

| BitVector |
|---|
| <ul style="list-style-type: none"> - _data - _data_size - _element_number - _element_size |
| <ul style="list-style-type: none"> + BitVector() + BitVector() + BitVector() + setElementNumber() + ~BitVector() + getDataSize() + getElementNumber() + getElementSize() + size() + currentElementCapacity() and 36 more... + maximumOctetNumber() + maximumElementNumber() + copyBits() # _killInternalData() |

| Header |
|--|
| <ul style="list-style-type: none"> - _comment - _accession |
| <ul style="list-style-type: none"> + getComment() + setComment() + getAccession() + setAccession() |

-seq

-_header

| Sequence |
|--|
| <ul style="list-style-type: none"> - type - _encoding_type - _iupac - legalDNA - legalRNA - legalAmino - validType |
| <ul style="list-style-type: none"> + 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() |

| FastaRelated |
|---|
| <ul style="list-style-type: none"> - legalMultiHeaderChar - legalMonoHeaderChar - legalCommentChar - blankChar |
| <ul style="list-style-type: none"> + isHeaderChar() + isMonoHeaderChar() + isMultiHeaderChar() + isCommentChar() + isBlankChar() |

| FastaSequence |
|--|
| <ul style="list-style-type: none"> - comments |
| <ul style="list-style-type: none"> + getAllComment() + FastaSequence() + insert() + eraseSeq() + addComment() + loadComments() + removeComment() + displaceComment() + isCommentedPos() + getComment() |