```
#include <iostream>
#include <fstream>
#include <cmath>
#include <valarray>
#include <string>
#include <sstream>
#include "bGrid.h"
#include <iomanip>
#include <string.h>
using namespace std;
const unsigned int DEBUG_COUNT=2;
/* Templates */
template< typename T, std::size_t N > inline
std::size_t size( T(&)[N] ) { return N ; }
 Constructors and Destructors
bGrid::bGrid(): isInRes_(0), residueCount_(0), res_(0.5), fit_(2), thk_(0) {
  for(unsigned int i = 0; i < size(min_); ++i) {</pre>
   min_[i] = 1000.0;
    max_[i] = -1000.0;
 }
bGrid::~bGrid() {}
  Read in Protein from File
bool bGrid::readPoints(char *file) {
  saveFilename(file); // save the filename (no extension)
  protPnt_ = _readPoints(file,prt_,1); // save the protein points
  return protPnt_;
bool bGrid::readPoints(char *file, char *file2) {
  saveFilename(file); // save the filename (no extension)
  protPnt_ = _readPoints(file,prt_,1); // save the protein points
  centPnt_ = _readPoints(file2,cnt_,0); // save the centroid points
  return (protPnt_ & centPnt_);
}
bool bGrid::_readPoints(char *file, std::valarray<float> &pnts, bool save) {
  // open the file
  ifstream ip;
  ip.open(file, ifstream::in);
  if(!ip) {
   return 0;
  printf("File: %s\n",file);
 // read in the file... not the most efficient, bleh
  // -- we read in w/o doing anything to get a cound of the residues
  vector<string> data;
  string bffr;
  for(int i = 0; getline(ip,bffr); ++i) {
    data.push_back(bffr);
  ip.close();
  // save how many points we have
  if(save) {
    residueCount_ = data.size();
```

```
}
  // resize the array and prepare the stream
  // -- this is an unfortunate side effect of using the valarray
  // => it clears everything upon resize =/
  pnts.resize(data.size()*3);
  // loop through each line and save the coordinates
  for(unsigned int i = 0; i < data.size(); ++i) {
    istringstream ss(data[i]);
   float flt;
   // loop through each value on the line
    for(int j = 0; ss >> flt; ++j) {
     pnts[(i*3) + j] = flt;
 }
  return 1;
/* Find min and max */
void bGrid::findMinMax() {
 // Loop over each axis and calculate the min and max values
  int numPnts = prt_.size()/3;
  for(int i = 0; i < 3; ++i) {
    min_[i] = (int)valarray<float>(prt_[slice(i,numPnts,3)]).min();
    max_[i] = (int)valarray<float>(prt_[slice(i,numPnts,3)]).max();
    cout << min_[i] << " : " << max_[i] << endl;
}
  Recenter protein completely within positive boundaries
bool bGrid::repositionPoints() {
  if(protPnt_) {
    // find max and min...and a few other things
    int numPnts = prt_.size()/3;
    for(int i=0; i<3; ++i) {
     // 1) find the min and max
     // 2) adjust them to account for desired grid spacing
     // -- i.e. go from protein point min and max to grid point
     min_[i] = (int)valarray<float>(prt_[slice(i,numPnts,3)]).min() - (fit_ + thk_);
     max_[i] = (int)valarray<float>(prt_[slice(i,numPnts,3)]).max() + (fit_ + thk_);
      cout << min_[i] << " : " << max_[i] << endl;
     // 3) save the correction amount -- still in angstroms
      correction_[i] = min_[i];
     // 4) reposition the min and max to positive numbers
     max_[i] -= min_[i];
     min_[i] = 0;
     cout << min_[i] << " : " << max_[i] << endl;
    // adjust the protein points
   this->protPnt_ = this->_repositionPoints(this->prt_);
  if(centPnt_) {
   this->centPnt_ = this->_repositionPoints(this->cnt_);
  return (this->protPnt_ & this->centPnt_);
```

```
bool bGrid::_repositionPoints(std::valarray<float> &pnts) {
  int numPnts = pnts.size() / 3; // get number of points
  for(int i=0; i<3; ++i) {
    slice s = slice(i,numPnts,3); // create a slice -- single axis
   pnts[s] = valarray<float>(pnts[s]) - correction_[i]; // adjust protein
   cout << i << ": " << correction_[i] << ", " << pnts[i] << endl;
 }
 return 1;
/* Initialize Grid */
void bGrid::initializeGrid() {
  // Check and see if we need extra space, i.e. the z-axis is too big
  // -- if so, double the grid size
  int addGridSpace = this->checkRes();
  cout << addGridSpace << endl;</pre>
  // adjust everything to resolution space
  this->setToResSpace();
  for(int i=0; i<3; ++i) {
   cout << min_[i] << " : " << max_[i] << endl;
 // initialize grid
  // -- the min should be 0 on all accounts
  // -- add 1 since max is the last index, not the size
  int grdDimensions = addGridSpace * (max_[0]+1) * (max_[1] + 1);
  grd_.resize(grdDimensions, 0x0);
  tmp_resize(grdDimensions, 0x0);
  cout << grdDimensions << endl;
  return;
}
/* Check required resolution */
int bGrid::checkRes() {
  numRes_{=} = 1;
  int resCheck = (max_[2] / res_);
  while(resCheck > 63) {
    resCheck -= 63 - ((fit_ + thk_) / res_);
    highRes_ = true;
    ++numRes_;
 }
  return numRes_;
/* set everything to resolution space */
bool bGrid::setToResSpace() {
  if(isInRes_) {
   return 0;
  prt_ /= res_; // protein points
  cnt_ /= res_;
  fit_ /= res_; // paramters
  thk_/= res_;
  for(int i=0; i<3; ++i) { // min and max
    min_[i] /= res_;
   max_[i] /= res_;
 }
```

```
isInRes_ = true;
  return 1;
}
/* set everything to normal space */
bool bGrid::setToNormalSpace() {
  if(!isInRes_) {
   return 0;
 }
  prt_ *= res_; // protein points
 cnt_ *= res_;
  fit_ *= res_; // parameters
  thk_ *= res_;
  for(int i=0; i<3; ++i) { // min and max
   min_[i] *= res_;
   max_[i] *= res_;
 }
 isInRes = false;
  return 1;
}
 Initialize Exclusion and Inclusion Matrices (or grids...yes, I know)
bool bGrid::initializeStamps() {
  int radius = fit_; // exclusion
 haveExc_ = this->_initializeStamp(radius,exc_,exsl_);
  radius = thk_ + fit_; // inclusion
  haveInc_ = this->_initializeStamp(radius,inc_,insl_);
  printStamp();
  return 1;
}
 Initialize Stamp
bool bGrid::_initializeStamp(int radius, std::valarray<unsigned long> &stamp, std::vector< valarray<size_t> > &stampSlice) {
 // radius from calling function: int radius = (fit_ / res_);
  int diamet = 2 * radius;
  // initialize bins
  int binCnt = 0;
  for(int i = 0; i <= (radius); ++i) {
   binCnt += i;
 }
  // resize! (and fill in with ones)
  stamp.resize(binCnt, 0);//(pow(2,diamet) - 1));
  stampSlice.resize(binCnt);
  // loop through each outer layer of the grid
  int index = 0;
  for(int k = 0; k < (radius); ++k) {
   // loop through each unique position on the outer layer
   for(int i = k; i < (radius); ++i) {
     // initialize the valarray
      stampSlice[index].resize(8);
```

```
int slc = 0;
     // calculate each of the eight points
     // note: these are mapped to the grid, not to the small box of the
     // exclusion. MEANING: we only need to add the x dimension (usually i
     // in our implementation) to the indirect arrays to get the
     // appropriate index on the grid.
     int n = max_[0]+1;
     int p = 2*radius - 1;
     stampSlice[index][slc] = i + (k*n);
     stampSlice[index][++slc] = (p-i) + (k*n);
     stampSlice[index][++slc] = (i*n) + k;
     stampSlice[index][++slc] = (i*n) + (p-k);
     stampSlice[index][++slc] = (p-i)*n + k;
     stampSlice[index][++slc] = (p-i)*n + (p-k);
     stampSlice[index][++slc] = (p-k)*n + i;
     stampSlice[index][++slc] = (p-k)*n + (p-i);
      ++index;
   }
 }
 // setup the middle point
 valarray<float> mid( (((float)diamet -1)/2), 3);
 // go through each point in the exclusion matrix...for the unique bins
 index = 0;
  float scaledLength = radius - res ;
  for(int k = 0; k < radius; ++k) {
   for(int i = k; i < radius; ++i) {</pre>
     for(int z = 0; z < radius; ++z) {
       // create temp valarray for the point
       float tmp[] = {(float)k, (float)i, (float)z};
       valarray<float> gpt(tmp,3);
       // check the distance between mid pt and tmp pt
       // -- if w/in range, add symmetrically
       if(pointDistance(mid,gpt) < scaledLength) {</pre>
         stamp[index] |= (unsigned long)pow(2.0,z);
         stamp[index] |= (unsigned long)pow(2.0,(diamet-1-z));
       }
     }
      ++index;
 }
 return 1;
 Calculate Point Distances
float bGrid::pointDistance(valarray<float> &a, valarray<float> &b) {
 valarray<float> c = a-b;
 c *= c;
 return sqrt( c.sum() );
 Stamp Protein Points with [In|Ex]clusion Grids
bool bGrid::stampPoints() {
 //~ for(unsigned int i=0; i< DEBUG_COUNT; ++i) {
   //\sim int index = i*3;
```

}

}

```
//~ cout << "P: ";
   //~ cout << prt_[index] << " : ";
   //~ cout << prt_[++index] << " : ";
   /\!/\sim cout << prt_[++index] << endl;
   //~ cout << "C: ":
   //~ cout << cnt_[index] << " : ";
   //~ cout << cnt_[++index] << " : ";
   //~ cout << cnt_[++index] << endl;
   //~ cout << endl;
  //~ }
  if(haveExc_) {
    int correction = fit ;
    cout << "Stamping Protein" << endl;
    stamped_ = this->_stampPoints(correction,prt_,exc_,exsl_,grd_);
    if(centPnt_) {
     cout << "Stamping Centroids" << endl;</pre>
     stamped_ = this->_stampPoints(correction,cnt_,exc_,exsl_,grd_);
   }
  if(havelnc_) {
    int correction = thk_ + fit_ - 1;
   cout << "Stamping Inclusions" << endl;</pre>
   //~ stamped_ = this->_stampPoints(correction,prt_,inc_,insl_,tmp_);
 // Reconcile the inclusion and exclusion
  grd_= grd_^ tmp_;
  print2dProtein();
  return stamped_;
}
  Stamp Grid Around a Point
bool bGrid::_stampPoints(int correction, std::valarray<float> &pnts,std::valarray<unsigned long> &stamp, std::vector< valarray<
 // Corection: given from calling function
 //~ int correction = (fit_/res_) - 1;
  // displacement and depth of exclusion grid
  int position = 0;
  int depth = 0;
  // Resize the protein to the proper resolution
  // -- we initialized the grid to the resolution
  //~ pnts /= res_;
  // shift point from the center box to the corner
  pnts -= correction;
  // create slices
  slice xs = slice(0,residueCount_,3);
  slice ys = slice(1,residueCount_,3);
  slice zs = slice(2,residueCount_,3);
  int allowableDepth = 63;
  if(highRes_) {
    allowableDepth = 63 - (fit_ + thk_);
```

```
// Loop through protein points
for(unsigned int i=0; i<DEBUG_COUNT;++i) { //pnts.size()/3; ++i) { // DEBUG_COUNT;++i) { //
 // a little ugly due to valarray implementation
  int index = i*3:
  int x = pnts[index];
  int y = pnts[++index];
  int z = pnts[++index];
 cout << "STAMPING:" << endl;
 cout << "\t" << x << endl;
 cout << "\t" << y << endl;
 cout << "\t" << z << endl;
 // save old positions
 int displacement = position;
  int perspective = depth;
 // calculate new positions
  position = ((y)*(max_[0]+1) + (x));
  depth = z;
 // check and adjust for high resolution
  if(z > allowableDepth) {
   adjustPlacement(depth,position);
   cout << "IN EXTRA!!!" << endl;
   cout << "\t" << position << endl;
 // adjust according to previous placement
  displacement = position - displacement;
 perspective = depth - perspective:
 cout << "Position: " << endl;
 cout << "\t" << position << endl;
 cout << "\t" << displacement << endl;
 cout << "Depth: " << endl;
 cout << "\t" << depth << endl;
 cout << "\t" << perspective << endl;
 // Loop through exclusion slice, displace each, and EXCLUDE
 // -- might it be faster to simply multiply rather than test?
 for(unsigned int g=0; g<stamp.size(); ++g) {</pre>
   stampSlice[g] += displacement;
   stamp[g] *= pow(2.0,perspective);
   //~ if(z>allowableDepth) {
     //~ cout << "Grid:" << endl;
     //~ cout << "\t" << stampSlice[g][0] << endl;
     //~ cout << "\t" << stamp[g] << endl;
     //~ cout << "\t" << paper[stampSlice[g][0]];
     //~ cout << endl;
   //~ }
     //~ cout << "\t";
     //~ printSingleHex(cout,stamp[g],63);
     //~ cout << "\t";
     //~ printSingleHex(cout,paper[stampSlice[g][0]],63);
   // EXCLUDE POINTS!!
   paper[stampSlice[g]] = stamp[g] | valarray<unsigned long>(paper[stampSlice[g]]);
```

```
//~ if(z>allowableDepth) {
       //~ cout << "\t";
       //~ printSingleHex(cout,paper[stampSlice[g][0]],63);
     //~ cout << endl;
     //~ }
   } // end loop through exclusion space
   cout << "END" << endl;
   //~ cout << endl;
 } // end loop through points
 // Reset protein to original location
  pnts += correction;
  //~ pnts *= res_;
     cout << "END" << endl;
  // Reset stamp to original location
  stamp /= pow(2.0,depth);
     cout << "END" << endl;
  for(unsigned int i=0; i<stampSlice.size(); ++ i) {</pre>
   stampSlice[i] -= position;
     cout << "END" << endl;
  return 1;
}
bool bGrid::adjustPlacement(int &depth, int &position) {
  int tooBig = 63 - (fit_ + thk_);
  int howBig = 1;
  while(depth > tooBig) {
    depth -= tooBig;
    ++howBig;
    if(howBig == numRes_) {
     tooBig = 63;
 position += (max_[0] + 1) * (max_[1] + 1);
  return 1;
}
  Find Nearby Points
int bGrid::countNearby(std::valarray<float>&a) {
 // adjust coordinates from center to corner
 a -= fit_ - 1;
  // initialize placement coordinates
  int position = (a[1] * max_[0]) + a[0];
  int depth = a[2];
  // check and adjust for high resolution
  int allowableDepth = 63 - (fit_ + thk_);
  if(depth>allowableDepth && highRes_) {
    depth -= allowableDepth;
   position += (max_[0] + 1) * (max_[1] + 1);
  // initialize counting variables
  int nearby = 0;
  std::valarray<unsigned long> overlap(exc_.size());
 //~ cout << "STAMPING:" << endl;
```

```
//~ cout << "\t" << a[0] << endl;
  //~ cout << "\t" << a[1] << endl;
  //~ cout << "\t" << a[2] << endl;
  //~ cout << "Position: " << endl;
 //~ cout << "\t" << position << endl;
  //~ cout << "Depth: " << endl;
  //~ cout << "\t" << a[2] << endl;
  for(unsigned int i=0; i<exsl_.size(); ++i) {</pre>
    exsl_[i] += position;
   exc_[i] *= pow(2.0,depth);
   //~ cout << endl;
   //\sim cout << i << endl;
   //~ cout << "Grid:" << endl;
   //~ cout << "\t" << exc_[i] << endl;
   //~ cout << "\t" << exsl_[i][0] << endl;
   //~ cout << "\t" << grd_[exsl_[i][0]];
   //~ cout << endl;
   //~ cout << "\t":
   //~ printSingleHex(cout,exc_[i],63);
   //~ cout << "\t":
   //~ printSingleHex(cout,grd_[exsl_[i][0]],63);
   //~ cout << "\t":
   //~ overlap[i] = exc_[i] & grd_[exsl_[i][0]];
   //~ printSingleHex(cout,overlap[i],63);
   overlap[i] = exc_[i] & grd_[exsl_[i][0]];
   overlap[i] >>= int(depth - fit_);
   //~ cout << "Hamming: " << overlap[i] << " : " << hamming_[overlap[i]] << endl;
    nearby += hamming_[overlap[i]];
    exc_[i] >>= (int)depth;
   exsl_[i] -= position;
  // return coordinates to corner
  a += fit_ - 1;
  return nearby;
/* Hamming table
  If we can store a lookup table of the hamming function of every 16 bit integer, we can do the following to compute the Hamming w
    static unsigned char wordbits[65536] = { bitcounts of ints between 0 and 65535 };
    static int popcount( unsigned int i )
      return( wordbits[i&0xFFFF] + wordbits[i>>16] );
*/
}
  Setup Hamming Table
void bGrid::setupHammingTable() {
  int a = pow(2,16);
  for(int i =0; i<a; ++i) {
    hamming_[i] = this->calculateHammingWeight(i);
```

```
}
 return;
}
  Calculate Hamming Weight
int bGrid::calculateHammingWeight(int x) {
  int i=0;
  for(i; x; ++i) {
   x &= x-1;
 }
 return i;
}
  Get Hamming Weight for 64 bit numbers
unsigned long bGrid::getHammingWeight(unsigned long x) {
  int weight = 0;
  for(int i=0; i<4 && x > 0; ++i) {
   //~ unsigned long b = 0xFFFF;
   //~ printSingleHex(cout,x,64);
   //~ printSingleHex(cout,b,64);
   weight += hamming_[int(x&0xFFFF)];
   //~ cout << "\tAND: " << (x&0xFFFF) << endl;
   //~ cout << "\tWEIGHT: " << hamming_[int(x&0xFFFF)] << endl;
   x >>= 16;
 }
  return weight;
}
  Print Grid
  ** Removed print 3d exclusion -- wasn't working correctly. Rewrite!
void bGrid::print3dStamp(ostream &out,int radius, valarray<unsigned long> &stamp,std::vector< valarray<size_t> > &stampSlice
  return;
}
void bGrid::print3dGrid(ostream &out, valarray<unsigned long> &grid, int x) { this->print3dGrid(out,grid,x,x); }
void bGrid::print3dGrid(ostream &out, valarray<unsigned long> &grid, int x, int d) {
 for(unsigned int k=0; k < (grid.size()/x); ++k) {</pre>
   for(int i=0; i < x; ++i) {
     for(int z = 0; z < d; ++z) {
       unsigned long a = pow(2.0,z);
       switch(a & grid[k*x + i]) {
         case 0: out << " " << 0;
           break;
         default: out << " " << 1;
           break;
       }
     }
     cout << endl;
   }
   cout << endl;
 }
void bGrid::printSingleHex(ostream &out,unsigned long &one,int z) {
  for(int i=0; i<z; ++i) {
    unsigned long a = pow(2.0,i);
    if(a & one) {
```

```
out << 1;
   }
   else {
     out << 0;
   }
 }
 out << endl;
 Create PyMol Python Script
void bGrid::printPyMol() {
 this->setToNormalSpace();
 // creat the filename
 char *spdb = (char*)malloc(sizeof(char)*64);
 char *sprt = (char*)malloc(sizeof(char)*64);
 strcpy(spdb,prtFile_);
 strcpy(sprt,prtFile_);
  strcat(spdb,".pdb");
  strcat(sprt,"_grid.py");
 strcat(prtFile_,"_grid.py");
 cout << "Writing to: " << sprt << endl;
 // open the file and check
  FILE *op = fopen(sprt, "w");
  if(!op) {
   cout << sprt << " did not open!\n" << endl;
 }
 // Reset to original coordinates
 for(int i=0; i<3; ++i) {
   slice s = slice(i,prt_.size()/3,3);
   prt_[s] = valarray<float>(prt_[s]) + correction_[i];
   s = slice(i,cnt_.size()/3,3);
   cnt_[s] = valarray<float>(cnt_[s]) + correction_[i];
 }
 // Write header
 fprintf(op,"from pymol import cmd\n");
  fprintf(op,"from pymol.cgo import *\n");
 fprintf(op,"\n");
 fprintf(op,"cmd.hide(\"everything\")\n");
  fprintf(op,"cmd.show(\"cartoon\")\n");
  fprintf(op,"cmd.set('transparency','1.0')\n");
  fprintf(op,"cmd.bg_color('white')\n");
 //~ fprintf(op,"FREEMOL=/usr/share/pymol/freemol-trunk/freemol\n");
 //~ fprintf(op,"export FREEMOL");
 fprintf(op,"\n");
 // Write pdb
 //\sim fprintf(op,"cmd.load(\"%s\")\n",spdb);
 //~ fprintf(op,"cmd.color(\"firebrick\")\n");
 //~ fprintf(op,"cmd.show(\"stick\")\n");
 // Write origin
 float mi[3];
 float ma[3];
  for(int i=0; i<3; ++i) {
    mi[i] = min_[i] + correction_[i];
   ma[i] = max_[i] + correction_[i];
 }
```

```
fprintf(op,"orgn = [\n");
fprintf(op,"\tCOLOR, 1.0, 1.0, 1.0,\n");
fprintf(op,"\tSPHERE, %.2f, %.2f, %.2f, %.2f,\n",mi[0],mi[1],mi[2], 1.0);
fprintf(op,"\tSPHERE, %.2f, %.2f, %.2f, %.2f,\n",mi[0],mi[1],ma[2], 1.0);
fprintf(op, "\tSPHERE, %.2f, %.2f, %.2f, \n", mi[0], ma[1], mi[2], 0.5);
fprintf(op,"\tSPHERE, %.2f, %.2f, %.2f, %.2f,\n",ma[0],mi[1],mi[2], 0.5);
fprintf(op, "\tSPHERE, %.2f, %.2f, %.2f, %.2f, \n", ma[0], ma[1], mi[2], 0.5);
fprintf(op,"\t]\n");
fprintf(op,"cmd.load_cgo(orgn,'orgn')\n");
// Write protein points
fprintf(op,"protein = [\n");
fprintf(op,"\tCOLOR, 0.0, 1.0, 0.0,\n");
for(unsigned int i=0; i<DEBUG_COUNT;++i) { //prt_.size()/3; ++i) {</pre>
  int index = i*3;
  fprintf(op,"\tSPHERE, %.2f, %.2f, %.2f, %.2f,\n",
    prt_[index], prt_[index+1], prt_[index+2], 0.2);
fprintf(op,"\t]\n");
fprintf(op,"cmd.load_cgo(protein,'protein')\n");
// Write centroid points
if(centPnt_) {
  fprintf(op,"centroid = [\n");
  fprintf(op,"\tCOLOR, 1.0, 0.5, 0.5,\n");
  for(unsigned int i=0; i<DEBUG_COUNT;++i) { //cnt_.size()/3; ++i) {</pre>
    int index = i*3;
    fprintf(op,"\tSPHERE, %.2f, %.2f, %.2f, %.2f,\n",
      cnt_[index], cnt_[index+1], cnt_[index+2], 0.1);
 fprintf(op,"\t]\n");
 fprintf(op,"cmd.load_cgo(centroid,'centroid')\n");
// Adjust grid points from resolution to angstroms
// Wait...bad idea...good chance we've already
// filled up all available bits...
// -- better do this individually =/
// grd_ *= res_;
// Write grid points
int allowableDepth = 63;
if(highRes_) {
  allowableDepth -= (fit_ + thk_)/res_;
fprintf(op,"grid = [\n");
fprintf(op,"\tCOLOR, 0.0, 0.0, 1.0,\n");
for(int k=0; k<=(max_[1]/res_); ++k) {
  for(int i=0; i<=(max_[0]/res_); ++i) {
    for(int z=0; z<=(max_[2]/res_); ++z) {
      int index = k*((max_[0]/res_)+1) + i;
      // adjust index and depth for highRes
      int depth = z;
      if(z > allowableDepth) {
        adjustPlacement(depth,index);
        //~ index += (max_[0]) * (max_[1]) / res_;
        //~ depth -= allowableDepth;
        //~ cout << "Printing other" << endl;
        //~ cout << "\t" << index << endl;
      if((grd_[index] & (unsigned long)pow(2.0,depth))) {
        fprintf(op,"\tSPHERE, %f, %f, %f, %.2f,\n",
          //\sim ((i*res_)+correction_[0]), ((k*res_)+correction_[1]), ((z*res_)+correction_[2]), 0.1);
```

```
((i*res_)+correction_[0]), ((k*res_)+correction_[1]), ((z*res_)+correction_[2]), 0.1);
       }
     }
   }
  }
  fprintf(op,"\t]\n");
  fprintf(op,"cmd.load_cgo(grid,'grid')\n\n");
  fprintf(op,"cmd.center('grid')\n");
  // Movie -- Rotate!
  // -- set up the frames
  fprintf(op,"cmd.mclear()\n");
  fprintf(op,"cmd.mset(\"1 x360\")\n");
  // -- simple 2-axis rotation
  fprintf(op, "for i in range(1,120):\n");
  fprintf(op,"\tcmd.mdo(i,'turn x,0.5; turn y,0.5; turn z,0.5')\n");
  // -- rotate and zoom in
  fprintf(op, "for i in range(120,240):\n");
  fprintf(op,"\tcmd.mdo(i,'turn x,0.5; turn y,0.5; turn z,0.5; move z,0.9')\n");
  // -- rotate and zoom out
  fprintf(op,"for i in range(240,360):\n");
  fprintf(op,"\tcmd.mdo(i,'turn x,0.5; turn y,0.5; turn z,0.5; move z,-0.9')\n");
  //~ fprintf(op,"cmd.mplay()\n");
  // Attempts to speed things up
  //fprintf(op,"cmd.set('orthoscopic','on')\n");
  //fprintf(op,"cmd.set('depth_cue','1')\n");
  //fprintf(op, "select cgo01, (all) and not ( (all) within 8 of cgo01) )");
  //fprintf(op, "hide everything, cgo01");
  //fprintf(op,"cmd.ray()");
  // Close the file handle
  fclose(op);
  // Run the command
  // ('cuz windows sucks...)
  //system("pymolwin -r \"D:\\My Dropbox\\grid\\1AWQ_alone_grid.py\\"");
}
void bGrid::saveFilename(char* file) {
  prtFile_ = (char*)malloc(sizeof(char)*64);
  strncpy(prtFile_,file,strrchr(file,'.')-file);
  return;
}
void bGrid::printProtein() {
  // Easy loop through grid -- SAVE
  for(int k=0; k<=max_[1]; ++k) {</pre>
    for(int i=0; i<=max_[0]; ++i) {</pre>
      int index = k*max_[0] + i;
      for(int z=0; z<=max_[2]; ++z) {
        bool t = grd_[index] & (unsigned long)pow(2.0,z);
        cout << t << " ";
      }
      cout << endl;
    }
    cout << endl;
void bGrid::printStamp() {
```

```
int index = 0;
  for(int i=0; i<fit_; ++i) {</pre>
    for(int j=0; j<i; ++j) {
      cout << " ";
    for(int j=i; j<fit_; ++j) {</pre>
      if(exc\_[index] > 0) {
        //~ cout << exc_[i] << endl;
        cout << "1 ";
      else {
        cout << "0 ";
      ++index;
    }
    cout << endl;
  cout << exc_.size() << " : " << index << endl;
}
void bGrid::print2dProtein() {
  //\sim for(int k=0; k<(max_[1]+1); ++k) {
    //~ printf("%2d: ",k);
    //\sim for(int i=0; i<(max_[0]+1); ++i) {
      //\sim int index = k^*(max_[0]) + i;
      //~ printf("%4d ",i);
    //~ }
    //~ printf("\t:%d\n",max_[0]);
  //~ }
  //~ printf("::%d\n\n",max_[1]);
  /\!\!/\sim for(int k=0; k<(max_[1]+1); ++k)  {
    //~ printf("%2d: ",k);
    //\sim for(int i=0; i<(max_[0]+1); ++i) {
      //\sim int index = k*(max_[0]+1) + i;
      //~ printf("%4d ",index);
    //~ }
   //~ printf("\t:%d\n",max_[0]);
  //~ }
  //~ printf("::%d\n\n",max_[1]);
  for(int k=0; k<(max_[1]+1); ++k) {
    printf("%2d: ",k);
    for(int i=0; i<(max_[0]+1); ++i) {
      int index = k*(max_[0]+1) + i;
      if(grd_[index] > 1) {
        printf("%d ",1);
      }
      else {
        printf("%d ",0);
   }
    printf("\t:%d\n",max_[0]);
  printf("::%d\n\n",max_[1]);
  printf("Correction");
  for(int i=0; i<3; ++i) {
    printf(": %f",correction_[i]);
 printf("\n\n");
```