```
file
```

Run → FastaToString.reader(file)

option

pattern
- remove
- browse
- sort (sa || sd)
- none

→ 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()				