

Expression of the Subunits of Protein Kinase CK2 During Oogenesis in *Xenopus laevis*

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Northern blot analysis of RNAs from different tissues demonstrated that the mRNA for the protein kinase CK2 α subunit is very abundant in the ovary of *Xenopus laevis*. The competitive reverse PCR technique has been used to quantitate the mRNA for both CK2 α and CK2 β subunits during oogenesis. The results obtained using eight different animals consistently show an increment of 2-3 fold in the mRNA for both subunits in vitellogenic oocytes (stages II-VI). Each stage-VI oocyte contains approximately 5×10^7 molecules CK2 α mRNA and 1×10^7 molecules CK2 β mRNA. These amounts are considerably higher than many other mRNAs analyzed in these cells. Activity measurements of CK2 using casein or a specific model peptide revealed increments of about 10-12 fold during oogenesis, and also indicated that the amount of enzyme in the nucleus accounted for 15-30% of the total enzyme in the oocyte at all stages. Western blot analysis of CK2 α indicated that the amount of this protein subunit also increased