```
taxon pair
+ taxon
+ protein start

    protein end

    listParseData

- listStructData

    listTaxa

+ getVariables()
+ print()
+ has data()
+ taxon pair()
+ taxon pair()
+ initList()
+ free taxonPairList()
+ init taxon pair()
+ assert class()
            listPair
     pipe bucket

    listParseData

    listStructData

   - listTaxa

    log

   - is inpa
   - listPair
   - list pair pos
   + pipe bucket()
   + operator()()
   + free data()
   + free mem()
   + assert class()
   - init taxon p()
```