Table 4 Proteins used to assess the method.

Codes are the Brookhaven PDB code postfixed with the chain identifier code and domain number in Roman numerals where appropriate. hnf3 is assigned to a structure not in the PDB.

Code	Protein Name	Class	Fold
hnf3 ‡	hepatocyte nuclear factor	$\alpha + \beta$	winged helix-turn-helix DNA binding motif
1mba	myoglobin	α	globin
1plc	plastocyanin	β	greek-key β sandwich
1rcb	interleukin–4	α	up-up-down-down 4–helix bundle
1shaa	*v–src* tyr kinase SH2 domain	$\alpha + \beta$	SH2 domain fold
1ubq	ubiquitin	$\alpha + \beta$	β–grasp
1wsya	α-subunit of trp synthase	α/β	α/β–barrel
2hmqa	Hemerythrin	α	up-down-up-down 4–helix bundle
2pgd_I	6–phosphogluconate dehydrogenase	α/β	Rossmann fold
2trxa	thioredoxin	α/β	thioredoxin
4fgf	basic fibroblast growth factor	β	β–trefoil