

Supplemental Figure 2. Sequences of N- and C-terminal phosphorylated sequences in AHA2 used to blast the GDG *Saccharomyces cerevisiae* database. Phosphorylated residues are underlined. No hits were found.

1-MSSLEDIKNET

894-FPEKGSYRELS

926-KGHVESVVKLLK

938-LDIETPSHYTV