```
/*
 * project: HIV quasispecies
 * file name: main.cc
 * date_created: 05-Dec-2010
 * date_modified: 10-Jul-2012
 * version: 1.3.0
*/

#include<stdio.h>
#include<cstdlib>
#include<math.h>
#include<iostream>
#include<time.h>
#include "mt19937-64.h" // Header file for Mersenne Twister random number generator

//#**** SIMULATION CONTROL ****#
// comment lines to switch on or off specific evolutionary processes
#define MUTATE_ON
#define RECOMBINE_ON
#define MULT_INF_ON //also change INF_DIST
#define DRIFT_ON
//#define FOUNDER_MUTATE_ON
#define FITNESS_SELECTION_ON

//#**** production and infection parameters ****#
#ifdef MULT_INF_ON
    #define MAXINF 3 // maximum number of infections per cell (define the specific distribution in next line)
    #define INF_DIST {0.0,0.0,100.0} // distribution of multiply infected cells
#else
    #define MAXINF 1
    #define INF_DIST {100.0}
#endif
#ifdef DRIFT_ON // multiple virions produced per cell
    #define MAXPRODUCE 10 // number of offspring viruses from each infected cell
#else
    #define MAXPRODUCE 1
#endif

//#**** simulation parameters ****#
#define MAXCYCLE 10000    // maximum number of generations
#define MAXRUNS 5 // maximum runs

//#**** system parameters ****#
#define L 10000      // Length of RNA sequence
#define MAXCELL 2400 // total number of cells
#define MAXVIRUS (MAXCELL*MAXPRODUCE) // total number of viruses

//#**** evolution parameters ****#
#ifdef MUTATE_ON // set specific mutation rate if MUTATE_ON switch is defined or set mutation rate to zero
    #define MUTATEINDEX 1.0e-3    // mutation rate (substitutions per site per replication)
#else
    #define MUTATEINDEX 0.0
#endif

#ifdef RECOMBINE_ON
    #define RECOMBINEINDEX 8.3e-4 // recombination rate (crossovers per site per replication)
#else
    #define RECOMBINEINDEX 0.0
#endif

#ifdef FOUNDER_MUTATE_ON
    #define FOUNDER_MUTATE_FRAC 0.1 // fraction of sites mutated from fittest sequence to form founder sequence
#else
    #define FOUNDER_MUTATE_FRAC 0.0
#endif

#ifdef FITNESS_SELECTION_ON   // calculate fitness of viruses if FITNESS_SELECTION_ON switch is defined or set virus fitness to 1.0
    #define VIRAL_FITNESS fitness(i)
#else
    #define VIRAL_FITNESS 1.0
#endif

```

```
73  //**** viral fitness parameters ****#
74  #define fMIN 0.2433  // minimum viral fitness
75  #define N 3.0    // Hill coefficient
76  #define d50 30.3 // Hamming distance for which viral fitness is half maximal
77
78  using namespace std;
79
80  double percnt_mult_inf[MAXINF] = INF_DIST;
81  int mult_inf[MAXINF] = {0};
82  int quasi_structure[L+1] = {0};
83  FILE *fp_quasi;
84
85  // virus class
86  class c_virus
87    {
88    public:
89      int DNA[2*L]; // array containing two viral RNA sequences of length L
90      double fitness; // fitness of virus
91      bool available; // flag indicating whether virus is available for infection
92      c_virus(); // virus constructor
93    };
94
95  // virus class constructor (Default values when a virus is created)
96  c_virus::c_virus()
97  {
98    for (int i=0;i<2*L;i++)
99      {
100       DNA[i] = 0; // set all nucleotides to A initially (0,1,2,3 corresponds to A, C, G
    and T respectively)
101     }
102   fitness = 1.0;
103   available = true;
104 }
105
106 c_virus virus[MAXVIRUS]; // array containing viruses
107 c_virus fittest_virus;
108 c_virus founder_virus;
109
110 // T-cell class
111 class c_cell:public c_virus
112   {
113   public:
114     c_virus V[MAXINF];   // array containing viruses that infected the cell
115     int provirus[MAXINF][L]; // array for proviral DNA sequence
116     int multinf; // present multiple infection status of cell
117
118     void infect(int v,int inf); // function to infect the cell by virus with index
    number 'v' at multiple infection status 'inf'
119     void mutate(); // function to mutate proviral sequence
120     void recombine(); // function to create proviral sequence from viral RNA sequences by
    recombination
121   };
122
123 c_cell cell[MAXCELL]; // array containing cells
124
125 // infect cell by virus with index number 'v' at multiple infection status 'inf'
126 void c_cell::infect(int v,int inf)
127 {
128     // copy viral RNA sequence into cell
129   for (int i=0;i<2*L;i++)
130     {
131       V[inf].DNA[i]=::virus[v].DNA[i];
132     }
133
134   multinf = multinf+1; // increase multiple infection status by 1
135   //printf("multinf = %d ",multinf);
136 }//end of c_cell::infect()
137
138 // function to mutate proviral sequence
139 void c_cell::mutate()
140 {
141   int ch_no = 2;
142   // loop over present cell's multiple infection status (equal to number of proviral
    sequences)
143   for(int i=0;i<multinf;i++)
144     {
145       // loop over proviral sequence length
```

```
146         for (int j=0;j<L;j++)
147           {
148             // if random number is less than mutation rate change the nucleotide by 2 (A
    <-> G and C <-> T : only transitions are considered)
149             if (rand_no()<MUTATEINDEX)
150               {
151                 //printf("mutation occurred for provirus %d, position %d \n",i,j);
152                 provirus[i][j] = (provirus[i][j]+ch_no)%4;
153               }
154           }
155       }
156   }//end of c_cell::mutate()
157
158
159   // function to create proviral DNA sequence from viral RNA sequence by recombination
160   void c_cell::recombine()
161   {
162     // loop over present multiple infection status
163     for (int i=0;i<multinf;i++)
164       {
165         int strand = 1;
166
167         // select first or second strand randomly
168         if (rand_no()>0.5)
169           strand = 1;
170         else
171           strand = 2;
172
173         // loop over proviral DNA sequence length
174         for (int j=0;j<L;j++)
175           {
176             // if random number is less than recombination rate switch sequence
177             if (rand_no() < RECOMBINEINDEX)
178                 {
179                   //printf("recombination occurred for provirus %d, position %d \n",i,j);
180                    if (strand == 1)
181                     strand = 2;
182                      else
183                        strand = 1;
184                 }
185
186             // copy nucleotide from selected viral RNA strand to proviral DNA sequence
187             if (strand == 1)
188               {
189                 provirus[i][j] = V[i].DNA[j];
190               }
191             else
192               {
193                 provirus[i][j] = V[i].DNA[L+j];
194               }
195           }
196       }
197   }
198
199
200   int main()
201   {
202     //int i,j,k;
203     int tempv;
204     int v1,v2;
205     int vcount;
206     int virus_prod_count = 0;
207
208     double fittest_virus_fitness = 0.0; // fitness of fittest virus in current generation
209
210
211     int time_now;
212     time_t rawtime;
213     struct tm * timeinfo;
214     cout<<"Time of Start is"<<int(time(0))<<endl;    // simulation start time
215
216     time_now = int(time(0));
217     time( &rawtime);
218     timeinfo = localtime ( &rawtime );
219     cout<<"Date :"<<asctime(timeinfo); // simulation start date and time
220
221     srand((long int)time(0)); // initialize random number generator
```

```
222
223     void display_viral_sequences();
224     void display_proviral_sequences();
225
226     double fitness(int v); // function to calculate fitness
227     void cal_quasi_structure(); // function to calculate quasispecies structure
228
229     void initialize_virus(); // clear information in all viruses
230     void initialize_cell(); // clear information in all cells
231     void initialize_fittest_virus(); // create fittest virus
232     void initialize_founder_virus(); // create founder virus
233
234     // file pointer for simulation result summary
235     FILE *fp_result;
236
237     if ((fp_result = fopen("result.txt","a+"))==NULL)
238       {
239         printf("error in opening result.txt \n");
240       }
241
242     if ((fp_quasi = fopen("quasi_structure.txt","a+"))==NULL)
243       {
244         printf("error in opening quasi_structure.txt \n");
245       }
246
247     fprintf(fp_result,"%d\nSTART: %s\n",time_now,asctime(timeinfo));
248
249     printf("L=%d, cell=%d, Maxruns=%d, Maxcycles=%d, Maxinf=%d, Maxprod=%d,
    mutateindex=%lf, recombine index=%lf, FOUNDER_MUTATE_FRAC=%lf \n",L,MAXCELL,MAXRUNS,
    MAXCYCLE, MAXINF, MAXPRODUCE, MUTATEINDEX, RECOMBINEINDEX,FOUNDER_MUTATE_FRAC);
250
251     fprintf(fp_result,"L=%d, cell=%d, Maxruns=%d, Maxcycles=%d, Maxinf=%d, Maxprod=%d,
    mutateindex=%lf, recombine index=%lf, FOUNDER_MUTATE_FRAC=%lf \n",L,MAXCELL,MAXRUNS,
    MAXCYCLE, MAXINF, MAXPRODUCE, MUTATEINDEX, RECOMBINEINDEX,FOUNDER_MUTATE_FRAC);
252
253     // display various switches' status
254     #ifdef MUTATE_ON
255       fprintf(fp_result,"MUTATION_ON\n");
256       printf("MUTATION_ON\n");
257     #endif
258     #ifndef MUTATE_ON
259       fprintf(fp_result,"MUTATION_OFF\n");
260       printf("MUTATION_OFF\n");
261     #endif
262
263     #ifdef RECOMBINE_ON
264       fprintf(fp_result,"RECOMBINATION_ON\n");
265       printf("RECOMBINATION_ON\n");
266     #endif
267     #ifndef RECOMBINE_ON
268       fprintf(fp_result,"RECOMBINATION_OFF\n");
269       printf("RECOMBINATION_OFF\n");
270     #endif
271
272     #ifdef MULT_INF_ON
273       fprintf(fp_result,"MULT_INF_ON\n");
274       printf("MULT_INF_ON\n");
275     #endif
276     #ifndef MULT_INF_ON
277       fprintf(fp_result,"MULT_INF_OFF\n");
278       printf("MULT_INF_OFF\n");
279     #endif
280
281     #ifdef DRIFT_ON
282       fprintf(fp_result,"DRIFT_ON\n");
283       printf("DRIFT_ON\n");
284     #endif
285     #ifndef DRIFT_ON
286       fprintf(fp_result,"DRIFT_OFF\n");
287       printf("DRIFT_OFF\n");
288     #endif
289
290     #ifdef FOUNDER_MUTATE_ON
291       fprintf(fp_result,"FOUNDER_MUTATE_ON\n");
292       printf("FOUNDER_MUTATE_ON\n");
293     #endif
294     #ifndef FOUNDER_MUTATE_ON
```

```
295       fprintf(fp_result,"FOUNDER_MUTATE_OFF\n");
296       printf("FOUNDER_MUTATE_OFF\n");
297     #endif
298
299     #ifdef FITNESS_SELECTION_ON
300       fprintf(fp_result,"FITNESS_SELECTION_ON\n");
301       printf("FITNESS_SELECTION_ON\n");
302     #endif
303     #ifndef FITNESS_SELECTION_ON
304       fprintf(fp_result,"FITNESS_SELECTION_OFF\n");
305       printf("FITNESS_SELECTION_OFF\n");
306     #endif
307
308     /*********************************************/
309
310     // calculate the number of cells of each multiple infected type from multiple infection
    distribution
311     mult_inf[MAXINF-1] = MAXCELL;
312     for(int i=0;i<MAXINF-1;i++)
313     {
314       mult_inf[i] = (int)rint(MAXCELL*percnt_mult_inf[i]/100);
315       printf("mult_inf[%d]=%d \n",i,mult_inf[i]);
316       mult_inf[MAXINF-1] = mult_inf[MAXINF-1]-mult_inf[i];
317     }
318
319       printf("mult_inf[%d]=%d \n",MAXINF-1,mult_inf[MAXINF-1]);
320
321       // display multiple infection cell numbers
322     for (int i=0;i<MAXINF;i++)
323     {
324       fprintf(fp_result,"mult_inf = %d \t percentage = %lf \t mult_inf_cells = %d\n",i+1,
    percnt_mult_inf[i],mult_inf[i]);
325     }
326
327     // loop over simulation runs
328     for (int run=1;run<=MAXRUNS;run++)
329       {
330         init_genrand64((unsigned long long)(rand())); //initialize Mersenne Twister random
    number generator with a random number generated from GCC random number generator
331         //printf("run =%d \n",run+1);
332
333         /***** system initialization *****/
334         initialize_virus();
335         initialize_cell();
336
337         initialize_fittest_virus(); // create fittest virus
338         initialize_founder_virus(); // create founder virus
339
340         /***** simulation start ******/
341         // loop over generations
342         for (int cycle=1;cycle<=MAXCYCLE;cycle++)
343           {
344
345             //printf("\n\n cycle start proviral sequences \n");
346             //display_proviral_sequences();
347
348             /*********** INFECTION **************/
349             //printf("infecting \n");
350              initialize_cell(); // clear all cell information to prepare for next round of
    infection
351             int cell_count = 0;
352
353             for (int i=1;i<=MAXINF;i++) // loop over multiple infected type
354               {
355                 for(int j=0;j<mult_inf[i-1];j++) // loop over cells of each multiple
    infected type
356                 {
357                 for (int inf=0;inf<i;inf++) // loop over number of multiple infections in
    each cell
358                   {
359                     do
360                       {
361                         tempv = (int)(rand_no()*MAXVIRUS); // select a virus randomly from
    surviving viral pool
362                       }
363                     while (virus[tempv].available==false);
364
```

```
                    //printf("cell_count=%d ",cell_count);
                    cell[cell_count].infect(tempv,inf); // infect the cell with the selected virus
                    virus[tempv].available = false; // set infecting virus to false so that it is not available for further infections
                    //printf("%d \n",tempv);
                  }
                //printf("  ");
                cell_count++; // increment cell count
                }
              }
            //printf("\n");

            /*
             printf("cell multiple infection status \n");
            for(int i=0;i<MAXCELL;i++)
            {
              printf("%d ",cell[i].multinf);
            }
            printf("\n");
            */


             // set all viruses as not available, once infection process is complete
            for (int i=0;i<MAXVIRUS;i++)
              {
                virus[i].available = false;
              }

            /*********** RECOMBINATION & MUTATION ************/
            for (int i=0;i<MAXCELL;i++) // loop over all cells
              {
                //printf("cell=%d ",i);
                cell[i].recombine(); // perform recombination to produce proviral DNA from viral RNA
                cell[i].mutate(); // perform mutation on proviral DNA
              }

            //printf("\n\nproviral sequences \n");
            //display_proviral_sequences(); //after infection & reverse transcription


            /********* PRODUCTION **********/
            //printf("produced ");
            virus_prod_count = 0;
            for (int i=0;i<MAXCELL;i++) // loop over all cells
              {
                for (int prod=0;prod<MAXPRODUCE;prod++) // loop over viral production per cell
                  {
                    do
                      {
                       // select a virus position from viral pool to copy offspring virus
                       // if the viral position is not already occupied
                        tempv = (int)(rand_no()*MAXVIRUS);
                      }
                    while (virus[tempv].available == true);

                     // select two proviral DNA strands existing inside cell randomly to copy into new offspring virus
                    v1 = (int)(rand_no()*cell[i].multinf);
                    v2 = (int)(rand_no()*cell[i].multinf);
                    //printf("cell=%d,%d virus=%d provirus=%d,%d \n",i,prod,tempv,v1,v2);

                     // copy proviral DNA strands to viral RNA strands
                    for (int l=0;l<L;l++)
                      {
                        virus[tempv].DNA[l] = cell[i].provirus[v1][l];
                        virus[tempv].DNA[L+l] = cell[i].provirus[v2][l];
                      }
                    virus[tempv].available = true; // set the virus as available for infection
                    virus_prod_count++; // increment viral production count
                    //printf("%d ",tempv);
                  }
                //printf("  ");
              }
```

```
436
437
438             //printf("\n\n");
439
440             //printf("proviral sequences \n");
441             //display_proviral_sequences();  //at the end of each cycle
442             //printf("\n\n viral sequences \n");
443             //display_viral_sequences();  //at the end of each cycle
444
445              // display information regarding current generation and simulation run
446         cout<<"run="<< run <<",cycle="<< cycle <<endl;
447             //printf("run=%d cycle=%d \n",run,cycle);
448
449           // calculate quasispecies structure of proviral DNA pool
450         cal_quasi_structure();
451
452             /*********** FITNESS SELECTION ********/
453         fittest_virus_fitness = 0.0; // set fittest virus fitness to zero
454             for (int i=0;i<MAXVIRUS;i++) // loop over all viruses
455               {
456                 virus[i].fitness = VIRAL_FITNESS; //fitness(i); calculate virus fitness
457                 //printf("fitness of virus[%d]=%lf \n",i,virus[i].fitness);
458
459                  // if the virus fitness is larger than fittest virus fitness
460                  // set fittest virus fitness value to present virus fitness value
461                 if(fittest_virus_fitness < virus[i].fitness)
462                   {
463                     fittest_virus_fitness = virus[i].fitness;
464                   }
465               }
466
467           vcount=0;
468
469           // fitness selection of viruses
470             for (int i=0;i<MAXVIRUS;i++) // loop over all viruses
471               {
472                 // if random number is greater than viral fitness normalized by the
    fittest virus fitness
473                  // then set present virus as not available for infection
474                 if (rand_no()>(virus[i].fitness/fittest_virus_fitness))
475                   {
476                     //printf("virus %d not available \n",i);
477                     virus[i].available = false;
478                     vcount++;
479                   }
480               }
481
482             cout<<"no of viruses surviving "<<(virus_prod_count-vcount)<<endl;
483             //printf("no of viruses surviving %d \n",(MAXVIRUS-vcount));
484
485         // flush file buffers for immediate data writing to file
486         fflush(fp_result);
487         fflush(fp_quasi);
488           } /***** end of cycle *****/
489
490         //display_viral_sequences();  //at the end of each run
491
492       } /***** end of runs *****/
493
494     // Time of simulation end
495     cout<<"Time of Ending is"<<int(time(0))<<endl;
496     time( &rawtime);
497     timeinfo = localtime ( &rawtime );
498
499     cout<<"Date :"<<asctime(timeinfo);  // date and time at the end of simulation
500     fprintf(fp_result,"\n\n%d\nEND: %s \n",int(time(0)),asctime(timeinfo));
501
502
503     // close all simulation result files
504     fclose(fp_result);
505     fclose(fp_quasi);
506
507     return(0);
508   }//end of main
509   /*********************************************************/
510
511
```

```
512  void display_viral_sequences()
513  {
514    for (int i=0;i<MAXVIRUS;i++)
515      {
516        for (int j=0;j<2*L;j++)
517          {
518            printf("%d\t",virus[i].DNA[j]);
519          }
520        printf("\n");
521      }
522  }
523
524  void display_proviral_sequences()
525  {
526    for (int i=0;i<MAXCELL;i++)
527      {
528        //for (int j=0;j<MAXINF;j++)
529        for (int j=0;j<cell[i].multinf;j++)
530          {
531            printf("cell=%d provirus=%d\n",i,j);
532            for (int l=0;l<L;l++)
533              {
534                printf("%d\t",cell[i].provirus[j][l]);
535              }
536            printf("\n");
537          }
538        printf("\n");
539      }
540  }
541
542  // calculate quasispecies structure
543  void cal_quasi_structure()
544  {
545    int hamming = 0;
546    long int nprovirus = 0;
547    long int count = 0;
548
549    for(int l=0;l<=L;l++)
550      {
551        quasi_structure[l] = 0;
552      }
553
554    for (int i=0;i<MAXCELL;i++) // loop over all cells
555      {
556        for(int j=0;j<cell[i].multinf;j++) // loop over number of provirues in each cell
557          {
558             count++;
559             hamming = 0;
560            for (int k=0;k<L;k++) // loop over proviral DNA length
561              {
562                // calculate number of differences between fittest virus
563                // and present provirus for classification into hamming classes
564              if (cell[i].provirus[j][k] != fittest_virus.DNA[k])
565                  hamming++;
566                  //printf("hamming = %d \n",hamming);
567              }
568               // increment hamming class count by 1 in quasispecies structure
569             quasi_structure[hamming] = quasi_structure[hamming] + 1;
570          }
571      }
572
573      // sum to calculate total number of proviruses in all cells
574    for(int inf=0;inf<MAXINF;inf++)
575      {
576        nprovirus = nprovirus + mult_inf[inf]*(inf+1);
577      }
578
579     //printf("nprovirus = %d count = %d\n",nprovirus,count);
580
581     // store quasispecies structure into file
582    for(int len=0;len<=L;len++)
583      {
584        fprintf(fp_quasi,"%d \t",quasi_structure[len]);
585        //printf("%d \t",quasi_structure[len]);
586      }
587    fprintf(fp_quasi,"\n");
588
```

```
}

// set all viral nucleotides to A for all viruses
void initialize_virus()
{
  for (int i=0;i<MAXVIRUS;i++)
    {
      for (int j=0;j<2*L;j++)
        {
          virus[i].DNA[j] = 0;
        }
    }
}

// create fittest virus
void initialize_fittest_virus()
{
  double rand = 0.0;

  // select each nucleotide position uniformly from A, G, C and T
  for (int k=0;k<L;k++)
    {
        rand = rand_no();

      if (rand < 0.25)
        {
          fittest_virus.DNA[k] = 0;
          fittest_virus.DNA[L+k] = 0;
        }
      else if (rand < 0.50)
        {
          fittest_virus.DNA[k] = 1;
          fittest_virus.DNA[L+k] = 1;
        }
      else if (rand < 0.75)
        {
          fittest_virus.DNA[k] = 2;
          fittest_virus.DNA[L+k] = 2;
        }
      else
        {
          fittest_virus.DNA[k] = 3;
          fittest_virus.DNA[L+k] = 3;
        }
    }

  fittest_virus.fitness = 1.0;
}

// create founder virus
void initialize_founder_virus()
{
    long int hamming = 0;
    int ch_no = 2;
    int rand_pos;

    for(int j=0;j<L;j++) //copying of fittest virus to founder virus
    {
      founder_virus.DNA[j] = fittest_virus.DNA[j];
      founder_virus.DNA[L+j] = fittest_virus.DNA[L+j];
    }

    /* mutating fraction of first virus genome length */
    for (int count=0;count<(L*FOUNDER_MUTATE_FRAC);)
     {
       rand_pos = (int)(rand_no()*L);
       //cout<<rand_pos<<" ";
       if(founder_virus.DNA[rand_pos] != fittest_virus.DNA[rand_pos])
       continue;
       else
        {
          founder_virus.DNA[rand_pos] = (fittest_virus.DNA[rand_pos]+ch_no)%4;
          founder_virus.DNA[L+rand_pos] = (fittest_virus.DNA[L+rand_pos]+ch_no)%4;
          count++;
        }
     }

```

```
666
667         /* founder virus fitness calculation */
668         for(int i=0;i<2*L;i++)
669         {
670             if (founder_virus.DNA[i] != fittest_virus.DNA[i])
671             hamming++;
672         }
673
674         hamming = (long int)(hamming/2.0);
675
676         founder_virus.fitness = 1.0 - (1.0-fMIN)*pow(hamming,N)/(pow(hamming,N) + pow(d50,N));
677
678        /* copy founder_virus to initial viral pool */
679         for (int i=0;i<MAXVIRUS;i++)
680         {
681             for(int j=0;j<2*L;j++)
682             {
683                 virus[i].DNA[j] = founder_virus.DNA[j];
684             }
685
686             virus[i].fitness = founder_virus.fitness;
687             virus[i].available = true;
688         }
689     }
690
691
692     // set all viral and proviral sequences to zero in all cells
693     void initialize_cell()
694     {
695       for (int i=0;i<MAXCELL;i++)
696         {
697           for (int j=0;j<MAXINF;j++)
698             {
699               for (int k=0;k<L;k++)
700                 {
701                   cell[i].V[j].DNA[k] = 0;
702                   cell[i].V[j].DNA[L+k] = 0;
703                   cell[i].provirus[j][k] = 0;
704                 }
705             }
706             cell[i].multinf = 0;
707         }
708     }
709
710     // calculate fitness of virus with index number 'v'
711     double fitness(int v)
712     {
713       long int hamming = 0;
714       double fitness = 0.0;
715
716       for (int i=0;i<2*L;i++)
717         {
718           if (virus[v].DNA[i] != fittest_virus.DNA[i])
719               hamming++;
720         }
721
722       hamming = (long int)(hamming/2.0);
723
724       fitness = 1.0 - (1.0-fMIN)*pow(hamming,N)/(pow(hamming,N) + pow(d50,N));
725
726       //printf("hamming value = %d fitness = %lf \n",hamming,fitness);
727       return(fitness);
728     }
729
730
```

```c
/*
 * project: HIV quasispecies
 * file name: mt19937-64.h
 * date: 31-10-08
 * version: 9.2
 * remarks: changed the name of the function "genrand64_real2(void)" to "rand_no(void)"
 */

#ifndef MT1993764_H_
#define MT1993764_H_

/*
   A C-program for MT19937-64 (2004/9/29 version).
   Coded by Takuji Nishimura and Makoto Matsumoto.

   This is a 64-bit version of Mersenne Twister pseudorandom number
   generator.

   Before using, initialize the state by using init_genrand64(seed)
   or init_by_array64(init_key, key_length).

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   References:
   T. Nishimura, ``Tables of 64-bit Mersenne Twisters''
     ACM Transactions on Modeling and
     Computer Simulation 10. (2000) 348--357.
   M. Matsumoto and T. Nishimura,
     ``Mersenne Twister: a 623-dimensionally equidistributed
       uniform pseudorandom number generator''
     ACM Transactions on Modeling and
     Computer Simulation 8. (Jan. 1998) 3--30.

   Any feedback is very welcome.
   http://www.math.hiroshima-u.ac.jp/~m-mat/MT/emt.html
   email: m-mat @ math.sci.hiroshima-u.ac.jp (remove spaces)
*/


#include <stdio.h>

#define NN 312
#define MM 156
#define MATRIX_A 0xB5026F5AA96619E9ULL
#define UM 0xFFFFFFFF80000000ULL /* Most significant 33 bits */
#define LM 0x7FFFFFFFULL /* Least significant 31 bits */


/* The array for the state vector */
```

```c
static unsigned long long mt[NN];
/* mti==NN+1 means mt[NN] is not initialized */
static int mti=NN+1;

/* initializes mt[NN] with a seed */
void init_genrand64(unsigned long long seed)
{
    mt[0] = seed;
    for (mti=1; mti<NN; mti++)
        mt[mti] =  (6364136223846793005ULL * (mt[mti-1] ^ (mt[mti-1] >> 62)) + mti);
}

/* initialize by an array with array-length */
/* init_key is the array for initializing keys */
/* key_length is its length */
//unsigned long long init_key[], key_length;
void init_by_array64(unsigned long long init_key[], unsigned long long key_length)
{
    unsigned long long i, j, k;
    init_genrand64(19650218ULL);
    i=1; j=0;
    k = (NN>key_length ? NN : key_length);
    for (; k; k--) {
        mt[i] = (mt[i] ^ ((mt[i-1] ^ (mt[i-1] >> 62)) * 3935559000370003845ULL))
          + init_key[j] + j; /* non linear */
        i++; j++;
        if (i>=NN) { mt[0] = mt[NN-1]; i=1; }
        if (j>=key_length) j=0;
    }
    for (k=NN-1; k; k--) {
        mt[i] = (mt[i] ^ ((mt[i-1] ^ (mt[i-1] >> 62)) * 2862933555777941757ULL))
          - i; /* non linear */
        i++;
        if (i>=NN) { mt[0] = mt[NN-1]; i=1; }
    }

    mt[0] = 1ULL << 63; /* MSB is 1; assuring non-zero initial array */
}

/* generates a random number on [0, 2^64-1]-interval */
unsigned long long genrand64_int64(void)
{
    int i;
    unsigned long long x;
    static unsigned long long mag01[2]={0ULL, MATRIX_A};

    if (mti >= NN) { /* generate NN words at one time */

        /* if init_genrand64() has not been called, */
        /* a default initial seed is used     */
        if (mti == NN+1)
            init_genrand64(5489ULL);

        for (i=0;i<NN-MM;i++) {
            x = (mt[i]&UM)|(mt[i+1]&LM);
            mt[i] = mt[i+MM] ^ (x>>1) ^ mag01[(int)(x&1ULL)];
        }
        for (;i<NN-1;i++) {
            x = (mt[i]&UM)|(mt[i+1]&LM);
            mt[i] = mt[i+(MM-NN)] ^ (x>>1) ^ mag01[(int)(x&1ULL)];
        }
        x = (mt[NN-1]&UM)|(mt[0]&LM);
        mt[NN-1] = mt[MM-1] ^ (x>>1) ^ mag01[(int)(x&1ULL)];

        mti = 0;
    }

    x = mt[mti++];

    x ^= (x >> 29) & 0x5555555555555555ULL;
    x ^= (x << 17) & 0x71D67FFFEDA60000ULL;
    x ^= (x << 37) & 0xFFF7EEE000000000ULL;
    x ^= (x >> 43);

    return x;
}

```

```
155  /* generates a random number on [0, 2^63-1]-interval */
156  long long genrand64_int63(void)
157  {
158      return (long long)(genrand64_int64() >> 1);
159  }
160
161  /* generates a random number on [0,1]-real-interval */
162  double genrand64_real1(void)
163  {
164      return (genrand64_int64() >> 11) * (1.0/9007199254740991.0);
165  }
166
167  /* generates a random number on [0,1)-real-interval */
168  //double genrand64_real2(void) ( ********* modification to function name ************)
169  double rand_no(void)
170  {
171      return (genrand64_int64() >> 11) * (1.0/9007199254740992.0);
172  }
173
174  /* generates a random number on (0,1)-real-interval */
175  double genrand64_real3(void)
176  {
177      return ((genrand64_int64() >> 12) + 0.5) * (1.0/4503599627370496.0);
178  }
179
180  /*
181  int main(void)
182  {
183      int i;
184      unsigned long long init[4]={0x12345ULL, 0x23456ULL, 0x34567ULL, 0x45678ULL}, length=4;
185      init_by_array64(init, length);
186      printf("1000 outputs of genrand64_int64()\n");
187      for (i=0; i<1000; i++) {
188        printf("%20llu ", genrand64_int64());
189        if (i%5==4) printf("\n");
190      }
191      printf("\n1000 outputs of genrand64_real2()\n");
192      for (i=0; i<1000; i++) {
193        printf("%10.8f ", genrand64_real2());
194        if (i%5==4) printf("\n");
195      }
196      return 0;
197  }
198
199  */
200
201  #endif /*MT1993764_H_*/
202
```