

Biostatistics 615/815 Lecture 19:

StatGen Library

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Outline

- StatGen Library Overview
- Building with the Library
- Useful Classes
 - InputFile
 - String
 - StringArray
 - Parameters
- Documentation
- Debugging

What is the StatGen Library?

- Written by various members of the department
- Contains:
 - C++ library classes
 - General Operation Classes including:
 - File input/output, string processing, parameter parsing
 - Statistical genetics specialized classes
 - Useful Statistical Genetics programs
- Compiles on Linux

Where is the StatGen Library?

- Stored in a git repository
 - <https://github.com/statgen/statgen>
 - Can browse code & history
- Benefits of a repository:
 - Code consolidated in one location so it is easy to find
 - Bugs can be fixed in one place and picked up by everyone
 - Enable reuse rather than always reinventing
 - Version Tracking
 - If it stops working, you can see what changed & revert

StatGen Library on GitHub

- If your machine has git (preferable) do the following:
 - *cd <directory where you want the repository located>*
 - *git clone https://github.com/statgen/statgen.git*
 - Creates a directory called statgen in the directory where you are located.
 - *cd statgen*
- In the future, update to the latest versions (advantage of using git):
 - *git pull*
 - Recompile from the top level (use a make clean first)
- If you do not have git, follow the instructions on the wiki:
 - <http://genome.sph.umich.edu/wiki/File:GithubWithoutGit.pdf>

Compiling StatGen Library

- From the statgen directory, build the repository:
 - For normal (optimized) mode, type:
 - *make*
 - For debug, instead use:
 - *make OPTFLAG="-ggdb -O0"*
 - Note: that is the letter O followed by the number zero.
 - When switching between debug & non-debug, prior to typing the new '*make*' call, type:
 - *make clean*

Building with the Library

- Assumes the library has been compiled and exists at: /home/mktrost/statgen (replace with the correct path to your library).
- Create and move into a directory for your code
 - Example: *mkdir ~/statgenDemo; cd ~/statgenDemo*
- Copy the Makefile.
 - *cp /home/mktrost/statgen/src/Makefile.src Makefile*
 - .o files will go in an obj directory that will be created

Makefile.tool

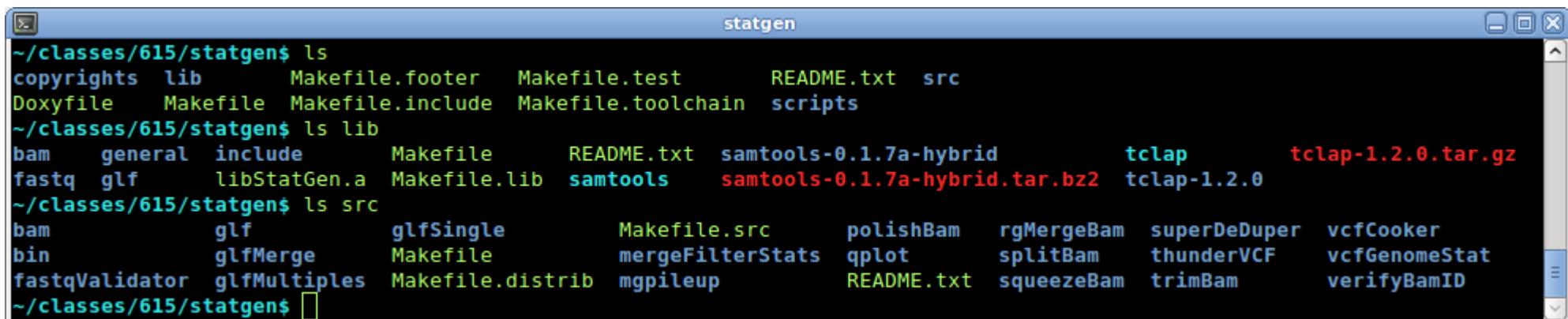
- Create Makefile.tool and set the following values:
 - EXE = yourExecutableName
 - TOOLBASE = File1 File2
 - SRCONLY = Main.cpp
 - HDRONLY = YourTemplate1.h
 - PATH_TO_BASE = /home/mktrost/statgen
 - BIN_DIR = .
 - USER_LIBS = /somePath/lib1.a
 - USER_INCLUDES = -I/somePath/IncDir
 - Example Makefile.tool:

```
EXE = libraryExample
TOOLBASE = LibraryExample
SRCONLY = Main.cpp
PATH_TO_BASE = /home/mktrost/statgen
BIN_DIR = .
```
 - Type *make* or *make OPTFLAG="-ggdb -O0"*

Executable name
Basename of files that have both cpp & h
Files with just .cpp and no .h
Files with just .h and no .cpp
Your statgen location
Where to put the EXE
Any additional libraries
Any additional include directories

StatGen Library Directory Structure

- copyrights – copyrights for various code pieces
- src – premade statgen programs
- lib – library code
 - **general** – general processing classes
 - include – library header files are linked here when compiled
 - specialized classes for processing different file types:
 - bam, fastq, glf
- samtools – samtools software
- tclap – alternate parameter processing library
- libStatGen.a – the compiled library



The screenshot shows a terminal window titled "statgen" displaying the directory structure of the StatGen library. The output of the "ls" command is as follows:

```
~/classes/615/statgen$ ls
copyrights  lib      Makefile.footer  Makefile.test      README.txt  src
Doxyfile    Makefile  Makefile.include  Makefile.toolchain  scripts

~/classes/615/statgen$ ls lib
bam        general  include      Makefile      README.txt  samtools-0.1.7a-hybrid          tclap        tclap-1.2.0.tar.gz
fastq     glf      libStatGen.a  Makefile.lib  samtools   samtools-0.1.7a-hybrid.tar.bz2  tclap-1.2.0

~/classes/615/statgen$ ls src
bam          glf      glfSingle     Makefile.src      polishBam   rgMergeBam   superDeDuper  vcfCooker
bin          glfMerge  Makefile      mergeFilterStats qplot       splitBam    thunderVCF   vcfGenomeStat
fastqValidator glfMultiples  Makefile.distrib  mpileup      README.txt  squeezeBam  trimBam    verifyBamID
~/classes/615/statgen$
```

InputFile/IFILE

- lib/general/InputFile.h(cpp)
- Read/Write different types of files
 - Uncompressed, BGZF, GZIP
- Reads file to determine type unless reading from stdin
- Flag to specify type for reading stdin and writing
 - Default:
 - gzip for files ending in .gz
 - uncompressed for all others
- IFILE is a pointer to the InputFile class
- Similar to the FILE* interface from the C library

IFILE

Functions

IFILE	ifopen (const char *filename, const char *mode, InputFile::ifileCompression compressionMode=InputFile::DEFAULT)	'-' for stdin/stdout	
	Open a file.		
int	fclose (IFILE file)		
	Close the file.		
unsigned int	ifread (IFILE file, void *buffer, unsigned int size)	"r" for read "w" for write	
	Read size bytes from the file into the buffer.		
int	ifgetc (IFILE file)		
	Get a character from the file.	You must preallocate buffer with at least size bytes	
void	ifrewind (IFILE file)		
	Reset to the beginning of the file.		
int	ifeof (IFILE file)		
	Check to see if we have reached the EOF.		
unsigned int	ifwrite (IFILE file, const void *buffer, unsigned int size)		
	Write the specified buffer into the file.		
long int	ftell (IFILE file)		
	Get current position in the file.		
bool	ifseek (IFILE file, long int offset, int origin)		
	Seek to the specified offset from the origin.		
int	fprintf (IFILE output, char *format,...)		
	Write to a file using fprintf format.		
IFILE	operator>> (IFILE stream, std::string &str)		
	Read a line from a file using streaming.		Used for writing or reading from stdin DEFAULT – gz extension, uses gzip, otherwise uncompressed UNCOMPRESSED GZIP – standard compressed format BGZF – specialized compressed format for easy "random" access

For more detailed descriptions:

http://www.sph.umich.edu/csg/mktrost/doxygen/march22_2011/InputFile_8h.html

Additional InputFile Methods

Handle buffering
of reads.
Low level I/O
detail you don't
need to worry
about

```
/// Set the buffer size for reading from files so that bufferSize bytes
/// are read at a time and stored until accessed by another read call.
/// This improves performance over reading the file small bits at a time.
/// Buffering reads disables the tell call for bgzf files.
/// Any previous values in the buffer will be deleted.
/// \param bufferSize number of bytes to read/buffer at a time,
/// default buffer size is 1048576, and turn off read buffering by setting
/// bufferSize = 1;
void bufferReads(unsigned int bufferSize = DEFAULT_BUFFER_SIZE);

/// Disable read buffering.
void disableBuffering();

/// Returns whether or not the file was successfully opened.
/// \return true if the file is open, false if not.
bool isOpen();

/// Get the filename that is currently opened.
/// \return filename associated with this class
const char* getFileName() const;
```

ifprintf format

- Beware of type safety!
- fprintf format (tons of formatting options):
 - <http://www.cplusplus.com/reference/clibrary/cstdio/fprintf/>
- printf example:

```
void testPrintf()
{
    string mystring = "Hello";
    const char* mycharptr = "Bye";
    int myint = 615;
    char mychar = 'Z';
    double mydouble = 1.23456789;
    printf("string %s, int %d, character %c, double %f, char* %s\n",
           mystring.c_str(), myint, mychar, mydouble, mycharptr);

}
```

Write string Hello, int 615, character Z, double 1.234568, char* Bye

- For ifprintf, add your file ptr as the first parameter:
 - ifprintf(myFilePtr, "int %d", myint);

Strings

- lib/general/StringBasics.h(cpp)
- Alternative to std::string
 - With advantage of easily adding integers/doubles.
 - Capability of reading a word or line from file.
 - Methods for finding substrings.
- Methods to:
 - *Clear();*
 - *IsEmpty();*
 - *Length();*
 - *ToUpper();*
 - *ToLower();*
 - *Reverse();*
 - *First();*
 - *Last();*
 - *Split(char splitChar) returns vector of strings*
 - *Trim() (from front and back)*

String Operators

- Overloaded operators
 - `=, +, +=`
 - Can set to or append characters, strings, integers, doubles, and unsigned integers.
 - `==, !=, <, >, >=, <=` to other strings.
 - *int, double, long double* – convert to int, double, long double
 - `[]` - return the character at the index specified in brackets
- Read/Write to file
 - *Read, ReadLine, Write, WriteLine*

String Example

```
void LibraryExample::testString()
{
    String myString;
    int val1 = 2;
    double val2 = 3.1;
    const char* cstring = "abcd";
    String string2 = "efgh";
    myString = 1;
    myString += "+";
    myString += val1;
    myString += " + ";
    myString += val2;
    myString += cstring;
    myString += string2;
    std::cerr << "myString = " << myString << "\n"
        << "length = " << myString.Length() << "\n"
        << "isEmpty = " << myString.IsEmpty() << "\n";
    if(myString.Length() > 1)
    {
        std::cerr << "Index 1 = " << myString[1] << "\n";
    }
    myString.ToUpper();
    std::cerr << "Upper myString = " << myString << "\n";
    myString.Reverse();
    std::cerr << "Reverse myString = " << myString << "\n";
    myString.ToLower();
    std::cerr << "Lower myString = " << myString << "\n";

    myString.Clear();
}
```

```
myString = 1+2 + 3.100 abcdefgh
length = 20
isEmpty = 0
Index 1 = +
Upper myString = 1+2 + 3.100 ABCDEFGH
Reverse myString = HGFEDCBA 001.3 + 2+1
Lower myString = hgfedcba 001.3 + 2+1
```

StringArray

- lib/general/StringArray.h(cpp)
- Useful for tokenizing a string.
 - *int ReplaceColumns(const String & s, char ch = '\t');*
 - *int AddColumns(const String & s, char ch = '\t');*
 - *void Clear();*
 - *int Length() const;*
 - *Index operator []*
- Notes:
 - Read reads the entire file one line into each array entry
 - Write writes each array entry on a separate line.
 - WriteLine writes each array entry on the same line, tab delimited.

Parameters

- One easy way to read parameters
 - lib/Parameters.h(cpp)
- Use for
 - bool, int, double, String
- Long Parameters uses macros & is easy to use
 - Include Parameters.h
 - Declare & initialize the variables you want as parameters
 - Don't forget to initialize otherwise the values are undefined.
 - Set those variables with the type of parameter
 - Read the parameters

Long Parameters

MACRO Name	Parameters	Description
BEGIN_LONG_PARAMETERS	array	Start the parameter declaration creating the variable named array
LONG_PARAMETER_GROUP	label	Create a group named label
LONG_PARAMETER	label, boolptr	Store bool parameter --label in boolptr
LONG_INTPARAMETER	label, intptr	Store int parameter --label in intptr
LONG_DOUBLEPARAMETER	label, doubleptr	Store double parameter --label in doubleptr
LONG_STRINGPARAMETER	label, stringptr	Store String parameter --label in stringptr
END_LONG_PARAMETERS	none	End parameter declaration.

- Each label is the text string that the user should enter for that parameter as --label value
- The ptrs are the address(&) of a previously declared variable.
- Int, double, and String parameters read the next argument as the value.
- Bool parameters are flags that swap true/false each time it appears on the parameter line
 - if mybool is initialized to false, --mybool –mybool results in false
 - if mybool is initialized to true, --mybool –mybool results in true

ParameterList

- Declare a ParameterList
 - *ParameterList parmList;*
- Once the LONG_PARAMETERS have been defined, add them to the parameter list.
 - *paramList.Add(new LongParameters("Arguments", myParams);*
 - Replace "Arguments" with the title you want for your parameters
 - Replace myParams with the variable in BEGIN_LONG_PARAMETERS
- Read the parameters from argv
 - *paramList.Read(argc, argv);*
- Optionally print parameter settings
 - *paramList.Status();*

Long Parameters Example

```
//////////  
// Create variables to store the parameters  
String inFile = ""; // String from statgen/lib/general/  
String outFile = "-";  
bool booll = false; // bool parameter  
bool bool2 = true; // bool parameter  
int fieldNum = -1; // int parameter  
double expValue = -1; // double parameter  
  
// Declare the parameters  
BEGIN_LONG_PARAMETERS(longParameterList)  
    // using a parameter group is optional.  
    LONG_PARAMETER_GROUP("Required Arguments") // 1st parameter group  
    LONG_STRINGPARAMETER("in", &inFile)  
    LONG_PARAMETER_GROUP("Optional Arguments") // 2nd parameter group  
    LONG_STRINGPARAMETER("out", &outFile)  
    LONG_PARAMETER("booll", &booll)  
    LONG_PARAMETER("bool2", &bool2)  
    LONG_INTPARAMETER("fieldNum", &fieldNum)  
    LONG_DOUBLEPARAMETER("expValue", &expValue)  
    // End of the parameters.  
END_LONG_PARAMETERS();  
  
// Create a parameter list  
ParameterList paramList;  
// Add the parameters to the list.  
paramList.Add(new LongParameters ("Arguments",  
                                longParameterList));  
  
// Read the parameters from argv.  
paramList.Read(argc, argv);  
  
// Print the parameter status (This step is not necessary).  
paramList.Status();
```

Output of program with no arguments specified.

```
The following parameters are available. Ones with "[]" are in effect:  
  
Arguments  
    Required Arguments : --in []  
    Optional Arguments : --out [-], --booll, --bool2 [ON], --fieldNum [-1],  
                        --expValue [-1.0e+00]
```

Usage Example

```
void LibraryExample::handleFiles(const String& inFile, const String& outFile)
{
    IFILE inputFile = ifopen(inFile, "r"); // Open the input file.
    IFILE outputFile = ifopen(outFile, "w"); // Open the output file.

    // Read a line from the file.
    StringArray splitLine; String currentLine; int bufferSize = 10000; char buffer[bufferSize];
    // String to print at the end of each line.
    const char* endLine = "\tend of line\n"; unsigned int endLineLen = strlen(endLine);

    // Until the end of the input file.
    while(!ifeof(inputFile))
    {
        currentLine.ReadLine(inputFile); // Read a line using String.
        splitLine.ReplaceColumns(currentLine, ':'); // Tokenize on ':' using StringArray
        // For each column, write the column# & its contents to the output file.
        for(int i = 0; i < splitLine.Length(); i++)
        {
            ifprintf(outputFile, "Col %d: %s\t", i+1, splitLine[i].c_str());
        }
        if(ifwrite(outputFile, endLine, endLineLen) != endLineLen) // Write endLine
        {
            std::cerr << "ERROR writing\n"; // failed to write the correct amount.
        }
    }

    if(strcmp(inputFile->getFileName(), "-") != 0) // Can't rewind when reading from stdin
    {
        ifrewind(inputFile); // Go back to the beginning.
        int bytesRead = ifread(inputFile, buffer, bufferSize); // Read into buffer.
        // Print the number of bytes read.
        std::cerr << "Number Bytes in File = " << bytesRead << "\n";
    }

    ifclose(inputFile); ifclose(outputFile); // Close the files
}
```

Usage Script Example

```
mkdir -p out

./libraryExample 2> out/output.txt

# Run the String output
./libraryExample --bool1 2>> out/output.txt

# Run the file processing with text input, text output
./libraryExample --in files/myTestFile.txt --out out/updatedFromTxt.txt 2>> out/output.txt
# Run the file processing with gzip input, text output
./libraryExample --in files/myTestFile.txt.gz --out out/updatedFromGzip.txt 2>> out/output.txt
# verify same results from both types of output
diff out/updatedFromTxt.txt out/updatedFromGzip.txt

# Run the file processing with text input, gzip output
./libraryExample --in files/myTestFile.txt --out out/updatedFromTxtToGz.txt.gz 2>> out/output.txt
# Run the file processing with gzip input, gzip output
./libraryExample --in files/myTestFile.txt.gz --out out/updatedFromGzipToGz.txt.gz 2>> out/output.txt
# verify same results from both types of output
diff out/updatedFromTxtToGz.txt.gz out/updatedFromGzipToGz.txt.gz

# Run the file processing with gzip input, stdout
./libraryExample --in files/myTestFile.txt.gz 2>> out/output.txt | wc
# Run the file processing with gzip input, stdout
./libraryExample --in files/myTestFile.txt.gz 2>> out/output.txt > out/stdOut.txt
# compare stdout to the normal text output
diff out/updatedFromTxt.txt out/stdOut.txt

# Read from stdin
cat files/myTestFile.txt | ./libraryExample --in - 2>> out/output.txt > out/stdIn.txt
# compare stdin to the normal text input
diff out/updatedFromTxt.txt out/stdIn.txt

# compare stderr to expected.
diff expectedOutput.txt out/output.txt
```

Example Input File

Lecture 1: Statistical Computing: (Handout mode - PDF:Presentation mode - PDF)
Lecture 2: C++ Basics and Precisions: (Handout mode - PDF:Presentation mode - PDF)
Lecture 3: Implementing Fisher's Exact Test: (Handout mode - PDF:Presentation mode - PDF)
Lecture 4: Classes and STLs: (Handout mode - PDF:Presentation mode - PDF)
Lecture 5: Divide and Conquer Algorithms: (Handout mode - PDF:Presentation mode - PDF)
Lecture 6: Linear Sorting Algorithms and Elementary Data Structures: (Handout mode - PDF:Presentation mode - PDF)
Lecture 7: Data Structures: (Handout mode - PDF:Presentation mode - PDF)
Lecture 8: Hash Tables: (Handout mode - PDF:Presentation mode - PDF)
Lecture 9: Dynamic Programming: (Handout mode - PDF:Presentation mode - PDF)
Lecture 10: Boost Libraries and Graphical Algorithms: (Handout mode - PDF:Presentation mode - PDF)
Lecture 11: Hidden Markov Models: (Handout mode - PDF:Presentation mode - PDF)
Lecture 12: Hidden Markov Models: (Handout mode - PDF:Presentation mode - PDF)
Lecture 13: Matrix Computation: (Handout mode - PDF:Presentation mode - PDF)
Lecture 14: Implementing Linear Regression: (Handout mode - PDF:Presentation mode - PDF)
Lecture 15: Random Number Generation: (Handout mode - PDF:Presentation mode - PDF)
Lecture 16: Monte-Carlo methods and importance sampling: (Handout mode - PDF:Presentation mode - PDF)
Lecture 17: Numerical optimization: (Handout mode - PDF:Presentation mode - PDF)
Lecture 18: Numerical optimization II: (Handout mode - PDF:Presentation mode - PDF)
Lecture 19: StatGen Library: Notes coming soon...

Example Output File

Col 1: Lecture 1	Col 2: Statistical Computing	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 2	Col 2: C++ Basics and Precisions	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 3	Col 2: Implementing Fisher's Exact Test	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 4	Col 2: Classes and STLs	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 5	Col 2: Divide and Conquer Algorithms	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 6	Col 2: Linear Sorting Algorithms and Elementary Data Structures	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 7	Col 2: Data Structures	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 8	Col 2: Hash Tables	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 9	Col 2: Dynamic Programming	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 10	Col 2: Boost Libraries and Graphical Algorithms	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 11	Col 2: Hidden Markov Models	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
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Col 1: Lecture 17	Col 2: Numerical optimization	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 18	Col 2: Numerical optimization II	Col 3: (Handout mode - PDF)	Col 4: Presentation mode - PDF)	end of line
Col 1: Lecture 19	Col 2: StatGen Library	Col 3: Notes coming soon...	Col 4:	end of line

Library Documentation

- Wiki pages:
 - For all software, library & tools:
 - <http://genome.sph.umich.edu/wiki/Software>
 - Library specific documentation:
 - http://genome.sph.umich.edu/wiki/C%2B%2B_Library:_libStatGen
- Doxygen
 - http://www.sph.umich.edu/csg/mktrost/doxygen/march22_2011/
 - For list of Classes:
 - Select "Classes"

