

Nucleotide sequence of a cDNA encoding the ribulose-1,5-bisphosphate carboxylase/oxygenase from sunflower (*Helianthus annuus*)

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Ribulose -1,5- bisphosphate carboxylase/oxygenase (RuBisCO) (E.C. 4.1.1.39) is a bifunctional enzyme which catalyses the carboxylation and the oxygenation of the ribulose -1,5- bisphosphate. This chloroplast enzyme is the most abundant protein in plants (1). The holoenzyme is composed of eight large subunits (LSU) encoded by chloroplast DNA and synthesized in the chloroplast and eight small subunits (SSU) encoded on free cytoplasmic poly-somes, transported to the chloroplast, matured and assembled with the LSUs to form the active enzyme. Nucleotide sequences from 4 SSU cDNAs have been reported (2). Here we present the complete nucleotide sequence of the cDNA encoding the RuBisCO SSU from sunflower. A sunflower cDNA library has been prepared (3) and screened with corresponding cDNA from pea (a gift from N.H. CHUA, Rockefeller University). The longest cDNA was sequenced on each strand by the chain termination method (4) : see figure below.

The nucleotide sequence includes 21 bp of the 5' - untranslated region, 534 bp of the coding region and 116 bp of the 3' - untranslated region. The transit and mature peptides have respectively 55 and 123 amino acids. Comparison with the coding region of pea RuBisCO reveals 68 % homology at the nucleotide level (5) (mature peptide only) and, at the amino acid level, 62 % for the transit peptide and 76 % for the mature peptide.

M	A	S	I	S	S	S	V	A	T	V	S	R	T	A	P	A	Q
CGAAAGACAAAGATTATCGTAATGCCCTCGACTCTCTCAGTCGCACCGTGTAGCCGACGCCCTTGCTCA																	
10	20	30	40	50	60	70											
A	N	H	V	A	P	F	T	G	L	K	S	N	A	A	F	T	T
85	95	105	115	125	135	145											
S	T	L	P	S	N	G	R	D	C	H	K	V	W	P	P	L	G
160	170	180	190	200	210	220											
T	L	S	Y	L	P	L	E	T	R	L	A	K	E	V	D	Y	L
235	245	255	265	275	285	295											
V	P	C	L	E	F	E	L	E	H	G	F	V	Y	R	E	N	A
310	320	330	340	350	360	370											
G	R	Y	W	T	M	H	N	K	L	M	F	G	T	D	S	A	D
385	395	405	415	425	435	445											
E	C	K	K	E	Y	P	D	A	W	I	R	I	G	F	D	N	V
460	470	480	490	500	510	520											
M	F	I	I	A	S	R	P	D	G	Y	***						
535	545	555	565	575	585	595											
CGAATGGTTGATCTTGGGTTCTCGTCATTCCTTCATAATTGGAAATTCTCTTGGATGATGGTCAGTCAATGTATC																	
610	620	630	640	650	660	670											

Figure Legend : Nucleotide sequence of the cDNA and the corresponding amino acid sequence from sunflower RuBisCO SSU.

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