txt");
std::cout << params << "\n";</pre>
// Read data
std::string infile = argv[1];
if(! std::__fs::filesystem::exists(infile)){
    std::cout << "File " << infile << " not found! \nQuit" << std::endl;</pre>
    return 0;
}
std::vector<MOMAdata> cells = getData(infile,
                                         config.time_col,
                                         config.length col,
                                         config.fp_col,
                                         config.delm);
// genealogy
build cell genealogy(cells);
print_cells(cells);
// minimization for tree starting from cells[0]
minimize_wrapper(&total_likelihood, cells[0], params);
// DEMO on how this works for all trees
// get the "trees" starting from all root cells
std::vector<MOMAdata *> root_cells = get_roots(cells);
std::vector<double> dummy_params;
for(long j=0; j<root_cells.size(); ++j){</pre>
    print_generation_tree(dummy_params, *root_cells[j]);
}
```

Likelihhod Calculation

- apply function recursively
- every cell is accessed once and after its parent is calculated

```
/* applies the function func to the cell cell and the other cells in the genealogy
    * such that the parent cell has already been accessed when the function is applied
    * to the cell.
    * Example (number implies the order in which)
              1
void likelihood_recr(const std::vector<double> & params_vec, MOMAdata *cell, double
   /*
    * Recursive implementation that applies the function func to every cell in the genea
    * not meant to be called directly, see wrapper below
    */
    if (cell == nullptr)
        return;
    sc_likelihood(params_vec, *cell, total_likelihood);
    likelihood recr(params vec, cell->daughter1, total likelihood);
    likelihood_recr(params_vec, cell->daughter2, total_likelihood);
}
double total_likelihood(const std::vector<double> &params_vec, std::vector<double> &grad
    * total_likelihood of cell tree, to be maximized
    */
    // MOMAdata cell = *(MOMAdata *) c;
    double total_likelihood = 0;
    likelihood_recr(params_vec, (MOMAdata *) c, total_likelihood);
    return total_likelihood;
```

Minimizer

• nlopt

with connfig file containing:

- bound variable: gamma_lambda
- free variable: var_lambda
- fixed varibale mean_q

Current minimizer: COBYLA

- Constrained Optimization By Linear Approximation (COBYLA)
- Implementation of Powell's method:
 - pick initial x0 and two directions h1, h2
 - starting from x0 1D optimization along first direction h1 -> find x1
 - starting from x1 1D optimization along first direction h2 -> find x2
 - h3 connects x0 and x2
 - starting from x2 1D optimization along first direction h3 -> find x3
 - o ...

Parameters

- Growth rate fluctualtions params:
 - mean_lambda;
 - gamma_lambda;
 - var_lambda;
- gfp fluctuation params
 - mean_q;
 - gamma_q;
 - var_q;
 - beta:
- variance guess for length and gfp
 - var_x;
 - var_g;

- cell division:
 - var_dx;
 - var_dg;