



Supplementary Figure S1. Alignment of the sequences in the N-terminal GAR domain of FBL. The FBL sequences in yeast (*S. cerevisiae*), nematodes (*C. elegans*) and fruit fly (*D. melanogaster*) and 5 vertebrate species (*D. rerio*, *X. tropicalis*, *A. carolinensis*, *H. sapiens* and *M. musculus*) are aligned and the ligments of the GAR domains are shown. The alignment was conducted by ClustalW with manual adjustments. The start and end RG(G) of the GAR domains are boxed.