file

Run

FastaToString.reader(file)

Creates class
Sequences for
all Sequence
entries

option

Option is a
shorthand for
the relevant
Sequences method

pattern
- remove
- browse

sort
(sa || sd)

outFile

Class Sequences

Methods		Attributes	Class Sequence	
			Methods	**Attributes**
• Constructor	• browse(String)	• size		
• add()	• removeSeqs(List<Sequence>)	• elements	• Constructor	• id
• getIds()	• sortLength(String)	• seqList (List<Sequence>)		• seq
• getSeqs()	• writeFasta(String)			• length
• getLength()				