

- Claros, M.G. & von Heijne, G. (1994) TopPred II: an improved software for membrane protein structure predictions. *Comput. Applic. Biosci.* 10, 685-686
- von Heijne, G. (1992) *J. Mol. Biol.* 225, 487-494
(algorithm for transmembrane segments, GvH1 scale and topology of prokaryotic proteins)
- Sipos,L. & von Heijne, G. (1993) *Eur. J. Biochem.* 213, 1333-1340
(topology of eukaryotic proteins)
- Persson, B. & Argos, P. (1994) *J. Mol. Biol.* 237, 182-192
(the method used with the PA scale to determine transmembrane domains)
- Jähnig, F. (1990) *Trends Biochem. Sci.* 15, 93-95
(a different method to find amphiphilic transmembrane domains with alpha o beta folding)
- P. Klein, M. Kanehisha, C. DeLisi (1985) *Biochim. Biophys. Acta* 815, 468-476
(classification into integral, peripheral or non-membrane hydrophobic protein)