

Table S2. List of proteins identified in the **gel slice b** (18-12 kDa) for each extraction procedures. Statistical parameters, number of unique peptides found for each protein, total spectra number assigned to each protein, protein molecular weight and Citrus Genome Database sequence or Uniprot ID are reported. NCBI description and accession ID obtained through BLAST search (blastp) of the recognized clementine proteins are also shown. The gray lanes specify the putative allergens identified.

Procedure	rank	log(e)	log(I)	% measured	% corrected	unique peptides	spectra n°	MW	Citrus Genome/Uniprot Database ID	NCBI description and accession
1	1	-103,6	5	49	100+	14	28	24,2	clementine0.9_021092m	germin-like protein [Citrus limon] AFN02126.1
	2	-33	4,76	29	42	5	13	17	clementine0.9_024225m	major allergen Pru ar 1 [Vitis vinifera] XP_002274108.2
	3	-23,8	4,66	53	56	5	10	16,3	clementine0.9_024678m	nucleoside diphosphate kinase 1 [Oryza sativa Indica Group] ABR25647.1
	4	-21	4,63	28	53	3	12	19	clementine0.9_023327m	predicted protein [Populus trichocarpa] >gb ABK93810.1 XP_002307094.1
	5	-20,2	4,41	9	10	3	7	47,8	clementine0.9_010237m	2-phospho-D-glycerate hydrolase [Citrus trifoliata] ADD12953.1
	6	-18	4,08	15	48	3	6	16,3	clementine0.9_024571m	putative ribosomal protein S14 [Wolffia arrhiza] ADV38313.1
	7	-14	4	23	40	2	3	19	clementine0.9_023429m	conserved hypothetical protein [Ricinus communis] >gb EEF39253.1 XP_002523068.1
	8	-12,8	4	17	21	3	4	16,8	tr Q9MBF3 Q9MBF3_CITUN Glycine-rich RNA-binding protein	
	9	-11,2	4,49	18	26	2	7	17,8	clementine0.9_024170m	heat shock protein [Citrus unshiu] BAK61831.1
2	1	-46,9	4,57	41	93	8	11	24,2	clementine0.9_021092m	germin-like protein [Citrus limon] AFN02126.1
	2	-21,5	4,22	25	30	4	8	16,8	tr Q9MBF3 Q9MBF3_CITUN Glycine-rich RNA-binding protein	

3	-18	4	34	36	4	7	16,3	clementine0.9_024678m	nucleoside diphosphate kinase 1 [Oryza sativa Indica Group] ABR25647.1
4	-16,2	4	25	33	3	4	16	clementine0.9_024848m	predicted protein [Populus trichocarpa] >gb EEE85908.1 XP_002305397.1
5	-12,7	4	21	40	2	6	19	clementine0.9_023327m	predicted protein [Populus trichocarpa] >gb ABK93810.1 >gb EEE94090.1 XP_002307094.1
6	-11,9	4	6,3	7	2	5	47,8	clementine0.9_010237m	2-phospho-D-glycerate hydrolase [Citrus trifoliata] ADD12953.1
7	-8,1	3,65	14	46	2	3	16,3	clementine0.9_024571m	putative ribosomal protein S14 [Wolffia arrhiza] ADV38313.1
8	-7	4,17	4,7	7	2	4	56,1	clementine0.9_007715m	Aspartic proteinase precursor, putative [Ricinus communis] >gb EEF32480.1 XP_002529926.1

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1	-29,9	4,68	12	17	4	16	39	clementine0.9_014258m	predicted protein [Populus trichocarpa] >gb EEE80730.1 XP_002301457.1
2	-26	4,33	28	30	5	6	20,3	clementine0.9_023186m	ribulose-1,5-bisphosphate carboxylase/oxygenase small subunit [Coffea arabica] CAD11990.1
3	-15,6	4	27	61	3	6	24,2	clementine0.9_021092m	germin-like protein [Citrus limon] AFN02126.1
4	-12,4	4,3	23	25	3	5	16,3	clementine0.9_024678m	nucleoside diphosphate kinase 1 [Oryza sativa Indica Group] ABR25647.1
5	-11,5	3,96	6,3	7	2	4	47,8	clementine0.9_010237m	2-phospho-D-glycerate hydrolase [Citrus trifoliata] ADD12953.1
6	-9,6	4,12	23	38	2	4	14,1	clementine0.9_025303m	Profilin-3 Q9M7N0.1

	7	-9	3,98	24	31	2	5	14,8	clementine0.9_025332m	PREDICTED: 40S ribosomal protein S15a-1-like [Glycine max] >ref XP_003529602.1 NP_001238465.1
4	1	-26,8	4,18	33	75	5	5	24,2	clementine0.9_021092m	germin-like protein [Citrus limon] AFN02126.1
	2	-11,6	4,01	21	40	2	4	19	clementine0.9_023327m	predicted protein [Populus trichocarpa] >gb ABK93810.1 >gb EEE94090.1 XP_002307094.1
	3	-9,9	4,25	25	26	3	6	16,3	clementine0.9_024678m	nucleoside diphosphate kinase 1 [Oryza sativa Indica Group] ABR25647.1