

```
#include <set>
#include <list>
#include <vector>
#include <fstream>
#include <iostream>
#include <string>
```

```
// Written by Eric Wasiolek Copyright 2008
```

```
using namespace std;
```

```
class neuron{
    private:
        string name;
        string lineage;
        string maintype;
        string subtype1;
        string subtype2;
        set<string> genes;

    public:
        neuron()
        {}
        void setName(string x)
        {
            name = x;
        }
}
```

```
void setLineage(string x)
{
    lineage = x;
}
void setMainType(string x)
{
    maintype = x;
}
void setSubType1(string x)
{
    subtype1 = x;
}
void setSubType2(string x)
{
    subtype2 = x;
}
void addGene(string gene)
{
    genes.insert(gene);
}
string getName()
{
    return name;
}
string getLineage()
{
    return lineage;
}
string getMainType()
```

```

    {
        return maintype;
    }
    string getSubType1()
    {
        return subtype1;
    }
    string getSubType2()
    {
        return subtype2;
    }
    set<string> getGenes()
    {
        return genes;
    }
};

```

// SET FUNCTION IMPLEMENTATIONS

```

template <typename T>
bool operator== (const set<T>& lhs, const set<T>& rhs)
{
    typename set<T>::const_iterator myself = lhs.begin(), other = rhs.begin();

    // return false if the sets do not have the same size
    if (lhs.size() == rhs.size())
    {
        // compare until encounter end of the sets or

```

```

        // find two elements that are not equal
        while (myself != lhs.end() && *myself++ == *other++);

        // if we left the loop before reaching the end
        // of the sets, they are not equal
        if (myself != lhs.end())
            return false;
        else
            return true;
    }
else
    return false;
}

template <typename T>
set<T> operator+ (const set<T>& lhs, const set<T>& rhs)
{
    // construct union
    set<T> setUnion;

    // iterators that traverse the sets
    typename set<T>::const_iterator lhsIter = lhs.begin(), rhsIter = rhs.begin();

    // move forward as long as we have not reached the end of
    // either set
    while (lhsIter != lhs.end() && rhsIter != rhs.end())
        if (*lhsIter < *rhsIter)
            // *lhsIter belongs to the union. insert and
            // move iterator forward
            setUnion.insert(*lhsIter++);
}

```

```

else if (*rhsIter < *lhsIter)
    // *rhsIter belongs to the union. insert and
    // move iterator forward
    setUnion.insert(*rhsIter++);
else
{
    // the two values are equal. insert just one and
    // move both iterators forward
    setUnion.insert(*lhsIter++);
    rhsIter++;
}

// flush any remaining items
if (lhsIter != lhs.end())
    while (lhsIter != lhs.end())
        setUnion.insert(*lhsIter++);
else if (rhsIter != rhs.end())
    while (rhsIter != rhs.end())
        setUnion.insert(*rhsIter++);

return setUnion;
}

```

```

template <typename T>
set<T> operator* (const set<T>& lhs, const set<T>& rhs)
{
    // construct intersection
    set<T> setIntersection;

    // iterators that traverse the sets

```

```

typename set<T>::const_iterator lhsIter = lhs.begin(), rhsIter = rhs.begin();

// move forward as long as we have not reached the end of
// either set
while (lhsIter != lhs.end() && rhsIter != rhs.end())
    if (*lhsIter < *rhsIter)
        // *lhsIter is in lhs and not in rhs. move iterator
        // forward
        lhsIter++;
    else if (*rhsIter < *lhsIter)
        // *rhsIter is in rhs and not in lhs. move iterator
        // forward
        rhsIter++;
    else
    {
        // the same value is in both sets. insert one value
        // and move the iterators forward
        setIntersection.insert(*lhsIter);
        lhsIter++;
        rhsIter++;
    }

    return setIntersection;
}

```

```

template <typename T>
set<T> operator- (const set<T>& lhs, const set<T>& rhs)
{
    // construct difference

```

```

set<T> setDifference;

// iterators that traverse the sets
typename set<T>::const_iterator lhsIter = lhs.begin(), rhsIter = rhs.begin();

// move forward as long as we have not reached the end of
// either set
while (lhsIter != lhs.end() && rhsIter != rhs.end())
    if (*lhsIter < *rhsIter)
        // *lhsIter belongs to lhs but not to rhs. put it in
        // the difference
        setDifference.insert(*lhsIter++);
    else if (*rhsIter < *lhsIter)
        // *rhsIter is in the rhs but not in the lhs. pass
        // over it
        rhsIter++;
    else
    {
        // the same value is in both sets. move the
        // iterators forward
        lhsIter++;
        rhsIter++;
    }

// flush any remaining items from lhs
if (lhsIter != lhs.end())
    while (lhsIter != lhs.end())
        setDifference.insert(*lhsIter++);

return setDifference;

```

```
}
```

```
set<string> findGeneSet(string neuronname, neuron neuron[], int numneurons)
```

```
{
```

```
    for(int i=0; i < numneurons; ++i)
```

```
        if(neuronname == neuron[i].getName())
```

```
            return neuron[i].getGenes();
```

```
}
```

```
bool find(string genename, set<string>& geneset)
```

```
{
```

```
    set<string>::iterator geneiter, endgenes;
```

```
    geneiter = geneset.begin();
```

```
    endgenes = geneset.end();
```

```
    int counter = 0;
```

```
    if(geneset.empty())
```

```
        return false;
```

```
    while(geneiter != geneset.end())
```

```
    {
```

```
        ++counter;
```

```
        if(genename == *geneiter)
```

```
            return true;
```

```
        else if(counter == geneset.size())
```

```
            return false;
```

```
        ++geneiter;
```

```
    }
```

```
}
```

```
template<typename Iterator>
```



```
void writeContainer(Iterator first, Iterator last, const string& seperator = " ")
```

```
{  
    Iterator iter = first;  
    while(iter != last)  
    {  
        cout << *iter << seperator;  
        iter++;  
    }  
}
```

```
void writeSet(set<string> s)
```

```
{  
    set<string>::iterator siter = s.begin(), esiter = s.end();  
  
    while(siter != esiter)  
    {  
        cout << *siter << " ";  
        siter++;  
    }  
    cout << endl;  
}
```

```
void writeBVector(vector<bool>& bvect)
```

```
{  
    int i = 0;  
    for(i = 0; i < bvect.size(); ++i)  
        cout << bvect[i];  
}
```

```

int main(int argc, char *argv[])
{
    // cout << "Entered main" << endl;
    ifstream neurons;
    string inputneuron, outputneuron;

    if(argc != 2)
    {
        cout << "Usage: bitgenes graphname " << endl;
        return 0;
    }
    else
    {
        neurons.open(argv[1]);
        if(!neurons)
        {
            cout << "Could not open graph file!" << endl;
            return 0;
        }
        else
            cout << "Opened graph file " << argv[1] << " successfully." << endl;
    }

    int numneurons = 0;
    string gene;

```

```

vector<neuron> vneurons(numneurons);

neurons >> numneurons;

string temp;

neuron n[numneurons];

int i = 0;

for(i = 0; i < numneurons; ++i)
{
    neurons >> temp;
    n[i].setName(temp);
    neurons >> temp;
    n[i].setLineage(temp);
    neurons >> temp;
    n[i].setMainType(temp);
    neurons >> temp;
    n[i].setSubType1(temp);
    neurons >> temp;
    n[i].setSubType2(temp);
    while(1)
    {
        neurons >> gene;
        if(gene == "end")
            break;
        n[i].addGene(gene);
    }
}

```

```

vector<set<string> > vs;

```

```

list<string>::iterator liter;
set<string>::iterator fsiter;
set<string>::iterator esiter;

// Create a vector of all of the gene sets
for(i = 0; i < numneurons; ++i)
{
    vs.push_back(n[i].getGenes());
}

// Create a union set with all of the unique genes in it
set<string> us;

us = vs[0];

for(i=0; i < vs.size(); ++i)
    us = us + vs[i];

cout << "There are " << us.size() << " neuron specific genes in C. Elegans" << endl;
cout << "And here are the genes: " << endl;
fsiter = us.begin();
esiter = us.end();
writeSet(us);

// Create bit strings showing which neurons have which genes
vector< vector<bool> > bvect(numneurons);

for(i = 0; i < numneurons; ++i)
{

```

```

// cout << "Diag: Bitstring for neuron " << i << endl;
while(fsiter != esiter)
{
    // The essence of this algorithm:
    if(find(*fsiter, vs[i])) // if you find the gene in the geneset iterated by fsiter in
the set of genes per neuron
        // vector of sets vs, set it to 1 else 0
        bvect[i].push_back(true);
    else
        bvect[i].push_back(false);
    ++fsiter;
}
fsiter = us.begin();
}

// Print out bit vectors
for(i = 0; i < numneurons; ++i)
{
    cout << "Neuron: " << n[i].getName() << " Bitstring: ";
    writeBVector(bvect[i]);
    cout << endl;
}

return 0;
}

```