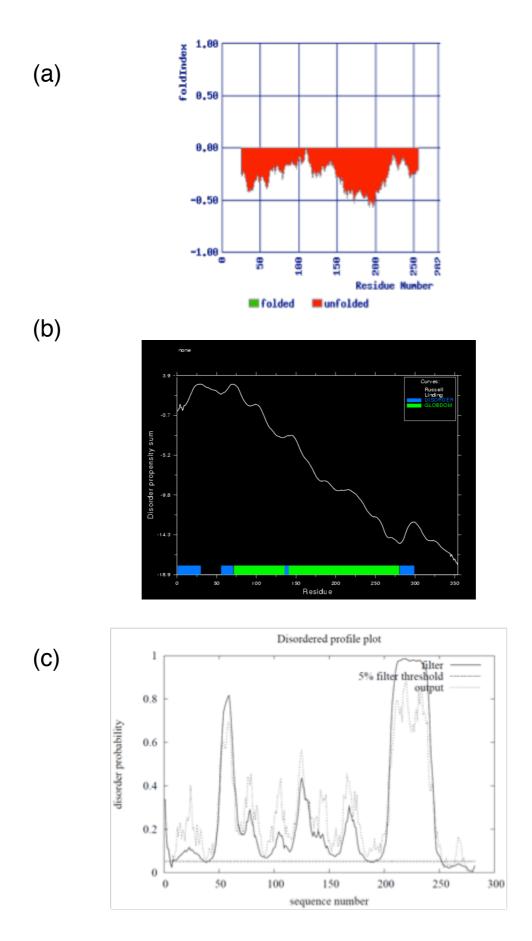
SUPPLEMENTARY INFORMATION

Juxtanodin is an intrinsically disordered F-actin binding protein

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Supplementary figure 1. Additional disorder predictors. (a) FoldIndex predicts no folded regions in juxtanodin. (b) GlobPlot suggests the presence of a globular domain (green) in addition to disordered regions (blue). (c) DisoPred predicts rather short segments of disorder, mainly between residues 50-70, 120-130, 160-180, and 200-250.