

The Structure of the Large Regulatory α -Subunit of Phosphorylase Kinase as
Determined by Hydrogen Deuterium Exchange and Modeling

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Supplementary Materials

Figure S1: Peptide map of deuterated peptides identified and analyzed for the α subunit, resulting in 74% coverage.

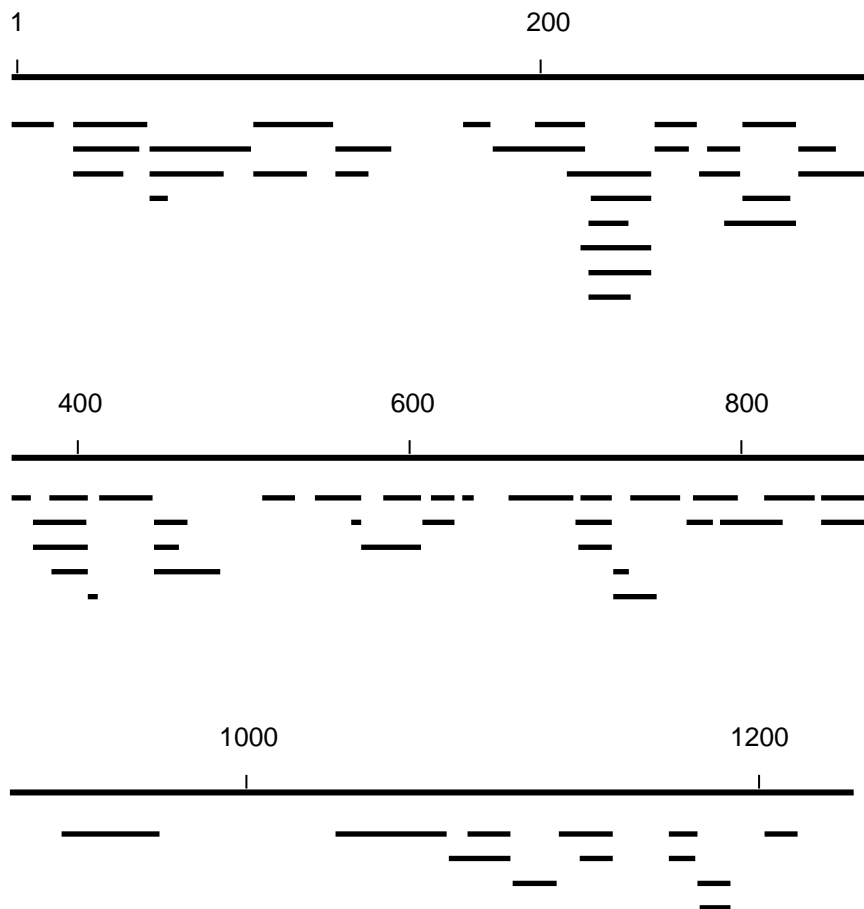


Figure S2: Time courses for deuterium incorporation into all identified peptides. Error bars are the average deviation of independent exchange experiments as described under *Materials and Methods*; error bars are present for all data points, although some are smaller than the symbols.

