

File name: Supplementary Movie 1

Description: Crystal structure of an MHC I–tapasin– ERp57 chaperone complex at 2.7 Å resolution. Important features of MHC I interaction, peptide proofreading, and loading are displayed. The editor loop of tapasin, which contributes to a widened F-pocket of peptidereceptive MHC I is shown with its electron density (contour level: 1.5 σ).