

Table S4. Putative cis-acting elements present in *SPL* promoters of *Codonopsis pilosula*.

	CGTCA-motif	TGACG-motif	ABRE	TCA-element	GARE-motif	ARE	MBS	LTR	GATA-motif	G-Box	GTGGC-motif	AE-box	TCT-motif	chs-CMA2a	CCAAT-box	O2-site	CAT-box	MSA-like
	MeJA-responsive	abscisic acid responsive	salicylic acid responsive	gibberellin-responsive	anaerobic induction	drought-inducibility	low-temperature responsive							MYBHv1 binding site	zein metabolism regulation	related to meristem expression	cell cycle regulation	
<i>CpSPL1</i>	1	0	1	1	2	3	0	0	2	0	0	1	0	0	0	1	2	
<i>CpSPL2</i>	1	0	0	0	0	2	0	0	0	1	0	0	0	0	0	1	0	
<i>CpSPL3</i>	2	2	2	0	0	2	0	0	0	0	0	0	0	0	0	0	0	
<i>CpSPL4</i>	2	2	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	
<i>CpSPL5</i>	4	2	0	0	0	0	0	1	2	0	0	0	0	0	0	0	0	
<i>CpSPL6</i>	1	0	4	0	1	0	2	0	0	6	0	0	0	0	0	0	1	
<i>CpSPL7</i>	3	1	4	0	1	0	0	0	0	1	1	0	0	0	3	0	0	
<i>CpSPL8</i>	2	2	2	0	0	1	0	0	0	0	0	0	0	1	0	0	0	
<i>CpSPL9</i>	0	0	1	0	0	2	0	0	0	0	0	0	0	0	0	1	0	
<i>CpSPL10</i>	2	2	3	0	1	1	4	0	0	0	0	0	0	0	0	0	1	
<i>CpSPL11</i>	1	0	0	0	0	1	1	0	0	0	0	0	0	1	0	0	0	
<i>CpSPL12</i>	1	2	1	0	0	1	0	0	0	0	0	0	1	0	0	1	0	
<i>CpSPL13</i>	0	0	0	0	0	1	1	1	0	1	0	0	0	1	0	0	0	
<i>CpSPL14</i>	6	0	1	0	1	2	7	0	1	0	0	0	0	3	4	1	0	
<i>CpSPL15</i>	1	5	1	0	0	2	0	0	0	0	0	0	0	0	1	0	0	