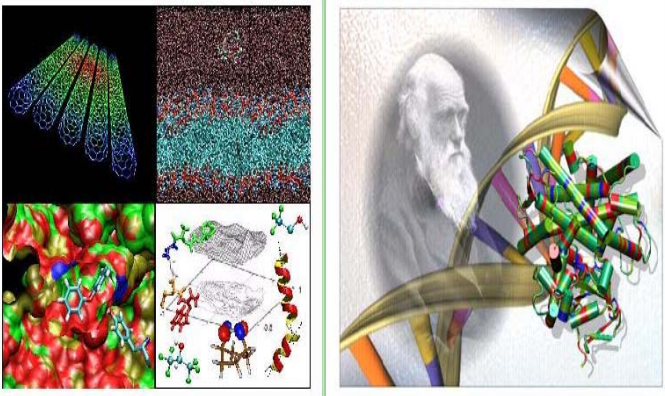


Location: <http://map.iu-bremen.de/>

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Contact: d.rocattano@iu-bremen.de and u.schwaneberg@iu-bremen.de

Location: <http://lynx.clm.u-bremen.de/SeSaM-Tv.html>

Transversion-favored Sequence Saturation Mutagenesis Program

Reference: [Disclaimer](#)

Step 1: Insert sequence (formats: FASTA, GenBank, raw sequence data):

From File:

Step 2: Select types of DNA fragments ([Help](#)):

☐ A ☐ T ☐ G ☐ C ☐ N

Step 3: Select types of degenerate bases ([Help](#)):

☐ P ☐ K ☐ I

For further details email: d.rocattano@iu-bremen.de, u.schwaneberg@iu-bremen.de

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Mutagenesis Assistant Program

Reference: Wong TS, Rocattano D, Zacharias M, Schwaneberg U. 2005. A statistical analysis of current random mutagenesis methods for directed protein evolution

[Disclaimer](#)

Insert sequence (formats: FASTA, GenBank, raw sequence data):

From File:

For further details email: d.rocattano@iu-bremen.de and u.schwaneberg@iu-bremen.de

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