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## CREWES Matlab® toolbox SEG-Y Input/Output update

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### ABSTRACT

The ability to read and write SEG-Y files into Matlab® using the CREWES toolbox has been evolving for a long time. Along the way, at least six separate attempts to code SEG-Y input/output (I/O) have been written by many people. None of this code was able to effectively deal with trace headers, custom trace headers, or with very large SEG-Y files. In addition, outputs were incompatible so that, for example, trace headers read from disk by one set of CREWES code could not be written to an output SEG-Y file using an unrelated set of CREWES code. This led to user frustration. A major re-write has been undertaken that combines ideas from legacy code in an object-oriented way that can handle very large datasets, trace headers and custom trace headers. Coding has also begun to support SEG-Y revision 2 which was released in January of 2017. Legacy code in the CREWES toolbox is being removed from the toolbox or being re-written as wrappers that call the new code for backwards compatibility. This report provides an overview of the current state of the new code, with examples of how to use it.

### INTRODUCTION

The ability to read and write SEG-Y files into Matlab® using the CREWES toolbox has been evolving for a long time. Along the way, at least six separate attempts to code SEG-Y input/output (I/O) have been written and modified by many people, some of whom documented their efforts (Hogan, 2004, Lloyd, 2010). All versions of SEG-Y code were stored concurrently in the CREWES toolbox, but none of it was compatible. This led to, for example, a SEG-Y file read from disk by one set of CREWES code could not be written to an output SEG-Y file using an unrelated set of CREWES code.

SEG-Y revision 0 (Barry et al., 1975) and revision 1 (Norris and Faichney, 2002) files were fairly well supported by CREWES code with the exception of trace header values. New code developed in 2010 was able to read trace headers (Lloyd et al., 2010), but was never optimized for speed. These codes were able to read IBM floating-point data, but not able to write it. CREWES code up to this point typically tried to read entire SEG-Y files into RAM, which was all right for small files but not good for files that were larger than the available memory on a given computer.

SEG-Y revision 2 (Hagelund and Stewart, 2017) as released in January of 2017, which prompted another look at these issues. A major re-write has been undertaken that combines ideas from legacy code in an object-oriented way that can handle very large datasets, trace headers and custom trace headers. This code aims to be more efficient, more robust, more flexible, and easier to maintain than past code.

Legacy code in the CREWES toolbox is being removed from the toolbox or being re-written as wrappers that call the new code for backwards compatibility. When this project is complete it will no longer be possible to read data with CREWES SEG-Y code only to be unable to write that data using a different set of CREWES SEG-Y code.

This report provides an overview of the current state of the new code. Rather than attempt to exhaustively document all features of code that is still under development, we show code examples that have been tested in Matlab® 2016b and 2017a. These examples can be copied from this document and pasted onto the Matlab® command line. All examples shown in this report are available in the file `crewes/seg/examplecode.m`

The example code that follows is organized by topic and numbered sequentially by the order in which the item appears in a SEG-Y file. For example, the textual file header examples in Example sections number 1.# and the binary file header examples are in sections 2.#, where # refers to 0) header definition information, 1) new header definition cell array, 2) new header value struct, 3) convert between struct and double, 4) file read and 5), file write.

The educational release of the CREWES Matlab® toolbox can be downloaded from <https://www.crewes.org/ResearchLinks/FreeSoftware>. The CREWES sponsor release can be downloaded from <https://www.crewes.org/ForOurSponsors/Software/matlab/crmatlab>.

## CLASSES AVAILABLE IN EDUCATIONAL RELEASE

### SegyTextHeader

Usage is: `thdr = SegyTextHeader(filename,permission,byteorder,segrevision,gui)`. See the Appendix for more details.

SegyTextHeader is a class for creating, reading and writing SEG-Y textual file headers. When a SegyTextHeader object is created for an existing SEG-Y file, the code attempts to guess if the text header is ASCII or EBCDIC format. The guess can be manually overridden setting the SegyRevision property after creating a SegyTextHeader object.

```
%TextHeader
%% Example 1.2: Create new textual file header
disp('*** Example 1.2: Create new textual file header ***')
thdr = SegyTextHeader; %Create a new SegyTextHeader object
txthdr=thdr.new; %Create a new SegyRevision 1 (default) textual file
header
thdr.SegyRevision = 0; %Update SegyRevision
txthdr=thdr.new; %Create a new SegyRevision 0 textual file header
thdr.SegyRevision = 2; %Update SegyRevision
txthdr=thdr.new; %Create a new SegyRevision 2 textual file header

%% Example 1.4: Read textual file header from disk
disp('*** Example 1.4: Read textual file header ***')
thdr = SegyTextHeader(ingyfile); %Create a new SegyTextHeader object,
TextFormat is guessed from file
txthdr = thdr.read; %Read textual file header from disk
thdr.TextFormat = 'ascii'; %Update TextFormat
txthdr = thdr.read; %Read textual file header from disk
thdr.TextFormat = 'ebcdic'; %Update TextFormat
txthdr = thdr.read; %Read textual file header from disk

%% Example 1.5: Write textual file header
disp('*** Example 1.5: Write textual file header ***')
```

```

thdr = SegyTextHeader(outsgyfile, 'w'); %Create a new SegyTextHeader
object with write permission
thdr.write(thdr.new); %Create and write a new textual file header to
disk
thdr.Permission = 'r'; %Update Permission
txthdr = thdr.read; %Read textual file header from disk

```

## SegyBinaryHeader

Usage is: *bhdr = SegyBinaryHeader(filename, permission, byteorder, segyrevision, gui)*. See the Appendix for more details.

SegyBinaryHeader is a class for creating, reading and writing SEG-Y binary file headers. When a SegyBinaryHeader object is created for an existing SEG-Y file, it guesses if the disk file is big ('b') or little ('l') endian byte-order by examining the data sample format code, which should be in the range 1-16. If the byte-order is incorrect, the format code will be read as a number greater than 255. This guess can be manually overridden by setting the ByteOrder property after creating a SegyBinaryHeader object.

Reading and writing a binary header word structure is governed by a binary header definition cell array with four columns, where the first column contains character strings that will be turned into struct fieldnames. The second column contains character strings specifying the data type for the SEG-Y header word. The data type can be any of the following: 'uint8', 'uint16', 'uint24', 'uint32', 'uint64', 'int8', 'int16', 'int24', 'int32', 'int64', 'ibm32', 'ieee32', and 'ieee64' where 'u' means unsigned, 'int' means integer, ibm means IBM floating point, ieee means IEEE floating point and the trailing number is the number of bits (8 bits in 1 byte) used to store the value. The third column contains the byte location (numeric) of the SEG-Y header word relative to the beginning of the binary file header. Note that the SEG-Y standard starts counting at byte 1, so the first header word, 'Job Identification Number', is at byte 1 in the standard, but this code starts counting at byte zero, so the start byte will be 0 in this case. The fourth column contains a long description of the header word as a character string

```

%% Binary Header
%% Example 2.0: Get binary header definition information
disp('*** Example 2.0: Get binary header definition information ***')
bhdr = SegyBinaryHeader; %Create a new SegyBinaryHeader object
bhdr.SegyRevision = 2; %Update SeGYRevision
bhdr.HdrDef; %Display current header definition cell array
[hwname, byteloc, idx] = bhdr.byte2word(20); %Return information about
header word closest to byte 20
bhdr.HdrDef(idx, :); %Display header word definition for row number idx

%% Example 2.1: Create a new binary header definition
disp('*** Example 2.1: Create new binary header definition ***')
bhdr = SegyBinaryHeader; %Create a new SegyBinaryHeader object
bindef = bhdr.newDefinition; %Create a new SegyRevision=1 (default)
header definition
bindef = bhdr.newDefinition(0); %Create a new SegyRevision 0 header
definition
bindef{50,1} = 'NewField'; %Update header word name for row 50 of the
definition
bhdr.HdrDef = bindef; %Update object's header definition

```

```
bhdr.HdrDef(50, :); %Display row 50 of object's definition

%% Example 2.2: Create a new binary header
disp('*** Example 2.2: Create new binary header ***')
bhdr = SegyBinaryHeader; %Create a new SegyBinaryHeader object
binhdr = bhdr.new; %Create a new binary header struct, using object
defaults
bhdr.SamplesPerTrace = 500; %Override number of samples per trace
bhdr.SampleInterval = 1000; %Override sample interval
[binhdr,bindef] = bhdr.new; %Create a new binary file header struct and
header definition

%% Example 2.4: Read binary file header
disp('*** Example 2.4: Read binary file header ***')
bhdr = SegyBinaryHeader(insgyfile); %Create a new SegyBinaryHeader
object, ByteOrder is guessed from file
binhdr = bhdr.read; %Read binary file header from disk
bhdr.ByteOrder = 'l'; %Update ByteOrder: 'l' is little-endian
binhdr = bhdr.read; %Read binary file header from disk
bhdr.ByteOrder = 'b'; %Update ByteOrder: 'b' is big-endian (SEG_Y
revision 0 and 1 standard)
binhdr = bhdr.read; %Read binary file header from disk

%% Example 2.5: Write binary file header to disk
disp('*** Example 2.5: Write binary file header to disk ***')
thdr = SegyTextHeader(outsgyfile, 'w'); %Create a new SegyTextHeader
object with write permission
thdr.write(thdr.new); %Create a new text header and write it to disk
bhdr = SegyBinaryHeader(outsgyfile, 'a'); %Create a new SegyBinaryHeader
object with append permission
nsamp = 10; %Samples per trace
sampint = 1000; %Sample interval in microseconds
bhdr.write(bhdr.new(nsamp, sampint)); %Write a new binary file header to
disk, override bhdr.SamplesPerTrace and bhdr.SampleInterval
bhdr.Permission = 'r'; %Update Permission
bhdr.read; %Read binary file header from disk
```

### SegyExtendedTextHeader

This is a stub. No code has been written or tested for creating, reading or writing extended textual file headers (if any) for SEG-Y revision 1 and 2 files.

### SegyTrace

Usage is: *trc = SegyTrace(filename, permission, fmtcode, byteorder, segyrevision, gui)*. See the Appendix for more details.

SegyTrace is a class for creating, reading and writing SEG-Y trace headers and trace data. When a SegyTrace object is created it uses a temporary SegyBinaryHeader object to get basic information from the binary file header which enables calculation of the number of traces in the file. Some of this information, for example, the number of samples per trace may be incorrect in the binary header. These properties be overridden before conducting read operations. If the files Segy Revision number is 0 and the binary header data sample format code is 1 (4-byte floating point), the class constructor tests the trace data to guess if

it is IBM or IEEE floating point. If the latter, the objects FormatCode is updated to 5 (IEEE 4-byte floating point; see Hall, 2017) At this time, this class does not handle variable length traces, although they can still be read by manually overriding properties such as OFFSET and SamplesPerTrace.

Reading and writing trace headers words is governed by a cell array, similar to the SegyBinaryHeader class, however, a trace header definition cell array has five columns. The first column contains header word name character strings that will be turned into struct fieldnames. The second column contains character strings specifying the data type for the SEG-Y header word. Data types can be any of the following: 'uint8', 'uint16', 'uint24', 'uint32', 'uint64', 'int8', 'int16', 'int24', 'int32', 'int64', 'ibm32', 'iee32', and 'iee64' where 'u' means unsigned, 'int' means integer, ibm means IBM floating point, iee means IEEE floating point and the trailing number is the number of bits (8 bits in 1 byte) used to store the value. The third column contains the byte location (numeric) of the SEG-Y header word relative to the beginning of the trace header. Note that the SEG-Y standard starts counting at byte 1 so the first header word, 'Trace Number in Line', is at byte 1 in the standard, but this code starts counting from zero so the start byte will be 0 in this example. The fourth column contains the name of the scalar header word that will be applied to the current header word if the ApplyCoordScalars property is set to 1 (default). The scalar header word name must also exist in column 1. The fifth column contains a long description of the header word as a character string.

```
%% Trace
%% Example 4.0: Get trace header definition information
disp('*** Example 4.0: Get trace header definition information ***')
trc = SegyTrace; %Create a new SegyTrace object
trc.HdrDef; %Display the current trace header definition cell array
[hwname, byteloc, idx] = trc.byte2word(20); %Get info about header word
closest to byte 20
trc.word2byte('EnsembleNum'); %Get byte number for a given header word
name
trc.HdrDef(idx,:); %Display header word definition for row number idx

%% Example 4.1: Create new trace header definition cell array
disp('*** Example 4.1: Create new trace header definition ***')
trc = SegyTrace; %Create a new SegyTrace object
trc.HdrDef{80,1} = 'NewField'; %Update field name for row 80 of the
definition
trc.HdrDef(80,:) %Display row 80 of HdrDef
trcdef = trc.newDefinition(0); %Create a new SegyRevision 0 header
definition
trcdef = trc.newDefinition(2); %Create a new SegyRrevision 2 header
definition
%Modify trcdef manually or by using uiSegyDefinition (not shown)
trc.HdrDef = trcdef; %Override object's header definition

%% Example 4.2: Create new trace header struct, data array and
definition
disp('*** Example 4.2: Create new trace header struct, data array and
definition ***')
trc = SegyTrace; %Create a new SegyTrace object
trcdef = trc.newDefinition(); %Return a new trace header definition
```

```
[trchdr, trcdat, trcdef] = trc.new(ntrace, nsamp, sampint); %Return new
trace header, trace data and trace definition
trc.SegyRevision = 0; %Update SegyRevision
trcdef = trc.newDefinition(); %Create a new SegyRevision 0 trace header
definition
[trchdr, trcdat, trcdef] = trc.new (ntrace, nsamp, sampint); %Return
new trace header, trace data and trace definition
trc.SegyRevision = 2; %Update SegyRevision
trcdef = trc.newDefinition; %Create a new SegyRevision 2 trace header
definition
[trchdr, trcdat, trcdef] = trc.new (ntrace, nsamp, sampint); %Return
new trace header, trace data and trace definition
```

```
%% Example 4.3: Convert between trace header struct and array of
doubles
disp('*** Example 4.3: Convert between trace header struct and array of
doubles ***')
[hdrvals,fieldnames,datatypes] = trc.struct2double(trchdr); %Convert
trace header struct to array of doubles
trchdr = trc.double2struct(hdrvals,fieldnames,datatypes); %Convert
array of doubles to a trace header struct
```

```
%% Example 4.4: Read trace headers and data
disp('*** Example 4.4: Read trace headers and data ***')
trc = SegyTrace(insgyfile); %Create a new SegyTrace object
[trchdr,trcdat,trcdef] = trc.read; %Read all trace headers and trace
data and return trace definition used
[trchdr,trchdr,trcdef] = trc.read(10:20); %Read traces 10-20
[trchdr,trchdr,trcdef] = trc.read(20:-1:10); %Read traces 10-20 in
reverse order
[trchdr,trchdr,trcdef] = trc.read([1,5,8]); %Read traces 1, 5 and 8
trchdr = trc.read([1,5,8], 'headers'); %Read trace headers only
trcdat = trc.read([1,5,8], 'data'); %Read trace data only
tnl = trc.read(1:100, 'TrcNumLine'); %Read header word TrcNumLine from
traces 1-100
trc.SegyRevision = 1; %Update SegyRevision
trc.ByteOrder = 'b'; %Update ByteOrder
trc.SamplesPerTrace = 1001; %Update SamplesPerTrace
[trchdr,trcdat] = trc.read; %Read all trace headers and trace data and
return trace definition used
%NOTE, SegyRevision, ByteOrder and SamplesPerTrace are incorrect,
trchdr and trcdat will be garbage!
```

```
%% Example 4.5: Write trace headers and data to disk
disp('*** Example 4.5: Write trace headers and data to disk ***')
thdr = SegyTextHeader(outsgyfile, 'w'); %Create a new SegyTrace object
with write permission
thdr.write(thdr.new); %Create a new text header and write to disk
bhdr = SegyBinaryHeader(outsgyfile, 'a'); %Create a new SegyBinaryHeader
object with append permission
bhdr.write(bhdr.new(nsamp, sampint)); %Create and write a new binary
header to disk
trc = SegyTrace(outsgyfile, 'a'); %Create a new SegyTrace object with
append permission
[trchdr,trcdat] = trc.new(ntrace, nsamp, sampint); %Create new trace
header and trace data
trc.write(trchdr, trcdat); %Write trace header and trace data to disk
```

## SegyDataTrailer

This is a stub. No code has been written or tested for creating, reading or writing data trailers for SEG-Y revision 2 files.

## SegyFile

Usage is *sf* = *SegyFile(filename,permission,segrevision,sampint,nsamps,segfmt,...  
txfmt,bytorder,bindef,trcdef,gui)*

The *SegyFile* class provides a convenient interface for *SegyTextHeader*, *SegyBinaryHeader*, *SegyExtendedTextHeader*, *SegyTrace* and *SegyDataTrailer*. If the *SegyRevision* or *ByteOrder* properties are changed at the top level in a *SegyFile* object, the change is propagated to all sub-objects.

```
%% SegyFile
%% Example 6.2: Create new file headers and trace data array
disp('*** Example 6.2: Create new file headers, trace data array, and
header definitions ***')
sf = SegyFile; %Create a new SegyFile object
[txthdr, binhdr, exthdr, trchdr, trcdat, bindef, trcdef] = sf.new;
%Create new file and trace headers, trace data and header defintions
[txthdr, binhdr, exthdr, trchdr, trcdat] = sf.new(ntrace, nsamp,
sampint); %Create new file and trace headers, trace data and header
defintions over-riding nsamp and sampint
%NOTE that exthdr will be empty ([])

%% Example 6.4: Read an existing SEG-Y file
disp('*** Example 6.4: Read SEG-Y file ***')
%In the following examples, exthdr will always be empty (exthdr = []).
This
%is a placeholder for when the ExtendedText Header class has been fully
written and tested.u
if exist('uiSegyFile.m','file')
    sf = uiSegyFile(insgyfile); %Create a new SegyFile object and
inspect SEG-Y file with GUI (sponsors toolbox release only)
end
sf = SegyFile(insgyfile); %Create a new SegyFile object
[txthdr, binhdr, exthdr, trchdr, trcdat, bindef, trcdef] = sf.read;
%Read entire file
[txthdr, binhdr, exthdr, trchdr, trcdat, bindef, trcdef] =
sf.read(1:2); %Read traces 1 and 2
txthdr = sf.TextHeader.read; %Read just the textual file header
binhdr = sf.BinaryHeader.read; %Read just the binary header
exthdr = sf.ExtendedTextHeader.read; %Read just the extended text file
header(s)
trchdr = sf.Trace.read(1,'headers'); %Read trace header 1
trchdr = sf.Trace.read([], 'headers'); %Read all trace headers
trcdat = sf.Trace.read(1:2:10,'data'); %Read data from traces 1-10 by
two's
trcdat = sf.Trace.read([], 'data'); %Read all trace data
[trchdr,trcdat] = sf.Trace.read; %Read all trace headers and trace data

%% Example 6.5: Write a new SEG-Y file: PROBLEMS
disp('*** Example 6.5: Write new SEG-Y file ***')
```

```
sf = SegyFile(insgyfile); %Create a new SegyFile object
[txthdr, binhdr, exthdr, trchdr, trcdat, bindef, trcdef] = sf.read;
%Read entire file
gui=1; %Command line prompts
% gui=[]; %GUI prompts
sf2 = SegyFile(outsgyfile, 'w', sf.SegyRevision, sf.SampleInterval, ...
    sf.SamplesPerTrace, sf.FormatCode, sf.TextFormat, sf.ByteOrder, ...
    bindef, trcdef, gui); %Create a new SegyFile object with write
permissions
sf2.write(txthdr, binhdr, exthdr, trchdr, trcdat, ...
    bindef, trcdef); %Write SEG-Y file to disk
```

## Wrappers

A number of wrapper functions are provided for convenience and for backwards compatability:

```
%% Wrappers
%% Example 7.0: Get trace header information
disp('*** Example 7.0: Get trace header information ***')
tracebyte2word(232); %Get header name closest to byte 20, revision 1
trace header definition (default)
tracebyte2word(232,0); %Get header name closest to byte 20, revision 0
trace header definition
tracebyte2word(232,2); %Get header name closest to byte 20, revision 2
trace header definition
traceword2byte('Unassigned01'); %Get byte location for header name,
revision 1 trace header definition (default)
traceword2byte('Unassigned14',0); %Get byte location for header name,
revision 0 trace header definition
traceword2byte('TrcHdrName',2); %Get byte location for header name,
revision 2 trace header definition
traceheaderdump(trchdr); %List all header names in trchdr struct that
contain non-zero values
[dump,words,inotempty]=traceheaderdump(trchdr); %Return all header
values, names and indices in trchdr struct for non-zero values
traceheaderdump_g(trchdr); %Display GUI that can plot values for up to
three separate header words

%% Example 7.4: Read an SEG-Y file
disp('*** Example 7.4: Read SEG-Y file ***')
%Read SEG-Y file with no overrides, display uiSegyFile() GUI if it
exists
[trcdat, segyrev, sampint, fmtcode, txtfmt, bytord, txthdr, ...
    binhdr, exthdr, trchdr, bindef, trcdef] = ...
    readsegy(insgyfile);
%Read SEG-Y file using all available overrides
%Update input paramters:
trcrange = []; %Empty => all traces
gui = 1; %Command line prompts, no GUI
nsamps = []; %Empty => Determine number of samples per trace from file
on disk
%Read SEG-Y file using all overrides, display uiSegyFile() GUI if it
exists
[trcdat, segyrev, sampint, fmtcode, txtfmt, bytord, txthdr, binhdr, exthdr, trchdr, ...
    dr, ...
```



```

    bindef, trcdef] =
readseggy(insgyfile, trcrange, segyrev, sampint, nsamps, ...
    fmtcode, txtfmt, bytord, bindef, trcdef, gui);

%% Example 7.5: Write a new SEG-Y file
disp('*** Example 7.5: Write SEG-Y file ***')
writeseqy(outsgyfile, trcdat); %Write Seg-Y revision 1 file using just
trcdat and defaults
writeseqy(outsgyfile, trcdat, segyrev, sampint, fmtcode, txtfmt, ...
    bytord, txthdr, binhdr, exthdr, trchdr, bindef, trcdef); %Write SEG-Y
file using all overrides

```

### ADDITIONAL CODE AVAILABLE IN SPONSOR RELEASE

Two graphical user interfaces (GUI) have been written to aid setting properties for SegyFile() objects. These are called uiSegyFile() and uiSegyDefinition().

#### uiSegyDefinition

Usage is *hdrdef = uiSegyDefinition()* or *hdrdef = uiSegyDefinition(hdrdef)*.

The uiSegyDefinition GUI displays a binary file header or trace header definition cell array in an editable table (Figure 1). Users may check the validity of the definition at any time by clicking the ‘Check Definition’ button. If the definition is not valid, the ‘Continue’ button will be disabled and changed from green to a red background color. Three pull-down menus are provided: 1) ‘File’ allows you to save the current displayed definition to a .mat file on disk, and to read it back into the table, 2) ‘New’ replaces the current definition with a new one, and 3) ‘Edit’ allows insertion, deletion and sorting (based on the valued of Start Byte) of rows in the table. The GUI returns the edited table as a header definition cell array when the ‘Continue’ button is clicked.

#### uiSegyFile

Usage is *sf = uiSegyFile(filename)* or *uiSegyFile()*

The uiSegyFile GUI (Figure 2) allows interactive inspection of an existing SEG-Y disk file and returns a SegyFile object when the ‘Continue’ button is clicked or if ‘Close’ is selected from the ‘File’ pull-down menu. Three pull-down menus are available, 1) ‘File’ allows you to open a different SEG-Y file, 2) ‘Edit’ calls uiSegyDefinition for either the current binary header definition or the current trace header definition, and 3) ‘Plot’ plots either a single trace header word or trace data values for up to 100 traces on either side of the current trace, using plotimage(). ‘Plot’ takes into account the radio buttons for ‘Time’, ‘Frequency’, ‘Gain Correction’, and ‘Bandpass Filter’ on the ‘Trace Display’ panel. ‘Gain Correction’ applies an automatic envelope correction using the aec() function in the CREWES toolbox. ‘Bandpass Filter’ applies a 10-15-55-60 Hz Ormsby filter using the filtorm() function in the CREWES toolbox. Bandpass frequency limits are not customizable at this time.

Figures 2 through 7 show some examples of using uiSegyfile() for a correlated Vibroseis source gather file.

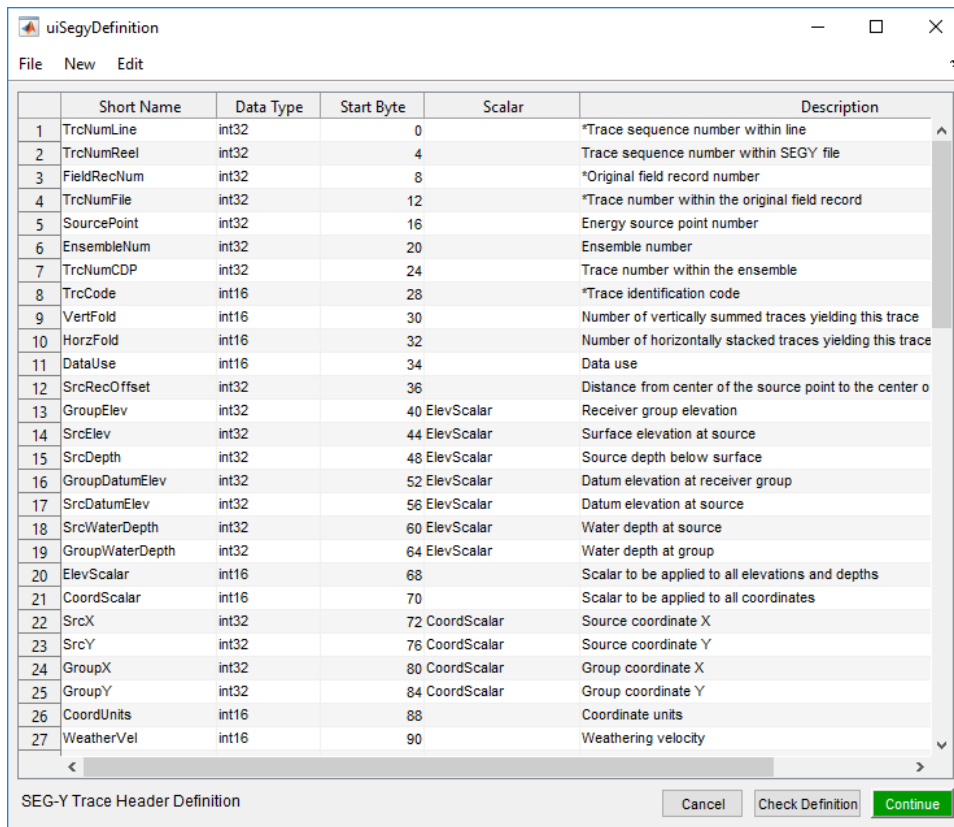


FIG. 1. GUI displayed for >> trcdef = uiSegyDefinition(sf.Trace.HdrDef)

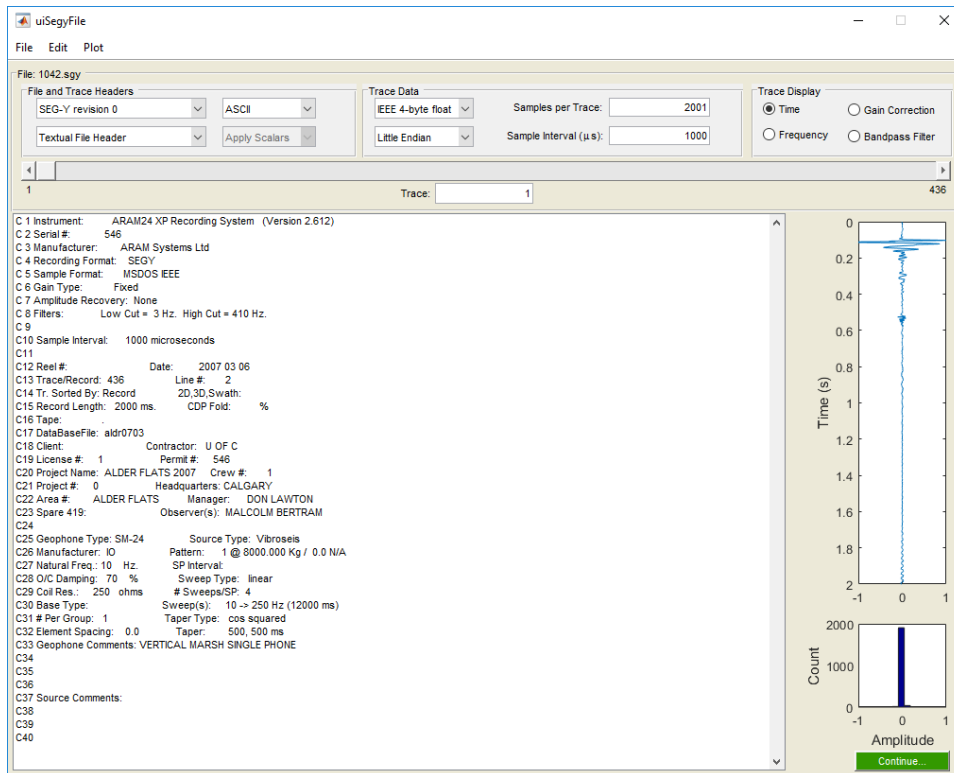


FIG. 2. Textual file header (table) and trace one data (right) are displayed.

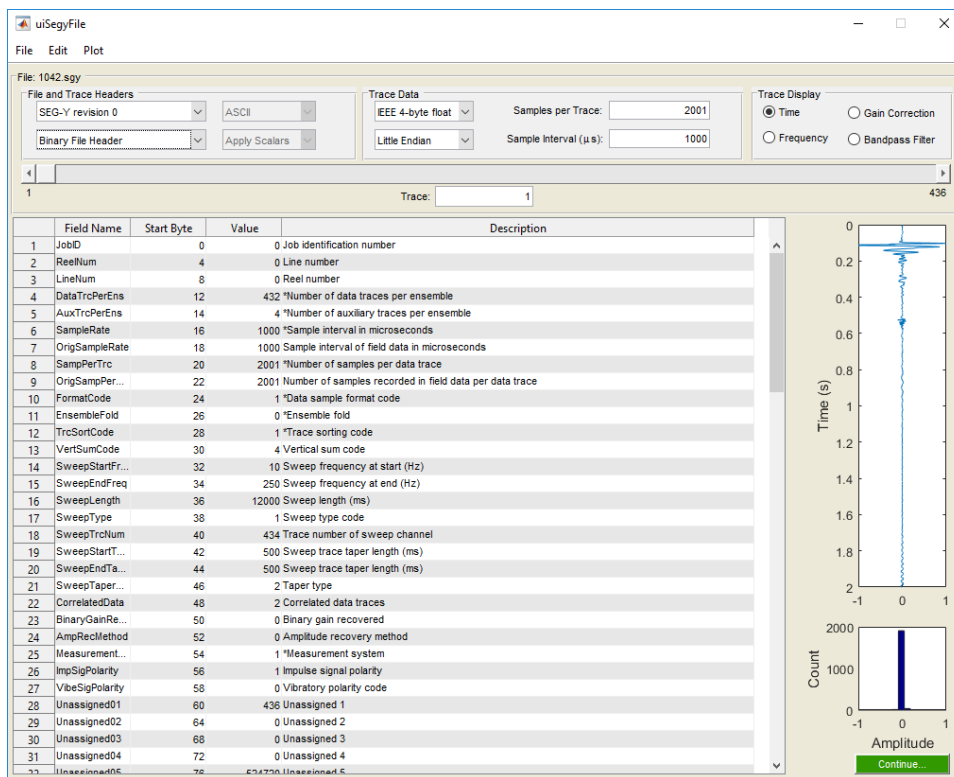


FIG. 3. Binary file header (table) and trace one data read correctly using little-endian byte order.

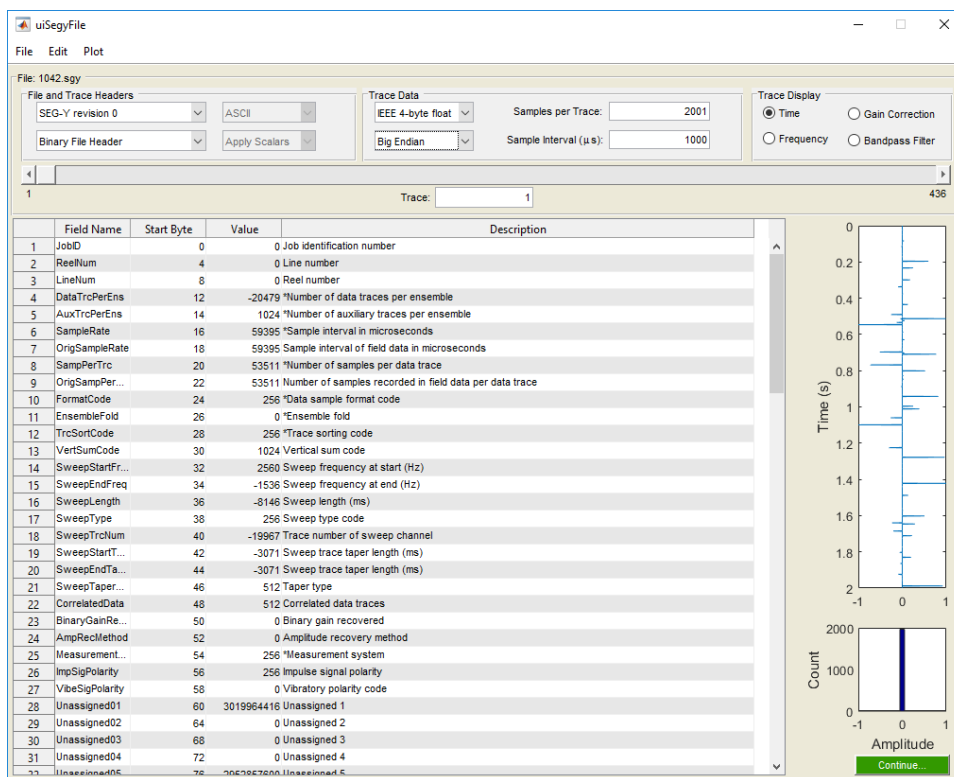


FIG. 4. Binary file header (table) and trace one data read incorrectly using big-endian byte order.

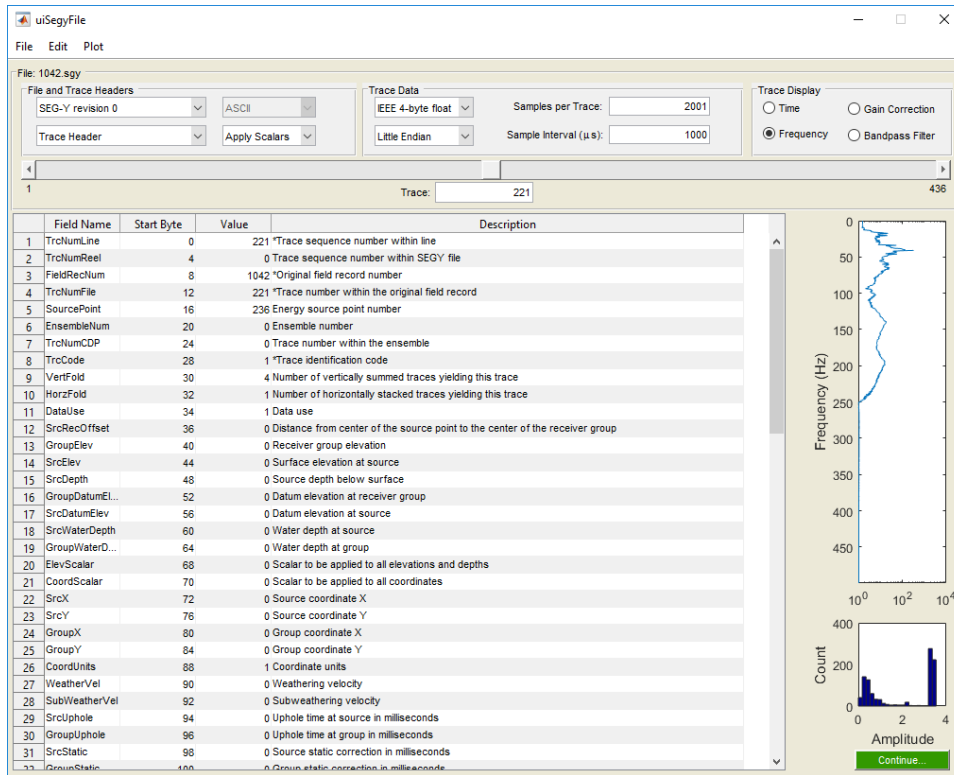


FIG. 5. Trace 221 header (table) and data read correctly using little-endian byte order, trace data displayed in frequency rather than time.

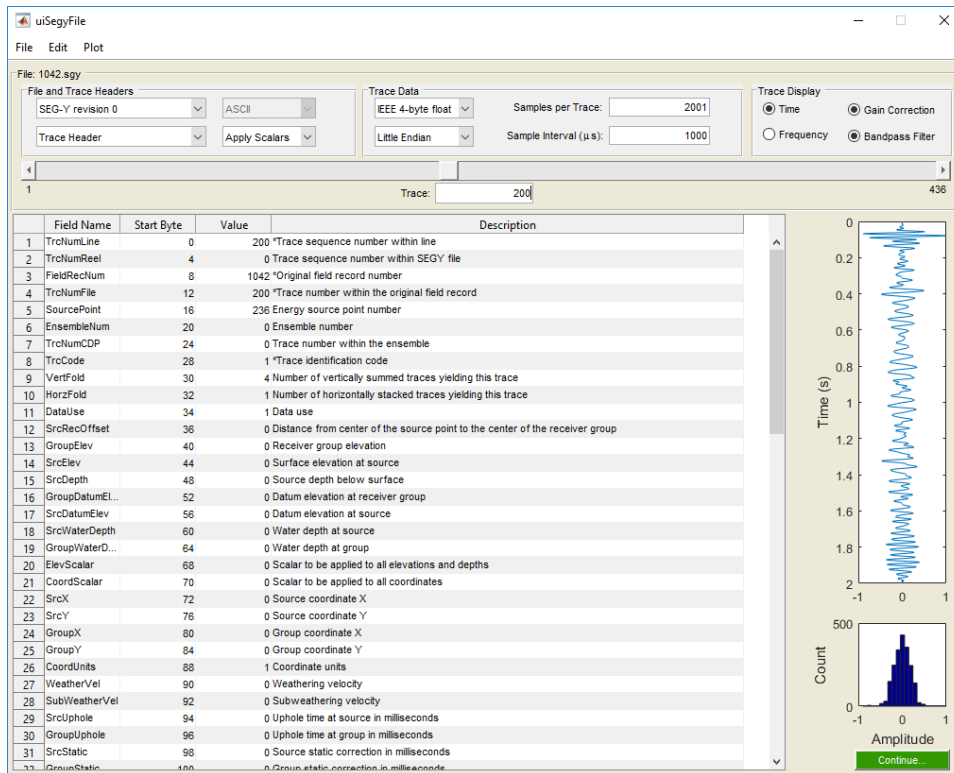


FIG. 6. Trace 200 trace header and trace data displayed with gain correction and bandpass filter.

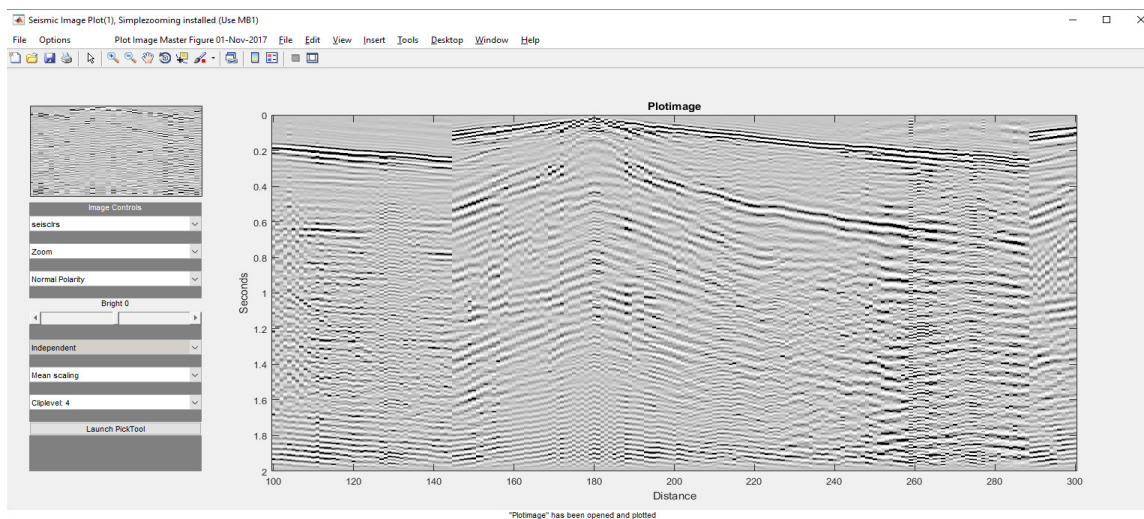


FIG. 7. Traces 100-300 plotted with gain correction and bandpass filter using the Plot->Traces pull-down menu,

## CONCLUSIONS AND FUTURE WORK

New object-oriented code to read and write SEG-Y files has been written and tested, but is not yet complete. However, it is more internally consistent, robust, useable and maintainable than any previous code released via the CREWES Matlab® toolbox. Development is on-going. All SEG-Y revision 0, 1 and 2 trace data formats are supported for reading and writing except for format code 4 (Fixed point with gain). Variable trace lengths in a file are not yet supported. SEG-Y revision 1 and 2 extended textual file headers are not yet supported. SEG-Y revision 2 extended trace headers and data trailers are not yet supported.

Future work includes writing code to address the issues listed above, removing incompatible code from the CREWES toolbox and writing wrappers for some older code for backwards compatibility (eg. `altreadsey()` and `altwriteseGY()`).

## ACKNOWLEDGEMENTS

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## **APPENDIX**

The following are class summaries that were auto-generated for the SegyTextHeader, SegyBinaryHeader SegyTrace and SegyFile classes by Matlab® based on help and comment lines contained within the code.

**MATLAB File Help:**  
SegyTextHeader

[View code for  
SegyTextHeader](#)

[Go to online doc for  
SegyTextHeader](#)

[Default  
Topics](#)

## SegyTextHeader

```
classdef SegyTextHeader
```

```
    SEG-Y textual file header class
```

```
Usage: thdr = SegyTextHeader(filename,permission,byteorder,segysrevision)
```

Where:

```
filename      = string containing full file name or a file id from fopen
permission    = string containing file permissions to use with fopen
                (optional), default is 'r' (see help fopen)
byteorder     = string containing 'b' or 'l' for big- or little-endian
                (optional), default is 'n' (see help fopen)
segysrevision = 0, 1, or 2
gui           = 0: no prompts,
                1: text prompts
                figure handle or empty: gui prompts
```

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### Class Details

**Superclasses**    [File](#)  
**Sealed**            false  
**Construct on load** false

### Constructor Summary

[SegyTextHeader](#)    classdef SegyTextHeader

### Property Summary

<a href="#">ByteOrder</a>	Byte Order, 'l' for little endian, 'b' for big endian
<a href="#">FileID</a>	File handle output from fopen
<a href="#">FileName</a>	FileName corresponding to FileID
<a href="#">GUI</a>	GUI flag: 0: no prompts, 1: text prompts (default), figure handle: gui prompts
<a href="#">OFFSET</a>	Header offset from beginning of file in bytes
<a href="#">Permission</a>	File Permission, eg. 'r', 'w', 'a'
<a href="#">SIZE</a>	Header size in bytes
<a href="#">SegyRevision</a>	SEG-Y revision number: 0, 1 (default), or 2

[TextFormat](#) Text format: 'ascii' or 'ebcdic'

## Method Summary

	<a href="#">addlistener</a>	Add listener for event.
	<a href="#">delete</a>	Delete a handle object.
	<a href="#">eq</a>	== (EQ) Test handle equality.
	<a href="#">fclose</a>	'fclose()' with error checking
	<a href="#">findobj</a>	Find objects matching specified conditions.
	<a href="#">findprop</a>	Find property of MATLAB handle object.
	<a href="#">fopen</a>	'fopen()' with error checking
	<a href="#">fprintf</a>	'fprintf()'
	<a href="#">fread</a>	'fread()' with error checking and additional data types
	<a href="#">freopen</a>	'freopen()' with error checking
	<a href="#">fseek</a>	'fseek()' with error checking
	<a href="#">fsize</a>	File size in bytes
	<a href="#">ftell</a>	'ftell()' with error checking
	<a href="#">fwrite</a>	'fwrite()' with error checking and addition datatypes
	<a href="#">ge</a>	>= (GE) Greater than or equal relation for handles.
	<a href="#">gt</a>	> (GT) Greater than relation for handles.
	<a href="#">guessTextFormat</a>	Guess if SEG-Y file textual file header is ASCII or EBCDIC
Sealed	<a href="#">isvalid</a>	Test handle validity.
	<a href="#">le</a>	<= (LE) Less than or equal relation for handles.
	<a href="#">listenByteOrder</a>	ByteOrder has changed, freopen file for file operations
	<a href="#">listenFileName</a>	FileName has changed, freopen file for file operations
	<a href="#">listenPermission</a>	Permission has changed, freopen file for file operations
	<a href="#">lt</a>	< (LT) Less than relation for handles.
	<a href="#">ne</a>	~= (NE) Not equal relation for handles.
	<a href="#">new</a>	Create and populate a new textual file header based on the SEG-Y revision number
	<a href="#">notify</a>	Notify listeners of event.
	<a href="#">read</a>	`Read a textual file header from a SEG-Y file
	<a href="#">write</a>	`Write a textual file header to a SEG-Y file

## Event Summary

<a href="#">ByteOrderChanged</a>	Notifies listeners that the ByteOrder has changed
<a href="#">FileNameChanged</a>	Notifies listeners that the FileName has changed
<a href="#">ObjectBeingDestroyed</a>	Notifies listeners that a particular object has been destroyed.
<a href="#">PermissionChanged</a>	Notifies listeners that the Permission has changed



## SegyBinaryHeader

```
classdef SegyBinaryHeader
```

```
    SEG-Y binary file header class
```

```
Usage:
```

```
    bh = SegyBinaryHeader(filename,permission,byteorder,segysrevision,gui)
```

```
Where:
```

```

filename    = string containing full file name or a file id from fopen
permission  = string containing file permissions to use with fopen
              (optional), default is 'r' (see help fopen)
byteorder   = string containing 'b' or 'l' for big- or little-endian
              (optional), default is 'n' (see help fopen)
segysrevision = 0, 1, or 2
gui         = 0: no prompts,
              1: text prompts
              figure handle or empty: gui prompts

```

```
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```

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### Class Details

**Superclasses** [File](#)  
**Sealed** false  
**Construct on load** false

### Constructor Summary

[SegyBinaryHeader](#) classdef SegyBinaryHeader

### Property Summary

<a href="#">ByteOrder</a>	Byte Order, 'l' for little endian, 'b' for big endian
<a href="#">FileID</a>	File handle output from fopen
<a href="#">FileName</a>	FileName corresponding to FileID
<a href="#">GUI</a>	GUI flag: 0: no prompts, 1: text prompts (default), figure handle: gui prompts
<a href="#">HdrDataTypes</a>	HdrDef col 2 (Dependent)
<a href="#">HdrDef</a>	Cell array containing the binary file header definition
<a href="#">HdrFieldNames</a>	HdrDef col 1 (Dependent)
<a href="#">HdrLongNames</a>	HdrDef col 4 (Dependent)
<a href="#">HdrStartBytes</a>	HdrDef col 3 (Dependent)
<a href="#">OFFSET</a>	Header offset from beginning of file in bytes
<a href="#">Permission</a>	File Permission, eg. 'r', 'w', 'a'

<a href="#">SIZE</a>	Header size in bytes
<a href="#">SegyRevision</a>	SEG-Y revision number: 0, 1 (default), or 2

## Method Summary

	<a href="#">addlistener</a>	Add listener for event.
	<a href="#">byte2word</a>	Return the header word nearest a byte location in the binary header
	<a href="#">check</a>	Determine if input is struct or cell array and pass control to checkDefinition or checkStruct
	<a href="#">checkDefinition</a>	Check a binary header definition cell array for validity
	<a href="#">checkStruct</a>	Compare a binary header struct to the current binary header definition
	<a href="#">delete</a>	Delete a handle object.
	<a href="#">eq</a>	== (EQ) Test handle equality.
	<a href="#">fclose</a>	'fclose()' with error checking
	<a href="#">findobj</a>	Find objects matching specified conditions.
	<a href="#">findprop</a>	Find property of MATLAB handle object.
	<a href="#">fopen</a>	'fopen()' with error checking
	<a href="#">fprintf</a>	'fprintf()'
	<a href="#">fread</a>	'fread()' with error checking and additional data types
	<a href="#">freopen</a>	'freopen()' with error checking
	<a href="#">fseek</a>	'fseek()' with error checking
	<a href="#">fsize</a>	File size in bytes
	<a href="#">ftell</a>	'ftell()' with error checking
	<a href="#">fwrite</a>	'fwrite()' with error checking and addition datatypes
	<a href="#">ge</a>	>= (GE) Greater than or equal relation for handles.
	<a href="#">gt</a>	> (GT) Greater than relation for handles.
	<a href="#">guessByteOrder</a>	Use format code to guess if file is big or little-endian
Sealed	<a href="#">isvalid</a>	Test handle validity.
	<a href="#">le</a>	<= (LE) Less than or equal relation for handles.
	<a href="#">listenByteOrder</a>	ByteOrder has changed, freopen file for file operations
	<a href="#">listenFileName</a>	FileName has changed, freopen file for file operations
	<a href="#">listenPermission</a>	Permission has changed, freopen file for file operations
	<a href="#">listenSegyRevision</a>	SegyRevision has changed, reset the header definition
	<a href="#">lt</a>	< (LT) Less than relation for handles.
	<a href="#">ne</a>	~= (NE) Not equal relation for handles.
	<a href="#">new</a>	Create and populate a new binary header struct for use with write()
	<a href="#">newDefinition</a>	Create a new binary header definition based on the SEG-Y revision number
	<a href="#">notify</a>	Notify listeners of event.
	<a href="#">read</a>	'Read a binary header struct from a SEG-Y file using the current binary header definition
	<a href="#">readExtSampleInterval</a>	Read extended sample interval [rev 2] (Does not use the Header Definition)
	<a href="#">readExtSamplesPerTrace</a>	Read extended samples per trace [rev 2] (Does not use the Header Definition)
	<a href="#">readExtTracesPerRec</a>	Read extended traces per record [rev 2] (Does not use the Header

	Definition)
<a href="#">readFileInfo</a>	Read basic information from a SEG-Y file (Does not use header definition)
<a href="#">readFixedTrcLength</a>	Read fixed trace length flag [rev 1+] (Does not use the Header Definition)
<a href="#">readFormatCode</a>	Read data sample format code (Does not use the Header Definition)
<a href="#">readHeaderWord</a>	Read a single header word from a given byte position and data type
<a href="#">readIntegerConstant</a>	Read integer constant [rev 2] (Does not use the Header Definition)
<a href="#">readNumDataTrailers</a>	Read number of data trailers [rev 2] (Does not use the Header Definition)
<a href="#">readNumExtHeaders</a>	Read number of extended textual file headers [rev 1+] (Does not use the Header Definition)
<a href="#">readNumExtTrcHeaders</a>	Read number of extended trace headers [rev 2] (Does not use the Header Definition)
<a href="#">readSampleInterval</a>	Read sample interval (Does not use the Header Definition)
<a href="#">readSamplesPerTrace</a>	Read samples per trace (Does not use the Header Definition)
<a href="#">readSegyRevision</a>	Read Segy Revision [rev 1+] (Does not use the Header Definition)
<a href="#">readTraceOneOffset</a>	Read trace one offset [rev 2] (Does not use the Header Definition)
<a href="#">readTracesPerRec</a>	Read traces per record (Does not use the Header Definition)
<a href="#">write</a>	Write a binary file header structure to a SEG-Y file

### Event Summary

<a href="#">ByteOrderChanged</a>	Notifies listeners that the ByteOrder has changed
<a href="#">FileNameChanged</a>	Notifies listeners that the FileName has changed
<a href="#">ObjectBeingDestroyed</a>	Notifies listeners that a particular object has been destroyed.
<a href="#">PermissionChanged</a>	Notifies listeners that the Permission has changed
<a href="#">SegyRevisionChanged</a>	Notifies listeners that the SegyRevision has changed

## SegyTrace

```
classdef SegyTrace
```

```
Class for SEG Y Traces (Headers and Data)
```

```
Usage:
```

```
trc = SegyTrace(filename,permission,fmtcode,byteorder,segrevision,gui)
```

```
Where:
```

```

filename      = string containing full file name or a file id from fopen
permission    = string containing file permissions to use with fopen
                (optional), default is 'r' (see help fopen)
byteorder     = string containing 'b' or 'l' for big- or little-endian
                (optional), default is 'n' (see help fopen)
segrevision   = 0, 1, or 2
gui           = 0: no prompts,
                1: text prompts
                figure handle or empty: gui prompts

```

```
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```

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### Class Details

**Superclasses**    [File](#)  
**Sealed**            false  
**Construct on load** false

### Constructor Summary

```
SegyTrace    classdef SegyTrace
```

### Property Summary

<a href="#">ApplyCoordScalars</a>	Apply Coordinate Scalars, default=true
<a href="#">ByteOrder</a>	Byte Order, 'l' for little endian, 'b' for big endian
<a href="#">BytesPerSample</a>	Bytes Per Sample, eg. uint8 is 1, uint32 is 4,...
<a href="#">ExtSampleInterval</a>	Extended Sample interval [rev 2], default = 0
<a href="#">ExtSamplesPerTrace</a>	Extended Samples Per Trace [rev2], default = 0
<a href="#">ExtTracesPerRec</a>	Extended Traces Per Record [rev 2], default = 0
<a href="#">FileID</a>	File handle output from fopen
<a href="#">FileName</a>	FileName corresponding to FileID
<a href="#">FixedTrcLength</a>	Fixed Trace Length flag [rev 1+], default = 1
<a href="#">FormatCode</a>	Format Code, default = 5, IEEE 4-byte float
<a href="#">FormatCodeType</a>	Text equivalent to the numeric Format Code
<a href="#">GUI</a>	GUI flag: 0: no prompts, 1: text prompts (default), figure handle: gui prompts

<a href="#">HdrDataTypes</a>	Header Data Types, HdrDef col 2
<a href="#">HdrDef</a>	Trace Header Definition (cell array)
<a href="#">HdrFieldNames</a>	Trace Header Field Names, HdrDef col 1
<a href="#">HdrLongName</a>	Header Descriptive (Long) name, HdrDef col 5
<a href="#">HdrScalars</a>	Header Scalars, HdrDef col 4
<a href="#">HdrStartBytes</a>	Header Word Start Bytes, HdrDef col 3
<a href="#">IntegerConstant</a>	Integer constant [rev 2], default = 0
<a href="#">NumExtTrcHeaders</a>	Number of Extended Trace Headers [rev 2], default = 0
<a href="#">NumExtTxtHeaders</a>	Number of Extended Textual File headers [rev 1+], default = 0
<a href="#">OFFSET</a>	Bytes from beginning of file to start of trace one
<a href="#">Permission</a>	File Permission, eg. 'r', 'w', 'a'
<a href="#">SIZE</a>	Size of trace header in bytes
<a href="#">SampleInterval</a>	Sample Interval in microseconds, default = 1000
<a href="#">SamplesPerTrace</a>	Samples Per Trace
<a href="#">SegyRevision</a>	SEG-Y revision number, 0, 1 (default), 2
<a href="#">TraceSize</a>	Size of one trace in bytes
<a href="#">TracesInFile</a>	Total number of traces in file
<a href="#">TracesPerRec</a>	Traces Per Record, includes data and aux traces

## Method Summary

	<a href="#">addlistener</a>	Add listener for event.
	<a href="#">applyCoordinateScalars</a>	Apply coordinate scalars to trace header values
	<a href="#">byte2word</a>	Return the header word name nearest a byte location in the trace header definition
	<a href="#">check</a>	Call checkDefinition(), checkStruct() or checkArray() as appropriate
	<a href="#">checkArray</a>	Check trace data values to see if they can be stored using the current FormatCode
	<a href="#">checkDefinition</a>	Check a trace header definition cell array for validity
	<a href="#">checkStruct</a>	Compare a trace header struct to the current trace header definition for validity
	<a href="#">delete</a>	Delete a handle object.
Static	<a href="#">double2struct</a>	Converts an array of doubles to a trace header struct
	<a href="#">eq</a>	== (EQ) Test handle equality.
	<a href="#">fclose</a>	'fclose()' with error checking
	<a href="#">findobj</a>	Find objects matching specified conditions.
	<a href="#">findprop</a>	Find property of MATLAB handle object.
	<a href="#">fopen</a>	'fopen()' with error checking
	<a href="#">fprintf</a>	'fprintf()'
	<a href="#">fread</a>	'fread()' with error checking and additional data types
	<a href="#">freopen</a>	'freopen()' with error checking
	<a href="#">fseek</a>	'fseek()' with error checking
	<a href="#">fsize</a>	File size in bytes
	<a href="#">ftell</a>	'ftell()' with error checking
	<a href="#">fwrite</a>	'fwrite()' with error checking and addition datatypes

	<a href="#">ge</a>	>= (GE) Greater than or equal relation for handles.
	<a href="#">gt</a>	> (GT) Greater than relation for handles.
	<a href="#">guessFormatCode</a>	Guess if trace data are IBM32 or IEEE32 for SegyRevision=0 and FormatCode=1
Sealed	<a href="#">isvalid</a>	Test handle validity.
	<a href="#">le</a>	<= (LE) Less than or equal relation for handles.
	<a href="#">listenByteOrder</a>	ByteOrder has changed, freopen file for file operations
	<a href="#">listenFileName</a>	FileName has changed, freopen file for file operations
	<a href="#">listenPermission</a>	Permission has changed, freopen file for file operations
	<a href="#">lt</a>	< (LT) Less than relation for handles.
	<a href="#">ne</a>	~= (NE) Not equal relation for handles.
	<a href="#">new</a>	Returns a new trace data matrix (all zeros) and a new trace header struct
	<a href="#">newDefinition</a>	Returns a new trace header definition cell array based on SegyRevision
Static	<a href="#">newHeaderWord</a>	Returns a new zero-filled vector of the appropriate data type
	<a href="#">notify</a>	Notify listeners of event.
	<a href="#">read</a>	Reads and returns trace data and/or a trace header struct from a SEG-Y file
	<a href="#">removeCoordinateScalars</a>	Remove coordinate scalars from trace header values
Static	<a href="#">struct2double</a>	Converts a trace header struct to an array of doubles
	<a href="#">word2byte</a>	Return the byte location for a given header word name
	<a href="#">word2idx</a>	Returns the index number in HdrDef for a given header word name
	<a href="#">write</a>	Writes a trace data array and a trace header structure to a SEG-Y file

### Event Summary

<a href="#">ByteOrderChanged</a>	Notifies listeners that the ByteOrder has changed
<a href="#">FileNameChanged</a>	Notifies listeners that the FileName has changed
<a href="#">ObjectBeingDestroyed</a>	Notifies listeners that a particular object has been destroyed.
<a href="#">PermissionChanged</a>	Notifies listeners that the Permission has changed

## SegyFile

```
function sf = SegyFile(filename,permission,segrevision,sampint,nsamp,...
    fmtcode,txtfmt,byteorder,bindef,trcdef,gui)
Optional Inputs ([] is accepted):
filename - SEG-Y disk file name
permission - 'r' (default), 'w', or 'a' (see help fopen)
segrevision - segy revision (0,1,2); Overrides SEG-Y Binary File Header on
disk.
sampint - Sample interval in (s); Overrides SEG-Y Binary File Header on
disk.
nsamps - Samples per trace; Overrides SEG-Y Binary File Header on
disk.
fmtcode - SEG-Y trace data format code. eg. 1 = IBM float, 5 = IEEE
float
txtfmt - Text format, 'ascii' or 'ebcdic'
byteorder - byte order of disk file 'l'=little-endian, 'b'=big-endian
bindef - 4 column binary header definition cell array such as provided by
@BinaryHeader/new; See uiSegyDefinition().
NOTE! writesegy will require the same bindef unless you
modify binhdr!
trcdef - 5 column trace header definition cell array such as provided by
@BinaryHeader/new; See uiSegyDefinition()
NOTE! writesegy will require the same trcdef unless you
modify trchdr!
gui - 0 (no progress bar), 1 (text progress bar),
[] (default; gui progress bar and warnings), figure handle
(same as [], but an attempt is made to center GUI popups on
the figure represented by the figure handle)

Outputs:
sf - A SEG-Y file object
```

### Example:

```
s=SegyFile('file.sgy','r')
s.TextHeader.read
s.BinaryHeader.read
s.Trace.read(1:2:100)
```

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## Class Details

**Superclasses** [File](#)  
**Sealed** false  
**Construct on load** false

## Constructor Summary

[SegyFile](#) function sf = SegyFile(filename,permission,segrevision,sampint,nsamp,...

## Property Summary

<a href="#">BinaryHeader</a>	Contains BinaryHeader object
<a href="#">ByteOrder</a>	Byte Order, 'l' for little endian, 'b' for big endian
<a href="#">ExtendedTextHeader</a>	Contains SegyExtendedTextHeader object
<a href="#">FileID</a>	File handle output from fopen
<a href="#">FileName</a>	FileName corresponding to FileID
<a href="#">FormatCode</a>	Data sample format code: 1,2,3,5,6,7,8,9,10,11,12,15,16
<a href="#">GUI</a>	GUI flag: 0: no prompts, 1: text prompts (default), figure handle: gui prompts
<a href="#">Permission</a>	File Permission, eg. 'r', 'w', 'a'
<a href="#">SampleInterval</a>	Sample interval in microseconds
<a href="#">SamplesPerTrace</a>	Number of data samples per trace
<a href="#">SegyRevision</a>	Segy revision number: 0, 1 (default), 2
<a href="#">TextFormat</a>	Text format: 'ascii' or 'ebcdic'
<a href="#">TextHeader</a>	Contains SegyTextHeader object
<a href="#">Trace</a>	Contains SegyTrace object

## Method Summary

<a href="#">addlistener</a>	Add listener for event.
<a href="#">delete</a>	Delete a handle object.
<a href="#">eq</a>	== (EQ) Test handle equality.
<a href="#">fclose</a>	'fclose()' with error checking
<a href="#">findobj</a>	Find objects matching specified conditions.
<a href="#">findprop</a>	Find property of MATLAB handle object.
<a href="#">fopen</a>	'fopen()' with error checking
<a href="#">fprintf</a>	'fprintf()'
<a href="#">fread</a>	'fread()' with error checking and additional data types
<a href="#">freopen</a>	'freopen()' with error checking
<a href="#">fseek</a>	'fseek()' with error checking
<a href="#">fsize</a>	File size in bytes
<a href="#">ftell</a>	'ftell()' with error checking
<a href="#">fwrite</a>	'fwrite()' with error checking and addition datatypes
<a href="#">ge</a>	>= (GE) Greater than or equal relation for handles.
<a href="#">gt</a>	> (GT) Greater than relation for handles.
Sealed	
<a href="#">isvalid</a>	Test handle validity.
<a href="#">le</a>	<= (LE) Less than or equal relation for handles.
<a href="#">listenByteOrder</a>	ByteOrder has changed, freopen file for file operations
<a href="#">listenFileName</a>	FileName has changed, fopen file for file operations
<a href="#">listenFormatCode</a>	FormatCode has changed, update binary header and trace objects
<a href="#">listenGUI</a>	GUI has changed, update header and trace objects
<a href="#">listenPermission</a>	Permission has changed, freopen file for file operations
<a href="#">listenSampleInterval</a>	Sample Interval has changed, update binary header and trace objects
<a href="#">listenSamplesPerTrace</a>	SamplesPerTrace has changed, update binary header and trace objects
<a href="#">listenSegyRevision</a>	SegyRevision has changed, update header and trace objects



---

<a href="#">listenTextFormat</a>	TextFormat has changed, update text header objects
<a href="#">lt</a>	< (LT) Less than relation for handles.
<a href="#">ne</a>	~= (NE) Not equal relation for handles.
<a href="#">new</a>	Returns new text header (char), binary header (struct), trace header (struct) and trace data (double)
<a href="#">notify</a>	Notify listeners of event.
<a href="#">read</a>	Reads a SEG-Y file
<a href="#">write</a>	Writes a new SEG-Y file

### Event Summary

<a href="#">ByteOrderChanged</a>	Notifies listeners that the ByteOrder has changed
<a href="#">FileNameChanged</a>	Notifies listeners that the FileName has changed
<a href="#">FormatCodeChanged</a>	Notifies listeners that the FormatCode has changed
<a href="#">GUIchanged</a>	Notifies listeners that the GUI has changed
<a href="#">ObjectBeingDestroyed</a>	Notifies listeners that a particular object has been destroyed.
<a href="#">PermissionChanged</a>	Notifies listeners that the Permission has changed
<a href="#">SampleIntervalChanged</a>	Notifies listeners that the SamplesInterval has changed
<a href="#">SamplesPerTraceChanged</a>	Notifies listeners that the SamplesPerTrace has changed
<a href="#">SegyRevisionChanged</a>	Notifies listeners that the SegyRevision has changed
<a href="#">TextFormatChanged</a>	Notifies listeners that the TextFormat has changed