A figure should be like it's own minipaper, says Doc Castner.

\* Comparison of full and partial model of 1KMO  
- Have the pymol sessions

Illustration of an in-plug - 1UYN

Illustration of PapC perhaps

\* Trimeric porin, showing what strands are at the interface, and what strands contact the constriction loop

\* Three types of Ezβ trend

Ez-β colored surface, with L2 latching loop labeled

Interface colored by hydrophobicity using AAIndex script from PyMOL wiki?

For the full/partial model - best would be a before/after thing from running a sequence through a structure prediction algortihm - I can sapre a half hour to see if I can get that. (until 3:36)  
none of the three stru prediction methods have predicted structures in their supplementary info, except for a protein without inplugs.