```
#include <stdio.h>
#include <stdlib.h>
#include <string.h>
#include <math.h>
#include "data structures.h"
#include "subroutines.h"
#include "InOut.h"
int main( void )
/***********************
*******
********
******************
********
  int lambda = 1;  // Parameter describing geometry of system (=1
for planar, =3 for spherical)
  double s_0 = 1.; // Initial interface position
  double R = 5.; // Position of far boundary
  int nAlpha = 100;
                      // Number of points in phase alpha
  double dAlpha = 1E-7;  // Diffusion coefficient of phase
alpha
  double initialAlpha = 0.8; // Initial concentration in phase
  double interAlpha = 0.6; // Interfacial concentration in phase
alpha
  int nBeta = 100;
  double dBeta = 1E-5;
  double initialBeta = 0.4;
  double interBeta = 0.0;
  double time step=0.1;
  int n time steps=10;  // As written, output written to maximum
100000000 steps
  double tol = 1.E-8;  // Used to determine whether linearisation
converges at each timestep
/**********************
*******
******** ENDS
**********
```

```
***********************
*********
   FILE *fpt;
   int i, tmp;
   two phase *whole system;
   whole system =(two phase *) calloc (1, sizeof(two phase));
   whole system->s = s \ 0;
   whole system->1 = R;
   whole system->old s = whole system->s;
   whole system->future s = whole system->s;
   // Alpha is on left
   whole system->left =(single phase *) calloc (1,
sizeof(single phase));
   whole system->left->n = nAlpha;
   whole system->left->d coeff = dAlpha;
   whole system->left->c boundary = interAlpha;
   whole system->left->u = (double *) calloc (whole system->left->n,
sizeof(double));
   whole system->left->c =(double *) calloc (whole system->left->n,
sizeof(double));
   whole system->left->future c = (double *) calloc (whole system-
>left->n, sizeof(double));
   for (i=0; i<whole system->left->n; i++)
   whole system->left->u[i] = double(i)/double(whole system->left->n
- 1);
   whole system->left->c[i] = initialAlpha;
   whole system->left->future c[i] = whole system->left->c[i];
   whole system->left->c[whole system->left->n-1] = whole system-
>left->c boundary;
   whole system->left->future c[whole system->left->n-1] =
whole system->left->c boundary;
   // Beta is on right
   whole system->right =(single phase *) calloc (1,
sizeof(single phase));
   whole system->right->n = nBeta;
   whole system->right->d coeff = dBeta;
   whole system->right->c boundary = interBeta;
```

```
whole system->right->u =(double *) calloc (whole system->right->n,
sizeof(double));
    whole system->right->c =(double *) calloc (whole system->right->n,
sizeof(double));
    whole system->right->future c = (double *) calloc (whole system-
>right->n, sizeof(double));
    for (i=0; i<whole system->right->n; i++)
    whole system->right->u[i] = double(i)/double(whole system->right-
>n - 1);
    whole system->right->c[i] = initialBeta;
    whole system->right->future c[i] = whole system->right->c[i];
    whole system->right->c[0] = whole system->right->c boundary;
    whole system->right->future c[0] = whole system->right-
>c boundary;
    fpt=fopen("results.txt","w");
    fprintf(fpt, "Time\tInterface Position\n");
    /**** Looping over time (for the given timestep) ****/
    for (i=0; i<n time steps+1; i++)</pre>
    {
        if(i<100)</pre>
            {out interface(whole system, double(i)*time step, fpt);}
        else if((i<1000) && (i%10 == 0))
            {out interface(whole system, double(i)*time step, fpt);}
        else if((i<10000) && (i%100 == 0))
            {out interface(whole system, double(i)*time step, fpt);}
        else if((i<100000) && (i%1000 == 0))
            {out interface(whole system, double(i)*time_step, fpt);}
        else if((i<1000000) && (i\%10000 == 0))
            {out interface(whole system, double(i)*time step, fpt);}
        else if((i<10000000) && (i%100000 == 0))</pre>
            {out interface(whole system, double(i)*time step, fpt);}
        else if((i<100000000) && (i%1000000 == 0))</pre>
            {out interface(whole system, double(i)*time_step, fpt);}
        if(lambda == 1)
            {tmp = take step planar(whole system, time step, tol);}
        else if(lambda == 3)
            {tmp = take_step_spherical(whole system, time step, tol);}
        else
            {printf("\nProblem with geometry\n\n");}
        if (tmp<0)</pre>
                   // Interface has moved beyond the end of the
system
            {printf("Finished at time = %lg", double(i)*time step);
fclose(fpt); return 0;}
```

```
if(i%1000 == 0)
            {printf("\nTimestep %d complete\n", i);}
        else if (i%10 == 0)
            {printf(".");}
    }
    free (whole system->left->u);
    free(whole system->left->c);
    free(whole system->left->future c);
    free(whole system->right->u);
    free(whole system->right->c);
    free(whole system->right->future c);
    free(whole system->left);
    free (whole system->right);
    fclose(fpt);
   printf("\n \n");
    return 1;
}
#include <stdio.h>
#include <stdlib.h>
#include <string.h>
#include "data structures.h"
#include "InOut.h"
void out profile(two phase *tp, FILE *fp, double time)
       // Prints:
            // time
            // x[0], c(x[0]),
            // x[1], c(x[1]),
            // ....
                       . . . .
            // x[n],
                      c(x[n]),
            // x[n+1], c(x[n+1]),
            // x[x+2], c(x[n+2]),
            // ...
            // x[n+N], c(x[n+N]),
            //
    int i;
    fprintf(fp, "%lg,\n", time);
    for (i=0; i<(tp->left->n); i++)
```

```
{
    fprintf(fp, "%lg,\t", tp->s*tp->left->u[i]);
    fprintf(fp, "%lg,\n", tp->left->c[i]);
}

for(i=0; i<(tp->right->n); i++)
{
    fprintf(fp, "%lg,\t", tp->s + (tp->l-tp->s)*tp->right->u[i]);
    fprintf(fp, "%lg,\n", tp->right->c[i]);
}
fprintf(fp, "\n");
}

void out_interface(two_phase *tp, double time, FILE *fp)
{
    fprintf(fp, "%lg\t%lg\n", time, tp->s);
}
```

```
//note: For first (implicit step), we require some first estimates
for solution variables
            // 1) the elements in future c to take the values of
'current ' c
            // 2) the value of future s to be the same as s
    double v;
    double diff 1, diff r;
    double rhs, lhs;
    if(pass nmb==0)
                               // FIRST PASS -
    tr(pass_inno==0)
{v=tp->s-tp->old_s;}
                                    // Use old interface position to
determine sign of interface velocity
    else
    {v=tp->future s-tp->s;} // SECOND (or higher) PASS -
                                    // Use estimate of future position
to determine sign of interface velocity
    diff l=(tp->left->c boundary - tp->left->future c[tp->left->n-
2])/(1. - tp->left->u[tp->left->n-2]);
    diff l= diff l*tp->left->d coeff / tp->future s;
    diff r=(tp->right->future c[1] - tp->right->c boundary)/(tp-
>right->u[1]);
    diff r= diff r*tp->right->d coeff / (tp->l-tp->future s);
    rhs = (diff r - diff l)*time step;
    if(v>=0)
                        //POSITIVE VELOCITY - Use points to the right:
    {
        lhs = tp->left->c boundary;
        lhs = lhs - tp->right->future c[1]*(1-tp->right->u[1]/2.);
        lhs = lhs - tp->right->c boundary*tp->right->u[1]/2.;
    }
                        //NEGATIVE VELOCITY - Use points to the left:
    else
        lhs = tp->left->future c[tp->left->n-2] * (0.5+tp->left->u[tp-
>left->n-2]/2.);
        lhs = lhs + tp->left->c boundary * (0.5-tp->left->u[tp->left-
>n-2]/2.);
        lhs = lhs - tp->right->c boundary;
    }
    tp->future s=tp->s + rhs/lhs;
   return 0;
}
```

```
/*************
   Given the concentration profiles and interface position at
timestep j and estimates for the interface at j+1,
   finds concentrations at j+1 in phase to the left of the boundary
void new concentration left planar(two phase *tp, double time step)
   int i;
   double tmpA, tmpB, left diff, right diff, left sum, right sum;
   trimatrix system tms;
   tms.dim = tp->left->n;
// RESERVE MEMORY FOR tms
   tms.lo = (double *) calloc(tms.dim, sizeof(double));
   tms.diag = (double *) calloc(tms.dim, sizeof(double));
   tms.up = (double *) calloc(tms.dim, sizeof(double));
   tms.rhs = (double *) calloc(tms.dim, sizeof(double));
   tms.c = tp->left->future c;
// FILL tms
   tmpA=tp->left->d coeff*time step / tp->future s;
   tmpB=(tp->future s-tp->s);
   if(tp->future s >= tp->s)
                                         // POSITIVE VELOCITY - use
points to the right:
       tms.lo[0] = 0.;
       tms.diag[0]=-tmpA/tp->left->u[1] - tp->future s*tp->left-
>u[1]/2.;
       tms.up[0] = tmpA/tp->left->u[1] + tmpB*tp->left->u[1]/2.;
       tms.rhs[0] =-tp->left->c[0]*tp->s*tp->left->u[1] / 2.;
       for(i=1; i<(tms.dim-1); i++)</pre>
           left diff = tp->left->u[i] - tp->left->u[i-1];
           left sum = tp->left->u[i] + tp->left->u[i-1];
           right diff= tp->left->u[i+1] - tp->left->u[i];
           right sum = tp->left->u[i+1] + tp->left->u[i];
           tms.lo[i] = tmpA/left diff;
           tms.diag[i]=-tmpA*(1/left diff + 1/right diff) -
tmpB*left sum/2.
                       - tp->future s*(right sum-left sum)/2.;
           tms.up[i] = tmpA/right diff + tmpB*right sum/2.;
           tms.rhs[i] =-tp->s*tp->left->c[i]*(right sum-left sum)/2.;
       }
   }
```

```
else
                                                                                                               // NEGATIVE VELOCITY - use
points to the left:
                    tms.lo[0] = 0.;
                    tms.diag[0] = -tmpA/tp - left - u[1] + tmpB*tp - left - u[1]/2. - tmpB*tp - u[1]/
tp->future s*tp->left->u[1]/2.;
                    tms.up[0] = tmpA/tp->left->u[1];
                    tms.rhs[0] =-tp->left->c[0]*tp->s*tp->left->u[1] / 2.;
                    for(i=1; i<(tms.dim-1); i++)</pre>
                              left diff = tp->left->u[i] - tp->left->u[i-1];
                              left sum = tp->left->u[i] + tp->left->u[i-1];
                              right diff= tp->left->u[i+1] - tp->left->u[i];
                              right sum = tp->left->u[i+1] + tp->left->u[i];
                              tms.lo[i] = tmpA/left diff - tmpB*left sum/2;
                              tms.diag[i]=-tmpA*(1/left diff + 1/right_diff) +
tmpB*right sum/2.
                                                            - tp->future s*(right sum-left sum)/2.;
                              tms.up[i] = tmpA/right diff;
                              tms.rhs[i] =-tp->s*tp->left->c[i]*(right sum-left sum)/2.;
                    }
          }
          tms.lo[tms.dim-1] = 0.;
          tms.diag[tms.dim-1]=-1.;
          tms.up[tms.dim-1] = 0.;
          tms.rhs[tms.dim-1] =-tp->left->c boundary;
// SOLVE tms
          solve trimatrix system(tms);
// FREE MEMORY RESERVED FOR tms
          free(tms.lo);
          free(tms.diag);
          free(tms.up);
          free(tms.rhs);
}
/*************
          Given the concentration profiles and interface position at
timestep j and estimates for the interface at j+1,
          finds concentrations at j+1 in phase to the right of the boundary
****************
void new concentration right planar(two phase *tp, double time step)
{
```

```
int i;
    double tmp, tmpA, tmpB, left diff, right diff, left sum,
right sum;
    trimatrix system tms;
    tms.dim = tp->right->n;
// RESERVE MEMORY FOR tms
    tms.lo = (double *) calloc(tms.dim, sizeof(double));
    tms.diag = (double *) calloc(tms.dim, sizeof(double));
    tms.up = (double *) calloc(tms.dim, sizeof(double));
    tms.rhs = (double *) calloc(tms.dim, sizeof(double));
    tms.c = tp->right->future c;
// FILL tms
    tmpA=tp->l-tp->future s;
    tmpA=tp->right->d coeff*time step/tmpA;
    tmpB=(tp->future s-tp->s);
    tms.lo[0] = 0.;
    tms.diag[0]=-1.;
    tms.up[0] = 0.;
    tms.rhs[0] =-tp->right->c boundary;
                                          // POSITIVE VELOCITY - use
    if(tp->future s >= tp->s)
points to the right:
    {
        for (i=1; i<(tms.dim-1); i++)</pre>
            left diff = tp->right->u[i] - tp->right->u[i-1];
            left sum = tp->right->u[i] + tp->right->u[i-1];
            right diff= tp->right->u[i+1] - tp->right->u[i];
            right sum = tp->right->u[i+1] + tp->right->u[i];
            tms.lo[i] = tmpA/left diff;
            tms.diag[i]=-tmpA*(1/right diff + 1/left diff) - tmpB*(1.-
left sum/2.)
                        - (tp->l-tp->future s)*(right sum-left sum)/
2.;
            tms.up[i] = tmpA/right diff + tmpB*(1. - right sum/2.);
            tms.rhs[i] = -(tp->l-tp->s)*tp->right->c[i]*(right sum-
left sum) / 2.;
        tmp=tp->right->u[tp->right->n-2];
        tms.lo[tms.dim-1] = tmpA/(1.-tmp);
        tms.diag[tms.dim-1] = -tmpA/(1.-tmp) - tmpB*(1. - (1.+tmp)/2.) -
(tp->1-tp->future s)*(1.-tmp)/2.;
        tms.up[tms.dim-1] = 0;
```

```
tms.rhs[tms.dim-1] =-tp->right->c[tp->right->n-1]*(tp->l-tp-
>s)*(1.-tmp) / 2.;
            }
           else
                                                                                                                      // NEGATIVE VELOCITY - use
points to the left:
           {
                       for (i=1; i<(tms.dim-1); i++)</pre>
                                   left diff = tp->right->u[i] - tp->right->u[i-1];
                                   left sum = tp->right->u[i] + tp->right->u[i-1];
                                   right diff= tp->right->u[i+1] - tp->right->u[i];
                                   right sum = tp->right->u[i+1] + tp->right->u[i];
                                   tms.lo[i] = tmpA/left diff - tmpB*(1.- left sum/2.);
                                   tms.diag[i]=-tmpA*(1/right diff + 1/left diff) + tmpB*(1.
- right sum/2.)
                                                                       - (tp->l-tp->future s)*(right_sum - left_sum)
/ 2.;
                                   tms.up[i] = tmpA/right diff;
                                   tms.rhs[i] = -(tp->l-tp->s)*tp->right->c[i]*(right sum-
left sum) / 2.;
                       }
                       tmp=tp->right->u[tp->right->n-2];
                       tms.lo[tms.dim-1] = tmpA/(1.-tmp) - tmpB*(1. - (1.+tmp)/2.);
                       tms.diag[tms.dim-1]=-tmpA/(1.-tmp) - (tp->l-tp->future s)*(1.-tms.diag[tms.dim-1]=-tmpA/(1.-tmp) - (tp->l-tp->future s)*(1.-tms.diag[tms.dim-1]=-tmpA/(1.-tmp) - (tp->l-tp->future s)*(1.-tms.diag[tms.dim-1]=-tmpA/(1.-tmp) - (tp->l-tp->future s)*(1.-tms.diag[tms.dim-1]=-tmpA/(1.-tmp) - (tp->l-tp->future s)*(1.-tms.diag[tms.dim-1]=-tmpA/(1.-tms.diag[tms.dim-1]=-tmpA/(1.-tms.diag[tms.dim-1]=-tmpA/(1.-tms.diag[tms.diag[tms.dim-1]=-tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms.diag[tms
tmp) /2.;
                       tms.up[tms.dim-1] = 0;
                       tms.rhs[tms.dim-1] =-tp->right->c[tp->right->n-1]*(tp->l-tp-
>s)*(1.-tmp) / 2.;
           }
// SOLVE tms
           solve trimatrix system(tms);
// FREE MEMORY RESERVED FOR tms
           free(tms.lo);
           free(tms.diag);
           free(tms.up);
           free(tms.rhs);
}
```

/*************

```
Takes system from time step j to timestep j+1
int take step planar(two phase *tp, double dt, double tolerance)
   int i, count;
   double error;
   double *tmp dbl ptr;
   /****** BOOKKEEPING
*************************
   // First prediction of interface position at j+1 is position at j
   tp->future s = tp->s;
   for(i=0; i<(tp->left->n); i++)
       {tp->left->future c[i] = tp->left->c[i];}
   for (i=0; i<(tp->right->n); i++)
       {tp->right->future c[i] = tp->right->c[i];}
   error=tolerance+1;
   count=0;
   while((error>tolerance) || (count==1)) // need at least one
implicit loop
   {
       error=tp->future s;
       // find new interface position
       new interface planar(tp, dt, count);
       // solve for concentration to left of interface:
       new concentration left planar(tp, dt);
       // solve for concentration to right of interface:
       new concentration right planar(tp, dt);
                                      // (old estimate >
       if(error>tp->future s)
new estimate)
          {error = error-tp->future s;}
                                      // (new estimate <=</pre>
       else
old estimate)
          {error = tp->future s-error;}
       count++;
   }
   /****** BOOKKEEPING
**********
   // Update s:
   tp->old s = tp->s;
```

```
tp->s = tp->future s;
    if((tp->s<0.) || (tp->s>tp->l))
    {// Interface has moved beyond the ends of the system
       printf("\n\nInterface has moved beyond the end of the
system\n");
       return -1;
    }
    else
    {// Update c:
        tmp dbl ptr = tp->left->c;
        tp->left->c
                           = tp->left->future c;
        tp->left->future c = tmp dbl ptr;
        tmp dbl ptr = tp->right->c;
        tp->right->c
                           = tp->right->future c;
        tp->right->future c = tmp dbl ptr;
        return count;
    }
}
```

```
#include <stdlib.h>
#include "trimatrix.h"

/********************************

algorithm for solving system of linear equations with tridiagonal
matrix

****************************

void solve_trimatrix_system (trimatrix_system tms)
{
    double *alf, *bet;
    int n;
    int i;

    n=tms.dim;
    alf = (double *) calloc(n+1, sizeof(double));
    bet = (double *) calloc(n+1, sizeof(double));

    alf[0]=bet[0]=0.0;
    for( i=0; i<=n-1; i++)
    {
        alf[i+1] = tms.up[i]/(-tms.diag[i] - tms.lo[i]*alf[i] );
    }
}</pre>
```

```
bet[i+1]=( tms.lo[i]*bet[i] - tms.rhs[i] )/(-tms.diag[i] -
tms.lo[i]*alf[i] );
}

tms.c[n-1] = bet[n];
for( i=n-2; i>=0; i--)
{
    tms.c[i] = alf[i+1]*tms.c[i+1]+bet[i+1];
}

free(alf);
free(bet);
}
```