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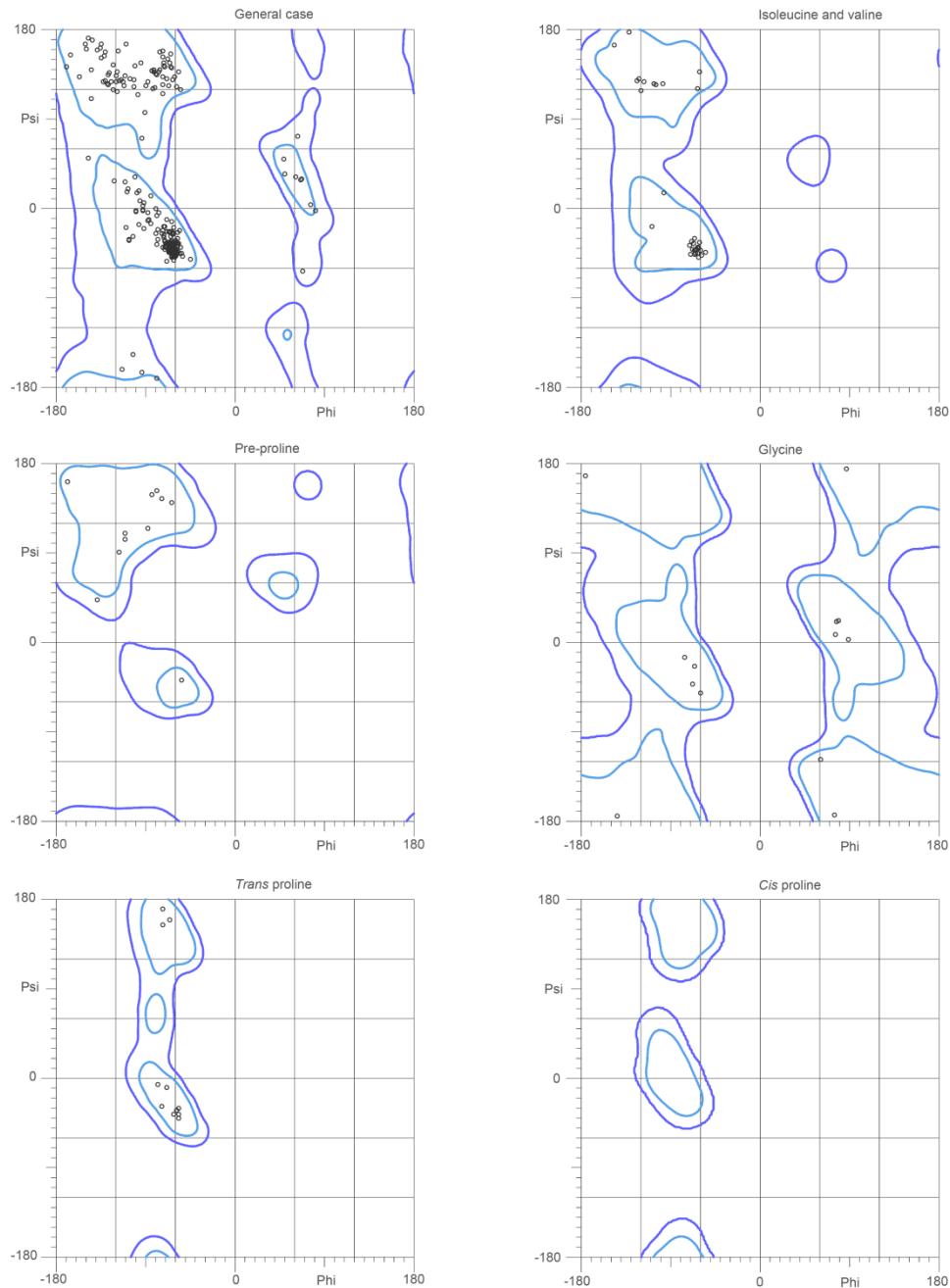
**Supporting information for article:**

**Tah1 helix-swap dimerization prevents mixed Hsp90 co-chaperone complexes**

**Rhodri M. L. Morgan, Mohinder Pal, S. Mark Roe, Laurence H. Pearl and Chrisostomos Prodromou**

## MolProbity Ramachandran analysis

4CQG, model 1



97.4% (299/307) of all residues were in favored (98%) regions.

100.0% (307/307) of all residues were in allowed (>99.8%) regions.

There were no outliers.

<http://kinemage.biochem.duke.edu>

Lovell, Davis, et al. Proteins 50:437 (2003)

**Figure S1** Ramachandran plot.