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**Título:** The Ten Most Wanted Solutions In Protein Bioinformatics

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**Precio:** \$1218.00

**Editorial:**

**Año:** 2005

**Tema:**

**Edición:** 1<sup>a</sup>

**Sinopsis**

**ISBN:** 9781584884910

Utilizing high speed computational methods to extrapolate to the rest of the protein universe, the knowledge accumulated on a subset of examples, protein bioinformatics seeks to accomplish what was impossible before its invention, namely the assignment of functions or functional hypotheses for all known proteins.