



This movie was generated from snapshots of the 130 ns MD simulation of binding process of a SH3 domain with its binder. Pymol was used to generate the picture with Povray.

Link to AVI version: High quality (30 MB) http://gepard.bioinformatik.uni-saarland.de/old_html/material/dl/SH3_large.avi

Ahmad, M., Gu, W., and Helms, V. (2008), *Angewandte Chemie International Edition*, Vol. 47, p. 7626-7630.
Mechanism of Fast Peptide Recognition by SH3 Domains.