Supplemental Figure 1:

Size-exclusion chromatography of purified GARP2 using analytical Superdex-200. Elution profiles of purified GARP2 under normal (—) and denaturing conditions (6M guanidinium HCl)(------).



Supplementary Fig. 1

Supplemental Figure 2:

Sequence analysis of the A1 subunit of the CNG channel and peripherin. *A*, PONDR of A1 subunit of the bovine rod cGMP-gated channel; *B*, PONDR of bovine peripherin. Note the pronounced disordered regions in the N-terminus of the A1 subunit and the C-terminus of peripherin.



Supplementary Fig. 2

Amino acid	GARP1	GARP2	Intrinsically unstructured proteins	Globular proteins ^a
% Disorder promoting amino acids				
Pro (P)	10	17.1	12.07	4.61
Glu (E)	23.9	12.7	14.26	5.98
Lys (K)	4.2	3.3	10.43	6.05
Ser (S)	6.9	6.7	6.91	6.31
Gln (Q)	8.5	8.0	4.46	3.69
% Order promoting amino acids				
Tyr (Y)	0	0	1.42	3.64
Phe (F)	0.5	0.3	1.66	3.95
Cys (C)	0.7	1	0.61	1.64
Ile (I)	2	2.7	3.67	5.43
Asn (N)	0	0.7	2.06	4.66
Leu (L)	6.2	8.7	5.44	8.37
Trp (W)	1.3	2.3	0.32	1.55

Supplemental Table. Amino acid frequencies of bovine GARPs compared with those of ordered and disordered proteins

^aData from Ref. (42)