

Table1 Location and Structure characteristics of GolS and RS members in rubber tree

Protein name	Subcellular localization			The secondary structure composition			
	Cytoplasmic	Chloroplast	Vacuolar	Alpha helix (%)	Extended strand (%)	Beta turn (%)	Random coil (%)
HbGolS1	7.98	1.12	0.51	38.53	11.76	3.53	46.18
HbGolS2	7.98	1.12	0.51	41.67	10.49	2.78	45.06
HbRS1	7.10	1.38	0.77	27.47	21.82	6.68	44.03
HbRS2	7.03	1.41	0.80	27.86	21.25	7.25	43.64