This folder contains work towards determining which BBTM matrices ot use at which sequence identities. It very much overlaps with the work in "deriving bbtm matrices" , "automated sequence alignment", and "ziff module": these four folders are all part of my effort to create and verify the usefulness of alignments for the hmomology modeling that goes into deriving Ezβ assymmetric.

# August 29, 2012

Downloaded TMout.zip from an e-mail David jimenez-Morales sent me on August 23. This is the same TMout folder that is seen in the "deriving bbtm matrices" folder and "automated sequence alignment" folder.