Sequence

- type
- seq
- _encoding_type
- _header
- _iupac
- legalDNA
- legalRNA
- legalAmino
- validType

+ begin()
+ end()
+ Sequence()
+ Sequence()
+ Sequence()
+ getSeq()
+ getType()
+ getEncodingType()
+ getIupac()
+ getTypeArray()
  and 19 more...
+ getSequenceSymbol()
+ getSymbolDNA()
+ isLegalNucleic()
+ isLegalAmino()
+ isLegalSymbol()
+ isDNA()
+ isRNA()
+ isAmino()
+ isNucleic()
+ isDnaSpecificNucleotide()
  and 14 more...
parseChar()

FastaSequence

- comments

+ getAllComment()
+ FastaSequence()
+ insert()
+ eraseSeq()
+ addComment()
+ loadComments()
+ removeComment()
+ displaceComment()
+ isCommentedPos()
+ getComment()