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//
// Iterative PBE Solver
// -----
// Solve discrete Smoluchowski Eqn with in/outflow and coagulation
// described by constant, additive and multiplicative kernels (analytic)
// in addition to a range of more physically realistic kernels.
//
// A. J. Smith (ajs224@cam.ac.uk)
//
// V4.5
//
//-----/

// To run with inflow rate=outflow rate=1, const kernel, 16 outer loops and a max cluster size of 2^10 use:
// time ./pbSolverIterative -alpha 1 -k constant -loops 16 -p 10
// To run with the additive kernel and default in/outflow rates use:
// time ./pbSolverIterative -k additive -loops 256 -p 20
// In order to achieve convergence with the additive (and multiplicative?) kernels lower the inflow rate:
// time ./pbSolverIterative -alpha 0.1 -k additive -loops 256 -p 10
//time ./pbSolverIterative -alpha 0.1 -k multiplicative -loops 64 -p 16

#include <iostream>
#include <sstream>
#include <fstream>
#include <iomanip>
#include <cstring>
#include <cstdlib>
#include <cmath>
//#include "mfa_functions.h"

using namespace std;
//using namespace mfa;

namespace mfaAnalytic
{
    enum kernelTypes { constant, additive, multiplicative, continuum, freemolecular, kinetic, shearlinear, shearnonlinear, settling, inertiasettling, berry, condensation, spmtest};

    kernelTypes kernelType; // Read from command line with -k flag. Default is constant

    const int noMoments=4; // Number of moments to compute
    const string dataDir="data/"; // Output directory

} //namespace mfaAnalytic

// Coagulation kernel header
double k(unsigned long i, unsigned long j);

// Coagulation kernel definition
double k(unsigned long i, unsigned long j)
{
    using namespace mfaAnalytic;

    switch (kernelType)
    {
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    case continuum: // Brownian motion (continuum regime)
        return (pow(i,1e0/3e0)+pow(j,1e0/3e0))*(pow(i,-1e0/3e0)+pow(j,-1e0/3e0));
        break;
    case freemolecular: // Brownian motion (free molecular regime)
        return pow(pow(i,1e0/3e0)+pow(j,1e0/3e0),2e0)*pow(pow(i,-1e0/3e0)+pow(j,-1e0/3e0)
,1e0/2e0);
        break;
    case kinetic: // Based on kinetic theory
        return (pow(i,1e0/3e0)+pow(j,1e0/3e0))*pow(i*j,1e0/2e0)*pow(i+j,-3e0/2e0);
        break;
    case shearlinear: // Shear (linear velocity profile)
        return pow(pow(i,1e0/3e0)+pow(j,1e0/3e0),3e0);
        break;
    case shearnonlinear: // Shear (nonlinear velocity profile):
        return pow(pow(i,1e0/3e0)+pow(j,1e0/3e0),7e0/3e0);
        break;
    case settling: // Gravitational settling
        return pow(pow(i,1e0/3e0)+pow(j,1e0/3e0),2e0)*abs(pow(i,1e0/3e0)-pow(j,1e0/3e0));
        break;
    case inertiasettling: // Inertia and gravitational settling
        return pow(pow(i,1e0/3e0)+pow(j,1e0/3e0),2e0)*abs(pow(i,2e0/3e0)-pow(j,2e0/3e0));
        break;
    case berry: // Analytic approximation of Berry's kernel
        return pow(i-j,2e0)*pow(i+j,-1e0);
        break;
    case condensation: // Condensation and/or branched-chain polymerisation
        return (i+2)*(j+2); // with constant c=2
        break;
    case additive:
        return i+j;
        break;
    case multiplicative:
        return i*j;
        break;
    case spmtest:
        return pow(i*j,1e0/3e0);
        break;
    default: // constant kernel
        return 1e0;
}
}

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int main(int argc, char *argv[])
{

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    using namespace mfaAnalytic;

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    unsigned int p=16; // Maximum cluster size  $N=2^p$ , default is 16 (over-ridden with the
-p flag)

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    // Sometimes we can improve the convergence by doing more than  $\log_2(N)$  iterations (ca
reful not to do too many though--div 0!)

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    unsigned int outerItLoops=4; // Try increasing this to around 256 for non constant ke
rnels (override with -loops flag)

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    kernelType=constant; // default kernel type

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    string kernelName;
    double * moments = new double[noMoments];

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    double alpha, beta; // In and outflow factors

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// Default values
alpha=1e-1;
beta=alpha;

bool numberDensityRep=true;

// Output blurb
cout << endl;
cout << "Iterative PBE Solver - A. J. Smith (ajs224@cam.ac.uk)" << endl;
cout << endl;
cout << "This code solves the discrete Smoluchowski equation with in/outflow" << endl;
;
cout << "and coagulation described by constant, additive and multiplicative kernels "
<< endl;
cout << "(admitting analytic solutions) in addition to a range of more physically " <
< endl;
cout << "realistic kernels (run with --help for additional information)." << endl;
cout << endl;

// Process command line arguments
for (int i=1; i<argc; ++i)
{
    if (strcmp(argv[i], "--help") == 0)
    {
        //cout << "This is the help message" << endl;
        cout << "Usage: " << argv[0] << " <flags>" << endl << endl;
        cout << "where <flags> is one or more of:" << endl << endl;
        cout << "\t" << "-alpha" << "\t\t" << "inflow factor (default is 1/10)" << en
dl;
        cout << "\t" << "-beta" << "\t\t" << "outflow factor (=alpha if omitted)" <<
endl;
        cout << "\t" << "-p" << "\t\t" << "power p, appearing in the maximum cluster
size, N=2^p (default is 16)" << endl;
        cout << "\t" << "-loops" << "\t\t" << "Can sometimes improve the convergence
by doing more than log2(N) " << endl;
        cout << "\t\t\titerations (careful not to do too many though! -- default is 4
)" << endl;
        cout << "\t" << "-mass" << "\t\t" << "Solves the PBE in mass flow form for mo
re direct comparison with" << endl;
        cout << "\t\t\tstochastic algorithms (default is in terms of number density)"
<< endl;
        cout << "\t\t\tN.B. In this case we expect m0=1, otherwise we expect m1=1 (us
eful convergence check)" << endl;
        cout << "\t" << "-k <type>" << "\t" << "kernel type, where <type> is one of:"
<< endl << endl;
        cout << "\t\t" << "constant" << "\t\t" << "constant kernel (default)" << endl
;
        cout << "\t\t" << "additive" << "\t\t" << "additive" << endl;
        cout << "\t\t" << "multiplicative" << "\t\t" << "multiplicative" << endl;
        cout << "\t\t" << "continuum" << "\t\t" << "Brownian motion (continuum regime
)" << endl;
        cout << "\t\t" << "freemolecular" << "\t\t" << "Brownian motion (free molecu
lar regime)" << endl;
        cout << "\t\t" << "kinetic" << "\t\t\t" << "Based on kinetic theory" << endl;
        cout << "\t\t" << "shearlinear" << "\t\t" << "Shear (linear velocity profile)
" << endl;
        cout << "\t\t" << "shearnonlinear" << "\t\t" << "Shear (nonlinear velocity pr
ofile)" << endl;
        cout << "\t\t" << "settling" << "\t\t" << "Gravitational settling" << endl;
        cout << "\t\t" << "inertiassetling" << "\t\t" << "Inertia and gravitational s
ettling" << endl;
        cout << "\t\t" << "berry" << "\t\t\t" << "Analytic approximation of Berry's k

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ernel" << endl;
    cout << "\t\t" << "condensation" << "\t\t" << "Condensation and/or branched-c
hain polymerisation" << endl;
    cout << "\t\t" << "sptest" << "\t\t\t" << "Kernel used to test the Single Pa
rticle Method (SPM)" << endl;
    cout << endl;
    cout << "Examples:" << endl << endl;
    cout << "* To run with inflow rate=outflow rate=1, const kernel, 16 outer loo
ps and a max cluster size of 2^10 use:" << endl;
    cout << "\tttime "<< argv[0] << " -alpha 1 -loops 16 -p 10" << endl;
    cout << "* To run with the additive kernel and default in/outflow rates use:"
<< endl;
    cout << "\tttime "<< argv[0] << " -k additive -loops 256 -p 20" << endl;
    cout << "* In order to achieve convergence with more complicated kernels lowe
r the inflow rate:" << endl;
    cout << "\tttime "<< argv[0] << " -alpha 0.1 -k multiplicative -loops 64 -p 16
" << endl;
    cout << "\tttime "<< argv[0] << " -alpha 0.05 -k freemolecular -loops 64 -p 16
" << endl;
    cout << endl;
    return 0;
}
else if (strcmp(argv[i], "-alpha") == 0)
{
    // Read inflow factor
    alpha = atof(argv[++i]); // default 1/10
    beta=alpha;
}
else if (strcmp(argv[i], "-beta") == 0)
{
    // Read outflow factor
    // If omitted inflow=outflow rate
    beta = atof(argv[++i]); // default 2
}
else if (strcmp(argv[i], "-mass") == 0)
{
    // Solves the equation in mass flow form
    numberDensityRep=false;
}
else if (strcmp(argv[i], "-k") == 0)
{
    // read constants appearing in multiplicative kernel  $k(x,y)=c*x^a*y^b$ 
    // c=argv[++i];
    // a=argv[++i];
    // b=argv[++i];

    // Just read one of the 3 basic kernel types with analytic solution for now
    char *kArg=argv[++i];
    if (strcmp(kArg, "additive") == 0)
        kernelType=additive;
    else if (strcmp(kArg, "multiplicative") == 0)
        kernelType=multiplicative;
    else if (strcmp(kArg, "continuum") == 0)
        kernelType=continuum;
    else if (strcmp(kArg, "freemolecular") == 0)
        kernelType=freemolecular;
    else if (strcmp(kArg, "kinetic") == 0)
        kernelType=kinetic;
    else if (strcmp(kArg, "shearlinear") == 0)
        kernelType=shearlinear;
    else if (strcmp(kArg, "shearnonlinear") == 0)
        kernelType=shearnonlinear;
    else if (strcmp(kArg, "settling") == 0)

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        kernelType=settling;
    else if (strcmp(kArg, "inertiasettling") == 0)
        kernelType=inertiasettling;
    else if (strcmp(kArg, "berry") == 0)
        kernelType=berry;
    else if (strcmp(kArg, "condensation") == 0)
        kernelType=condensation;
    else if (strcmp(kArg, "spmtest") == 0)
        kernelType=spmtest;
    else
        kernelType=constant; // actually this is default anyway
    }
else if (strcmp(argv[i], "-p") == 0)
{
    // Read p, where the maximum cluster size,  $N=2^p$ 
    p = atoi(argv[++i]); // default 16
}
else if (strcmp(argv[i], "-loops") == 0)
{
    // Read number of outer convergence loops
    outerItLoops = atoi(argv[++i]); // default 4
}
}

const unsigned long N=pow(2,p); // Max particle size, i.e.,  $i < N$  in  $n_i$ . # of particle
s is sum_i  $n_i$ 
// I choose a power of 2, because for pure coagulations the cluster sizes double with
each iteration
// so we can't do more than  $\log_2(N)=p$  iterations before gelation occurs.
const int L=outerItLoops*floor(log2(N)); // Number of iterations to perform

// Find out which kernel type is selected
switch (kernelType)
{
    case continuum:
        kernelName="continuum";
        break;
    case freemolecular:
        kernelName="freemolecular";
        break;
    case kinetic:
        kernelName="kinetic";
        break;
    case shearlinear:
        kernelName="shearlinear";
        break;
    case shearnonlinear:
        kernelName="shearnonlinear";
        break;
    case settling:
        kernelName="settling";
        break;
    case inertiasettling:
        kernelName="inertiasettling";
        break;
    case berry:
        kernelName="berry";
        break;
    case condensation:
        kernelName="condensation";
        break;
    // Analytic kernels
    case additive:
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    kernelName="additive";
    break;
case multiplicative:
    kernelName="multiplicative";
    break;
case spmtest:
    kernelName="spmtest";
    break;
default: // constant kernel
    kernelName="constant";
}

// Set up output file streams
ofstream outputFile;
ofstream momentsFile;
//ofstream diamsFile;

string outputFileName=dataDir+kernelName+"_data_";
string momentsFileName=dataDir+kernelName+"_moments_";
//string diamsFileName=dataDir+"diameters_";
string ext=".txt";

string desc;
stringstream out;

// For simplicity let's index stuff from 1 - N instead of 0 - (N-1)
double *n = new double[N+1]; // Allocate N ints and save ptr in n. Requires  $2^{p+3-1}$ 
0} Mb, i.e., ~8Gb when p=20.
double *nold = new double[N+1];
//double K[N][N];
double d, summa;

double n_in;

//vector<double> xDiam (N);

// Initialise PSD to a delta delta_{i1}, i.e., mono-dispersed
for (unsigned long i=1; i<=N; i++)
{
    n[i] = 0e0;
}
n[1]=1e0;

//out << N;
out << "p" << p << "_alpha" << alpha << "_beta" << beta << "_loops" << outerItLoops;
desc=out.str();

outputFileName+=desc+ext;
momentsFileName+=desc+ext;
//diamsFileName+=desc+ext;

outputFile.open(outputFileName.c_str(), ios::out);
momentsFile.open(momentsFileName.c_str(), ios::out);
//diamsFile.open(diamsFileName.c_str(), ios::out);

cout << "Running iterative solver (";
if(numberDensityRep)
    cout << "in terms of number density";
else
    cout << "in massflow form";
cout << ") with maximum particle size of " << N << " and " << L << " iterations." <<
endl << endl;
kernelName[0]=toupper(kernelName[0]); // capitalise

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cout << kernelName << " kernel selected." << endl;
cout << "Inflow rate (1/alpha):" << 1/alpha << endl;
cout << "Outflow rate (1/beta):" << 1/beta << endl;
cout << endl;

cout.precision(10);
cout.width(20);
//cout.fill('0');
//cout.setf(ios::showpos);
cout.setf(ios::scientific);
outputFile.precision(8);
momentsFile.precision(8);

// Output header
cout << "Iter\tm0\t\t\tm1\t\t\tm2\t\t\tm3" << endl;

int l=1;

// Iterate L times
while (l<=L)
{
    // Let's compute the moments of the distribution
    for(int moment=0;moment<noMoments;moment++)
    {
        moments[moment]=0e0;
        for(unsigned long i=1;i<=N;i++)
        {
            moments[moment]+=pow(i,moment)*n[i];
        }
    }

    /*
    for(int i=0;i<N;i++)
    {
        cout << n[i] << " ";
    }
    cout << endl;

    system("sleep 0");
    */

    cout << l << "\t" << moments[0] << "\t" << moments[1] << "\t" << moments[2] << "\t"
    << moments[3] << endl;
    momentsFile << l << "\t" << moments[0] << "\t" << moments[1] << "\t" << moments[2]
    << "\t" << moments[3] << endl;

    // Update old distribution to new distribution
    for(unsigned long i=1;i<=N;i++)
    {
        nold[i]=n[i];
    }

    for(unsigned long i=1;i<=N;i++) // Loop over N particle sizes
    {
        // Compute sums in numerator and denominator
        d=0e0;
        for(unsigned long j=1;j<=N;j++)
        {
            if(numberDensityRep)
                d+=k(i,j)*nold[j];
            else

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        d+=k(i,j)*nold[j]/j;
    }

    summa=0e0;
    for(unsigned long j=1;j<=i-1;j++)
    {
        //summa+=K[i-j][j]*nold[i-j]*nold[j];
        if(numberDensityRep)
            summa+=k(i-j,j)*nold[i-j]*nold[j];
        else
            summa+=k(i-j,j)*nold[i-j]*nold[j]/j;
    }

    //cout <<"i="<<i<<" , d="<<d<<" , summa="<<summa<<endl;

    // Evaluate n_in (here delta_{i1}), i.e., clusters of size 1 flow into the do
main
    if(i==1)
        n_in=1e0;
    else
        n_in=0e0;

    if(numberDensityRep)
        summa*=0.5;

    // Iterate baby!
    //n[i]=(n_in/alpha+0.5*summa)/(1e0/beta+d);
    n[i]=(n_in/alpha+summa)/(1e0/beta+d);

    //n[i]=0.5*summa/d; // Pure coagulation
}

l++; // Update iteration counter

}

// Dump steady-state PSD
for(unsigned long i=1;i<=N;i++) // Loop over N particle sizes
{
    outputFile << i << "\t" << n[i] << endl;
}

// Close files
outputFile.close();
momentsFile.close();

// Clean up memory like a good boy
delete [] n;
n = NULL;

delete [] nold;
nold = NULL;

delete [] moments;
moments = NULL;

//delete [][] K;
//K=NULL;

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    return 0;  
}
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