

夏不为利，Agriseraphotocynthesis antibody product line, big promotion!

产品货号	产品名称	价格	促销价
AS01 001	ClpC Chloroplastic form of HSP100	5712	3713
AS01 002	Lhcb3 LHCII type III chlorophyll a/b-binding protein	5712	3713
AS01 002PRE	Lhcb3 LHCII type III chlorophyll a/b-binding protein, pre-immune serum	1881.6	1223
AS01 003	Lhcb2 LHCII type II chlorophyll a/b-binding protein	5712	3713
AS01 004	Lhcb1 LHCII type I chlorophyll a/b-binding protein	4670.4	3036
AS01 004PRE	Lhcb1 LHCII type I chlorophyll a/b-binding protein, pre-immune serum	1965.6	1278
AS01 005	Lhca1 PSI type I chlorophyll a/b-binding protein	5745.6	3735
AS01 006	Lhca2 PSI type II chlorophyll a/b-binding protein	5712	3713
AS01 007	Lhca3 PSI type III chlorophyll a/b-binding protein	5712	3713
AS01 008	Lhca4 PSI type IV chlorophyll a/b-binding protein	5712	3713
AS01 009	Lhcb5 CP26 chlorophyll a/b-binding protein of plant PSII	5712	3713
AS01 009PRE	Lhcb5 CP26 chlorophyll a/b-binding protein of plant PSII, pre-immune serum	1881.6	1223
AS01 010	Lhcb6 CP24 chlorophyll a/b-binding protein of plant PSII	4670.4	3036
AS01 011	2 Set of 10 plant anti-Lhca and anti-Lhcb antibodies	30004.8	19503
AS01 011 CHLAMYDOMONAS	2 Set of 4 Chlamydomonas anti-Lhc antibodies	20244	13159
AS01 016	PsbA D1 protein of PSII, C-terminal (chicken)	6182.4	4019
AS01 016S	PsbA D1 positive control/quantitation standard	2990.4	1944
AS01 017	RbcL Rubisco large subunit, form I	6098.4	3964
AS01 017S	RbcL Rubisco positive control/quantitation standard	2990.4	1944
AS03 030	AtpB Beta subunit of ATP synthase (chloroplastic + mitochondrial) (chicken)	6098.4	3964
AS03 030S	AtpB Positive control/quantitation standard	2990.4	1944
AS03 032	PsbS 22 kDa Lhc-like PSII protein (chicken)	6098.4	3964

AS03 035A	SPS Sucrose phosphate synthase, global	6552	4259
AS03 035-ALP	SPS Sucrose phosphate synthase, global, ALP-conjugated (40 µg)	7173.6	4663
AS03 035-HRP	SPS Sucrose phosphate synthase, global, HRP-conjugated (40 µg)	7173.6	4663
AS03 036	Elip2 Early light inducible protein 2	6098.4	3964
AS03 037	RbcL Rubisco large subunit, form I and form II	4552.8	2959
AS03 037A	RbcL Rubisco large subunit, form I and form II (affinity purified)	4670.4	3036
AS03 037-ALP	RbcL Rubisco large subunit, form I and form II, ALP-conjugated (40 µg)	7173.6	4663
AS03 037B	RbcL Rubisco large subunit, form I and form II, Biotin conjugated (40 µg)	7173.6	4663
AS03 037-HRP	RbcL Rubisco large subunit, form I and form II, HRP-conjugated (40 µg)	7173.6	4663
AS03 037PRE	Rubisco large subunit, pre-immune serum	1965.6	1278
AS04 038	PsbB CP47 protein of PSII	6182.4	4019
AS04 038PRE	PsbB CP47 protein of PSII, pre-immune serum	1965.6	1278
AS04 042P	PsaC PSI-C core subunit of photosystem I (PsaC antibody + PsaC protein positive control)	6182.4	4019
AS04 042S	PsaC positive control/quantitation standard	4216.8	2741
AS04 043	cFBPase Cytosolic fructose-1,6-bisphosphatase (cytoplasm marker in photosynthetic tissues)	4552.8	2959
AS04 045	Lhcb4 CP29 chlorophyll a/b binding protein of plant PSII	5745.6	3735
AS04 045PRE	Lhcb4 CP29 chlorophyll a/b binding protein of plant PSII, pre-immune serum	1881.6	1223
AS04 047	PsaE PSI-E subunit of photosystem I (monocot)	5712	3713
AS04 048	PsaG PSI-G subunit of photosystem I	5712	3713
AS04 049	PsaK PSI-K subunit of photosystem I	5712	3713
AS04 050	PsaO PSI-O subunit of photosystem I	5712	3713
AS04 051	2 Photosynthesis Tool Kit - quantitation	15002.4	9752
AS05 059	PsbR 10 kDa protein of PSII	6098.4	3964
AS05 060	PsbW Small subunit W of PSII	6098.4	3964

AS05 066	PsbX small subunit X of PSII	6098.4	3964
AS05 067	POR Protochlorophyllide oxidoreductase	4670.4	3036
AS05 067-10	POR Protochlorophyllide oxidoreductase (10 µl)	2637.6	1714
AS05 070	2 Educational antibody kit - photosynthesis	25326	16462
AS05 071	AtpH ATP synthase subunit c (chloroplastic)	6098.4	3964
AS05 073	CAH3 Carbonic anhydrase	6098.4	3964
AS05 081	CPX2 Coproporphyrinogen III oxidase, isoform 2	6098.4	3964
AS05 082	Lhca5 PSI type V chlorophyll a/b-binding protein	5712	3713
AS05 084	PsbA D1 protein of PSII, C-terminal (rabbit antibody) (thylakoid membrane marker)	4670.4	3036
AS05 084A	PsbA D1 protein of PSII, C-terminal (affinity purified)	4670.4	3036
AS05 084A-ALP	PsbA D1 protein of PSII, C-terminal, ALP-conjugated (40 µg)	7173.6	4663
AS05 084A-HRP	PsbA D1 protein of PSII, C-terminal, HRP-conjugated (40 µg)	6820.8	4434
AS05 085	AtpB Beta subunit of ATP synthase (chloroplastic + mitochondrial) (rabbit)	4670.4	3036
AS05 085-10	AtpB Beta subunit of ATP synthase (chloroplastic + mitochondrial) (rabbit) (10 µl)	2536.8	1649
AS05 085PRE	AtpB Beta subunit of ATP synthase, pre-immune serum	1965.6	1278
AS05 086	UGPase UDP-glucose pyrophosphorylase (cytoplasm marker)	4670.4	3036
AS05 090	Lci5 low carbon dioxide induced protein number 5	6098.4	3964
AS05 092	PsbO 33 kDa of the oxygen evolving complex (OEC) of PSII (anti-peptide)	6098.4	3964
AS05 093	PrxQ Peroxiredoxin, thioredoxin reductase	6098.4	3964
AS05 094A	FtsH6 ATP-dependent zinc metalloprotease FtsH6 (chloroplastic)	6098.4	3964
AS06 104	PsaF PSI-F subunit of photosystem I	6098.4	3964
AS06 105	PsaH PSI-H subunit of photosystem I (plants)	6098.4	3964
AS06 108	PsaL PSI-L subunit of photosystem I	6098.4	3964
AS06 109	PsaN PSI-N subunit of photosystem I	6098.4	3964

AS06 111	CP43' IsiA homolog of plant CP43	6552	4259
AS06 112	PsbE Alfa subunit of Cytochrome b559 of PSII	4670.4	3036
AS06 113	PsbF beta subunit of Cytochrome b559 of PSII	6098.4	3964
AS06 114	PsbY Small subunit Y of PSII	6098.4	3964
AS06 115	PsbZ ycf9 protein of PSII	6098.4	3964
AS06 116	PGL35 Plastoglobulin 35; FIB1a; FBN1a	6552	4259
AS06 117	Lhcb4 CP29 (Lhcb4) homolog, Chlamydomonas	5712	3713
AS06 118	Pre-apoplastocyanin	6098.4	3964
AS06 119	Cyt f Cytochrome f protein (PetA) of thylakoid Cyt b6/f-complex (algal)	6098.4	3964
AS06 121	FDX1 Ferredoxin	4670.4	3036
AS06 122	CRD1 Cyanobacterial homolog of plant CHL27 cyclase	4670.4	3036
AS06 123	CPX1 coproporphyrinogen III oxidase, isoform 1	6098.4	3964
AS06 130	FtsH9 ATP-dependent zinc metalloprotease FtsH9 (chloroplastic)	6098.4	3964
AS06 141	PC Plastocyanin	4670.4	3036
AS06 142-16	PsbQ 16 kDa protein of the oxygen evolving complex (OEC) of PSII	6098.4	3964
AS06 142-23	PsbP 23 kDa protein of the oxygen evolving complex (OEC) of PSII (anti-protein)	6098.4	3964
AS06 142-33	PsbO 33 kDa of the oxygen evolving complex (OEC) of PSII (anti-protein)	4670.4	3036
AS06 143	PsaH PSI-H subunit of photosystem I, Chlamydomonas	6098.4	3964
AS06 144	PsaG PSI-G subunit of photosystem I, Chlamydomonas	6098.4	3964
AS06 145	LTP vesicle-inducing protein in plastids (VIPP1)	6098.4	3964
AS06 146	PsbD D2 protein of PSII	4670.4	3036
AS06 146PRE	PsbD D2 protein of PSII, pre-immune serum	1965.6	1278
AS06 147A	Elip Early light induced protein	4670.4	3036
AS06 157	PsbH Small subunit H of PSII	6098.4	3964

AS06 158	PsbI Small subunit I of PSII	6098.4	3964
AS06 159	NtrC Thioredoxin reductase (TR/TRxR) (algal)	6552	4259
AS06 163	Hcf101 High chlorophyll fluorescence phenotype protein	6098.4	3964
AS06 167	PsbP 23 kDa protein of the oxygen evolving complex (OEC) of PSII (anti-peptide)	6098.4	3964
AS06 171	PsbTn Tn protein of PSII	6098.4	3964
AS06 172	PsaA PSI-A core protein of photosystem I	4670.4	3036
AS06 176	CGE1 Alfa-CGE1, chloroplastic GrpE homolog	6098.4	3964
AS06 185	SPS Sucrose phosphate synthase (maize)	4670.4	3036
AS06 202	Cyt c6 Thylakoid lumen cytochrome c6 protein	6098.4	3964
AS07 218	Rubisco 557 kDa hexadecamer	5712	3713
AS07 223	Fdx6 ferredoxin 6	6098.4	3964
AS07 224	Fdx3 ferredoxin 3	6098.4	3964
AS07 226	HemF Coproporphyrinogen III oxidase	6098.4	3964
AS07 227	HemY protoporphyrinogen oxidase	6098.4	3964
AS07 228	HemH Protoporphyrin ferrochelatase	6098.4	3964
AS07 235	Toc34 chloroplast outer envelope membrane translocon complex protein (soluble domain)	6098.4	3964
AS07 236	Toc33 chloroplast outer envelope membrane translocon complex protein	6098.4	3964
AS07 237	Toc34 chloroplast outer envelope membrane translocon complex protein (denatured domain)	6098.4	3964
AS07 238	Toc34 chloroplast outer envelope membrane translocon complex protein (CTDcc domain)	6098.4	3964
AS07 239	Toc159 Chloroplast outer envelope membrane translocon complex protein	6098.4	3964
AS07 240	Psb29 Thylakoid membrane formation 1, THF1	6098.4	3964
AS07 241	PEPCK PEP carboxykinase	6098.4	3964
AS07 243	NtrC Thioredoxin reductase (TR/TRxR) (higher plants)	6552	4259
AS07 246	Deg1 chloroplastic DegP-type serine protease 1	6098.4	3964

AS07 250	LHL4 LHC-like protein 4	6098.4	3964
AS07 257	PRK ribulose-5-P-kinase Phosphoribulokinase	4670.4	3036
AS07 257-ALP	PRK ribulose-5-P-kinase phosphoribulokinase, ALP-conjugated (40 µg)	7173.6	4663
AS07 257-HRP	PRK ribulose-5-P-kinase Phosphoribulokinase, HRP-conjugated (40 µg)	7173.6	4663
AS07 259	RbcS Rubisco small subunit (SSU)	4670.4	3036
AS07 259A	RbcS Rubisco small subunit (SSU) (affinity purified)	4670.4	3036
AS07 270	DnaK chloroplast stromal chaperone	4670.4	3036
AS07 273	Ycf3 Photosystem I assembly protein ycf3	6098.4	3964
AS07 274	Ycf4 photosystem I assembly protein ycf4	6098.4	3964
AS08 277	APC allophycocyanin alpha and beta	6098.4	3964
AS08 278	R-PC R-phycocyanin	6098.4	3964
AS08 279	b-PE Phycoerythrobilin	6098.4	3964
AS08 280	B-PE phycoerythrobilin and phycourobilin	6098.4	3964
AS08 281	95 kDa Lcm1 (phycobilisome - to- thylakoid core linker with phycocyanobilin chromophore)	6098.4	3964
AS08 282	Lhc1 PS 1 light harvesting holo-complex	5712	3713
AS08 289	ZEP Zeaxanthin Epoxidase (anti-peptide)	4670.4	3036
AS08 292	Toc75 Outer chloroplast membrane translocon complex protein	6098.4	3964
AS08 293	Tic110 chloroplast inner envelope membrane translocon complex	6098.4	3964
AS08 300	1 PEB (4x) protein extraction buffer	1713.6	1114
AS08 304	AtpA Alpha subunit of ATP synthase, chloroplastic	6098.4	3964
AS08 305	PsbP 23 kDa protein of the oxygen evolving complex (OEC) of PSII (anti-monocot protein)	6098.4	3964
AS08 306	Cyt f Cytochrome f protein (PetA) of thylakoid Cyt b6/f-complex (higher plants)	4670.4	3036
AS08 312	AtpC Gamma subunit of ATP synthase (chloroplastic)	6098.4	3964
AS08 316	CURT1A Curvature thylakoid 1A	4670.4	3036

AS08 324A	PsaE PSI-E subunit of photosystem I (affinity purified)	4670.4	3036
AS08 330	PetC Rieske iron-sulfur protein of Cyt b6/f complex	4670.4	3036
AS08 330S	PetC Rieske iron-sulfur protein of Cyt b6/f complex, protein standard	4216.8	2741
AS08 345	Toc75 Protein TOC75-3, chloroplastic, POTRA domain 1	6098.4	3964
AS08 351	Toc75 Protein TOC75-3, chloroplastic, POTRA domain 3	4670.4	3036
AS08 369	Tlp18.3 Thylakoid lumen 18.3 kDa protein	6098.4	3964
AS08 370	ATPase ATP synthase, whole enzyme	6098.4	3964
AS08 371S	HSP70 Positive control/quantitation standard	2990.4	1944
AS08 375	HemC Porphobilinogen-Deaminase (PBG)	6098.4	3964
AS09 146S	PsbD D2 protein of PSII positive control/quantitation standard	4216.8	2741
AS09 312P	AtpC gamma subunit of ATP synthase, blocking peptide	1696.8	1103
AS09 407	Lhcb5 CP26 (Lhcb5) homolog, Chlamydomonas	4670.4	3036
AS09 408	Lhcbm5 Chlorophyll a-b binding protein of LHCI	4670.4	3036
AS09 409	2 Rubisco quantitation kit	10905.3	7088
AS09 457	AtpG ATPsynthase subunit II b' (chloroplastic)	6098.4	3964
AS09 458	PEPC Phosphoenolpyruvate carboxylase	4670.4	3036
AS09 458-ALP	PEPC phosphoenolpyruvate carboxylase, ALP-conjugated (40 µg)	7173.6	4663
AS09 458-HRP	PEPC phosphoenolpyruvate carboxylase, HRP-conjugated (40 µg)	7173.6	4663
AS09 458S	PEPC Phosphoenolpyruvate carboxylase positive control/quantitation standard	2990.4	1944
AS09 459	ClpB-P ClpB3	5896.8	3833
AS09 461	PsaD PSI-D subunit of photosystem I	6098.4	3964
AS09 512	Lhcb4 CP29 (Lhcb4) homolog (Ostreococcus tauri)	5712	3713
AS09 513	Lhcb5 CP26 (Lhcb5) homolog (Ostreococcus tauri)	5712	3713
AS09 522	Lhcb1 LHCI type I chlorophyll a/b-binding protein (Arabidopsis specific)	5712	3713

AS09 533	PsbS 22 kDa Lhc-like PSII protein (rabbit antibody)	4670.4	3036
AS09 591	AtpH ATP synthase subunit c (chloroplastic)	6098.4	3964
AS10 111S	CP43' IsiA homolog of plant CP43 positive control/quantitation standard	4216.8	2741
AS10 1581	Deg2 Chloroplastic DegP-type serine protease 2	7182	4668
AS10 1583	AtpI CFoIV subunit of ATP synthase	7182	4668
AS10 1584	TAK1 serine/threonine protein kinase (300 ul)	17028.9	11069
AS10 1586	AtpE Epsilon subunit of ATP synthase	7182	4668
AS10 1588	HtrA putative serine protease (300 ul)	17028.9	11069
AS10 1589	PPIase CYP37 peptidyl-prolyl cis-trans isomerase CYP37 (chloroplastic)	7182	4668
AS10 1590	AtpD CF1 delta subunit of ATP synthase (affinity purified)	7182	4668
AS10 1591	AtpD CF1 delta subunit of ATP synthase	7182	4668
AS10 1594	sII0408 Putative thylakoid lumen PPIase	7182	4668
AS10 1595	PPIase CYP37 peptidyl-prolyl cis-trans isomerase CYP37 (chloroplastic) (300 ul)	17028.9	11069
AS10 1598	SppA1 signal peptide peptidase (higher plants) (300 ul)	17028.9	11069
AS10 1599	Hik29 sensor histidine kinase (300 ul)	17028.9	11069
AS10 1600	Hik29 sensor histidine kinase	7182	4668
AS10 1601	STN8 Serine/threonine-protein kinase STN8 (chloroplastic)	4670.4	3036
AS10 1602	HtrA Putative serine protease	7182	4668
AS10 1603	HliA high light inducible protein (300 ul)	17028.9	11069
AS10 1604	AtpF CFoI subunit of ATP synthase	7182	4668
AS10 1605	SppA2 signal peptide peptidase (cyanobacterial) (300 ul)	17028.9	11069
AS10 1607	CYP38 Cyclophilin 38, peptidyl-prolyl cis-trans isomerase	7182	4668
AS10 1608	HhoA protease HhoA (SII1427; Cyanobase) (300 ul)	17028.9	11069
AS10 1609	sII0408 putative thylakoid lumen PPIase (300 ul)	17028.9	11069

AS10 1610	HliD High light inducible protein	7182	4668
AS10 1612	TAK1 Serine/threonine protein kinase TAK1	7182	4668
AS10 1613	SppA1 Signal peptide peptidase (higher plants)	7182	4668
AS10 1614	SppA1 signal peptide peptidase (cyanobacterial) (200 ul)	7182	4668
AS10 1615	HliD High light inducible protein (300 ul)	17028.9	11069
AS10 1616	HliA High light inducible protein	7182	4668
AS10 1617	HhoB protease HhoB (Sll1679; Cyanobase) (300 ul)	17028.9	11069
AS10 1618	CYP38 cyclophilin 38, peptidyl-prolyl cis-trans isomerase (300 ul)	17028.9	11069
AS10 1619	HhoA protease HhoA (Sll1427; Cyanobase) (100 ul)	7182	4668
AS10 1620	HhoB Protease HhoB (Sll1679; Cyanobase)	7182	4668
AS10 1624	SppA2 Signal peptide peptidase (cyanobacterial)	7182	4668
AS10 682	PsbP-like protein (sll1418) cyanobacterial	6098.4	3964
AS10 689	GluTR Glutamyl -tRNA reductase	6098.4	3964
AS10 692	PsbI Small subunit I of PSII (cyanobacterial)	6098.4	3964
AS10 693	HemH Protoporphyrin ferrochelatase (cyanobacterial)	6098.4	3964
AS10 695	PsaB PSI-B core subunit of photosystem I	4670.4	3036
AS10 700	RA Rubisco activase	4670.4	3036
AS10 703	Deg5 protease Do-like 5 (chloroplastic)	6098.4	3964
AS10 704	PsbA D1 protein of PSII, DE-loop	4670.4	3036
AS10 709	Tic40 Inner envelope membrane translocon complex protein (chloroplast)	4670.4	3036
AS10 725	PGRL1 PGR5-like protein 1A (chloroplastic)	6098.4	3964
AS10 936	CGL78 YCF54	6098.4	3964
AS10 939	PsaC PSI-C core subunit of photosystem I	4670.4	3036
AS11 1756	DegP7 protease Do-like 7	6098.4	3964

AS11 1757	Fd Ferredoxin 1 (chloroplastic)	4670.4	3036
AS11 1783	ACD1 Accelerated cell death 1	4670.4	3036
AS11 1786	PsbA D1 protein of PSII, N-terminal	4670.4	3036
AS11 1787	PsbC CP43 protein of PSII	4670.4	3036
AS12 1852	PSB33 Rieske (2Fe-2S) domain-containing protein	4670.4	3036
AS12 1853	CGL160 Conserved in green lineage 160	4670.4	3036
AS12 2117	VDR1 Violaxanthin de-epoxidase-related protein	4670.4	3036
AS12 2118	SNE18 Rossmann-fold NAD(P)-binding domain-containing protein	4670.4	3036
AS12 2613	CPN60A1 Chaperonin 60 subunit alpha 1 (chloroplastic)	4670.4	3036
AS12 2634	DRP5B Dynamin related protein 5B	4670.4	3036
AS13 2641	CGL40	4670.4	3036
AS13 2642	CNfu1 NifU-like protein 1 (chloroplastic)	4670.4	3036
AS13 2654	PSA2 Photosystem I assembly factor 2	4670.4	3036
AS13 2655	ClpP6 chloroplast ClpP6 proteolytic subunit	4670.4	3036
AS13 2669	PsbA D1 protein of PSII, phosphorylated	4670.4	3036
AS13 2676	ARC5 Dynamin-like protein ARC5	6066.9	3943
AS13 2704	Lhcb1-P LHCII type I chlorophyll a/b-binding protein, phosphorylated	5712	3713
AS13 2705	Lhcb2-P LHCII type II chlorophyll a/b-binding protein, phosphorylated	5712	3713
AS13 2709	LYC Lycopene beta cyclase (chloroplastic)	4670.4	3036
AS13 2710	IPP isomerase Isopentyl pyrophosphate isomerase	4670.4	3036
AS13 2711	NdhF NAD(P)H dehydrogenase subunit 5 (chloroplastic)	3685.5	2396
AS13 2712	NdhH NAD(P)H-quinone oxidoreductase subunit H (chloroplastic)	3685.5	2396
AS13 2720	MatK Maturase K	3685.5	2396
AS13 2725	SVR4 Suppressor of variegation 4	4670.4	3036

AS13 2726	SVR4-like Suppressor of variegation 4 - like	4670.4	3036
AS13 2727	GAPC2 Glyceraldehyde-3-phosphate dehydrogenase	5325.6	3462
AS13 2729	RAF2 Rubisco accumulation factor 2	4670.4	3036
AS14 2760	CsoS1A/B/C Major carboxysome shell protein 1A, AB, 1C	4670.4	3036
AS14 2766	LHCSR3	4670.4	3036
AS14 2767	Deg8 protease Do-like 8 (chloroplastic)	4670.4	3036
AS14 2782	EPSP synthase 3-phosphosnikimate 1-carboxyvinyltransferase (chloroplastic, Arabidopsis thaliana)	4670.4	3036
AS14 2786	PsbN Potosystem II reaction center protein N	4670.4	3036
AS14 2793	G4 Chlorophyll synthase (chloroplastic)	4670.4	3036
AS14 2808	Trxf1/2 Thioredoxin F1/F2 (chloroplastic)	4670.4	3036
AS14 2809	TrxM1/M2 Thioredoxin M1/M2 (chloroplastic)	4670.4	3036
AS14 2813	UGPase UDP-glucose pyrophosphorylase (cytoplasm marker) (Hordeum vulgare)	4670.4	3036
AS14 2819	LHCSR1	4670.4	3036
AS14 2820	RBP40 38 kDa RNA-binding protein	4670.4	3036
AS14 2824	PsbO1 33 kDa of the oxygen evolving complex (OEC) of PSII	4670.4	3036
AS14 2825	PsbO2 33 kDa of the oxygen evolving complex (OEC) of PSII	4670.4	3036
AS14 2829	UAGPase UDP-GlcNAc pyrophosphorylase	4670.4	3036
AS15 2830	SUS1 Sucrose synthase 1	4670.4	3036
AS15 2833	PAB protein in chloroplast atpase biogenesis	4670.4	3036
AS15 2834	At2g21960	4670.4	3036
AS15 2835	RIQ1 Reduced induction of non-photochemical quenching 1	4670.4	3036
AS15 2872	PSA3 Photosystem I Assembly 3	4670.4	3036
AS15 2873	SBPase Sedoheptulose-1,7-bis phosphatase	4670.4	3036
AS15 2873S	SBP Sedoheptulose-1,7-bis phosphatase positive control/quantitation standard	2990.4	1944

AS15 2894	GAPC1/2 Glyceraldehyde-3-phosphate dehydrogenase	4670.4	3036
AS15 2895	BAM1 Beta amylase 1 (chloroplastic)	4670.4	3036
AS15 2903	TKL1 transketolase (chloroplastic)	4670.4	3036
AS15 2909	FNR Ferredoxin-NADP+-oxidoreductase	4670.4	3036
AS15 2910	PntA (Slr1239) Pyridine nucleotide transhydrogenase alpha-subunit	4670.4	3036
AS15 2955	RbcL II Rubisco large subunit, form II	4670.4	3036
AS15 2955S	RbcL II Rubisco form II positive control/quantitation standard	2990.4	1944
AS15 2994	Rubisco ELISA quantitation kit	16764.3	10897
AS15 3077	pClpP Chloroplastic ATP-dependent Clp protease proteolytic subunit 1	6974.1	4533
AS15 3079	LCY Lycopene beta-cyclase (chloroplastic)	4670.4	3036
AS15 3080	STT7 Serine/threonine-protein kinase STT7 (chloroplastic)	4670.4	3036
AS15 3081	LhcSR1 (Physcomitrella patens)	4670.4	3036
AS15 3083	ARSA targeting of tail-anchored proteins	4670.4	3036
AS15 3084	LUT1 Beta-carotene hydroxylase	4670.4	3036
AS15 3085	LUT5 beta-carotene hydroxylase	4670.4	3036
AS15 3088	LHCb9 Light-harvesting complex	4670.4	3036
AS15 3089	TAB3 Nucleus-encoded factor involved in PsaB translation	4670.4	3036
AS15 3090	GUN4 Regulatory subunit of Mg-chelatase	4670.4	3036
AS15 3091	VDE Violaxanthin de-epoxidase	4670.4	3036
AS15 3092	ZEP Zeaxanthin Epoxidase (anti-protein)	4670.4	3036
AS16 3114	Fdx2 Ferredoxin 2	4670.4	3036
AS16 3119	EPSP synthase 5-phosphosuccinate 1-carboxyvinyltransferase (chloroplastic, higher plants)	4670.4	3036
AS16 3211	cpCK2 Casein kinase II subunit alpha (chloroplastic)	2923.2	1900
AS16 3692	PTOX Plastid terminal oxidase	5745.6	3735

AS16 3929	FtsH2 + FtsH8 ATP-dependent zinc metalloprotease FtsH2 + FtsH8 (chloroplasic)	4670.4	3036
AS16 3930	FtsH1 + FtsH5 ATP-dependent zinc metalloprotease FtsH1 + FtsH5 (chloroplasic)	4670.4	3036
AS16 3931	NDC1 Alternative NAD(P)H-ubiquinone oxidoreductase C1 (chloroplasic/mitochondrial)	4670.4	3036
AS16 3976	AtpB Beta subunit of ATP synthase, mitochondrial	4670.4	3036
AS16 3985	PGR5 Proton gradient regulation 5	4670.4	3036
AS16 4039	ClpB1 Chaperone protein ClpB1 (cyanobacterial)	4670.4	3036
AS16 4040	ClpB2 Chaperone protein ClpB2 (cyanobacterial)	4670.4	3036
AS16 4041	ClpP1 ATP-dependent Clp protease proteolytic subunit 1 (cyanobacterial)	4670.4	3036
AS16 4042	ClpP2 ATP-dependent Clp protease proteolytic subunit 2 (cyanobacterial)	4670.4	3036
AS16 4043	ClpP3 ATP-dependent Clp protease proteolytic subunit 3 (cyanobacterial)	4670.4	3036
AS16 4048	ClpD Chaperone protein ClpD (chloroplasic)	4670.4	3036
AS16 4049	ClpP1 ATP-dependent Clp protease proteolytic subunit 1 (chloroplasic)	4670.4	3036
AS16 4050	ClpP3 ATP-dependent Clp protease proteolytic subunit 3 (chloroplasic)	4670.4	3036
AS16 4051	ClpP4 ATP-dependent Clp protease proteolytic subunit 4 (chloroplasic)	4670.4	3036
AS16 4052	ClpP5 ATP-dependent Clp protease proteolytic subunit 5 (chloroplasic)	4670.4	3036
AS16 4053	ClpP6 ATP-dependent Clp protease proteolytic subunit 6 (chloroplasic)	4670.4	3036
AS16 4054	ClpK1 ATP-dependent Clp protease proteolytic subunit-related protein 1 (chloroplasic)	4670.4	3036
AS16 4055	ClpK2 ATP-dependent Clp protease proteolytic subunit-related protein 2 (chloroplasic)	4670.4	3036
AS16 4056	ClpK3 ATP-dependent Clp protease proteolytic subunit-related protein 3 (chloroplasic)	4670.4	3036
AS16 4057	ClpK4 ATP-dependent Clp protease proteolytic subunit-related protein 4 (chloroplasic)	4670.4	3036
AS16 4058	ClpT1 ATP-dependent Clp protease ATP-binding subunit CLPT1 (chloroplasic)	4670.4	3036
AS16 4064	NdhB NAD(P)H-quinone oxidoreductase subunit 2 (chloroplasic)	5254.2	3415
AS16 4065	NdhH NAD(P)H-quinone oxidoreductase subunit H (chloroplasic)	5254.2	3415
AS16 4066	NdhS NAD(P)H-quinone oxidoreductase subunit S (chloroplasic)	5254.2	3415

AS16 4082	RPL33 50S ribosomal protein L33 (chloroplastic)	4670.4	3036
AS16 4084	PPH1/TAP38 Protein phosphatase 1	4670.4	3036
AS16 4098	STN7 Serine/threonine-protein kinase STN7 (chloroplastic)	4670.4	3036
AS17 4115	Lhcx6 Lhcx subclade of fucoxanthin Chl a/c proteins	4670.4	3036
AS17 4116	FCP Fucoxanthin Chla/c protein	4670.4	3036
AS17 4117	Nucleomorph HLIP Nucleomorph high-light induced protein	4670.4	3036
AS17 4146	ClpS ATP-dependent Clp protease adapter protein ClpS	4670.4	3036
AS18 4169	Cyt b6 / PetB Thylakoid membrane cytochrome b6 protein, N terminal	4670.4	3036
AS19 4257	TROL Thylakoid rhodanese-like protein	5040	3276
AS19 4287	CURT1C Curvature thylakoid 1C	4670.4	3036
AS19 4293	PLGG1 Plastidal glycolate/glycerate translocator 1, (chloroplastic)	4670.4	3036
AS19 4297	NdbA Thylakoid Localized Type 2 NAD(P)H Dehydrogenase	4670.4	3036