A putative homology of U2AF\textsuperscript{65} in \textit{S.cerevisiae}

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The alignment of yeast YCL11c and human U2AF\textsuperscript{65} in Figure 1 incorrectly showed the RNP-2 submotifs and the limits of each proposed RRM. The correct figure is provided below. The authors regret any confusion this may have caused.

Figure 1. Alignment of YCL11c ORF protein (top line) and hU2AF\textsuperscript{65} (bottom line) using the GCG program Bestfit, with a gap weight 3.5 and gap extension 0.1 (10). One additional gap in the arginine-rich domain was introduced manually. Black boxes represent identical matches; stippled boxes represent conservative changes (V=L=I, F=Y=W, R=K, Q=N, D=E, S=T). RRMs are shown by horizontal lines; RNP-1 and RNP-2 motifs are boxed. The broken line for the C-terminal RRM indicates a weak match to the consensus, particularly for RNP-2 of hU2AF\textsuperscript{65}. 