

REFERENZE :

Software

- Protein Tools 7.6.1* (realizzato appositamente dall'autore e di imminente distribuzione negli archivi pubblici su Internet) Per averne una versione, sviluppata in Future Basic per Macintosh, contattare direttamente l'autore (paolo@icgeb.trieste.it).
- 3D Structure Analyzer 1.2b* (realizzato appositamente dall'autore) Per averne una versione di lavoro, sviluppata in Future Basic per Macintosh, contattare direttamente l'autore (paolo@icgeb.trieste.it).
- SOMA Classifier 6.0* (realizzato appositamente dall'autore) Per averne una versione di lavoro, sviluppata in Think C ++ e Unix C, contattare direttamente l'autore (paolo@icgeb.trieste.it).
- Netscape Navigator 2.0*. Per la connessione con le banche dati su Internet.
- Rasmol 2.0* Per la grafica molecolare.
- MacMolecule* Per la grafica molecolare.
- ADE Tools package* Per l'analisi multivariata.
- DALI* Per la classificazione di strutture tridimensionali di proteine.(EMBL Archive)
- GCG package* Per l'analisi delle strutture primarie

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