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Inter-residue hydrogen bonds in the mitochondrial ATP-binding site of the S-locus cysteine-protease SPO11. SPO11 is a key enzyme in meiosis. It is a cysteine protease located in the cytoplasm that cleaves the phosphorylated H2A histone at a specific inter-residue site, the crosslinking domain between adjacent nucleosomes. ATP hydrolysis by SPO11 is essential for the formation of the DSBs that trigger recombination. We have solved the structure of the isolated mitochondrial SPO11 and describe here the location of all the nucleotide-binding sites. Surprisingly, there is a nucleotide in one of the two nucleotide-binding sites, as shown by the presence of a strong electron density peak. Using site-directed mutagenesis, we have shown that the nucleotide-binding site, an essential conserved residue, is part of the catalytic center and that it also interacts with the S-locus cysteine protease SLO1, one of the two substrates of SPO11. The mutation of the two residues involved in the binding of the ATP molecule also disrupts the ATPase activity of the protein.

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