

Table S1

Number	Accession	Annotation	Organism	Taxonomy
1	62738704	Chain A, Crystal Structure Of Potato Tuber Adp-Glucose Pyrophosphorylase	<i>Solanum tuberosum</i>	dicot
2	27819107	ADP-glucose pyrophosphorylase small subunit	<i>Solanum tuberosum</i>	dicot
3	77416911	ADP-glucose pyrophosphorylase small subunit-like	<i>Solanum tuberosum</i>	dicot
4	556351	ADP-glucose pyrophosphorylase small subunit	<i>Solanum tuberosum</i>	dicot
5	15238933	glucose-1-phosphate adenyltransferase small subunit	<i>Arabidopsis thaliana</i>	dicot
6	297791999	hypothetical protein ARALYDRAFT_494882 agps	<i>Arabidopsis lyrata</i> subsp. <i>lyrata</i>	dicot
7	357495273	Glucose-1-phosphate adenyltransferase	<i>Medicago truncatula</i>	dicot
8	357462397	Glucose-1-phosphate adenyltransferase	<i>Medicago truncatula</i>	dicot
9	356552274	PREDICTED: glucose-1-phosphate adenyltransferase small subunit, isoform 1	<i>Glycine max</i>	dicot
10	356501687	PREDICTED: glucose-1-phosphate adenyltransferase small subunit, isoform 2	<i>Glycine max</i>	dicot
11	224131934	predicted protein	<i>Populus trichocarpa</i>	dicot
12	255567204	glucose-1-phosphate adenyltransferase, putative	<i>Ricinus communis</i>	dicot
13	225447450	PREDICTED: hypothetical protein	<i>Vitis vinifera</i>	dicot
14	13487709	ADP-glucose pyrophosphorylase small subunit	<i>Brassica rapa</i> subsp. <i>pekinensis</i>	dicot
15	556622	ADP-glucose pyrophosphorylase	<i>Beta vulgaris</i> subsp. <i>vulgaris</i>	dicot
16	1237080	ADP-glucose pyrophosphorylase	<i>Pisum sativum</i>	dicot
17	1237082	ADP-glucose pyrophosphorylase	<i>Pisum sativum</i>	dicot
18	13487787	ADP-glucose pyrophosphorylase small subunit CagpS1	<i>Cicer arietinum</i>	dicot
19	16950559	ADP-glucose pyrophosphorylase small subunit CagpS2	<i>Cicer arietinum</i>	dicot
20	2642636	ADP-glucose pyrophosphorylase small subunit	<i>Citrullus lanatus</i> subsp. <i>vulgaris</i>	dicot
21	2625084	ADP-glucose pyrophosphorylase small subunit	<i>Cucumis melo</i>	dicot
22	5917789	ADP-glucose pyrophosphorylase small subunit	<i>Citrus unshiu</i>	dicot
23	111660950	ADP-glucose pyrophosphorylase small subunit	<i>Citrus sinensis</i>	dicot
24	41350641	ADP-glucose pyrophosphorylase small subunit	<i>Fragaria x ananassa</i>	dicot

25	45505207	ADP-glucose pyrophosphorylase small subunit	<i>Ipomoea batatas</i>	dicot
26	45505205	ADP-glucose pyrophosphorylase small subunit	<i>Ipomoea batatas</i>	dicot
27	7671232	ADP-glucose pyrophosphorylase	<i>Perilla frutescens</i>	dicot
28	7671230	ADP-glucose pyrophosphorylase catalytic subunit	<i>Perilla frutescens</i>	dicot
29	29421116	ADP-glucose pyrophosphorylase small subunit PvAGPS1	<i>Phaseolus vulgaris</i>	dicot
30	440595	ADP-glucose pyrophosphorylase	<i>Vicia faba</i> var. minor	dicot
31	440593	ADP-glucose pyrophosphorylase	<i>Vicia faba</i> var. minor	dicot
32	162462257	ADP-glucose pyrophosphorylase small subunit	<i>Zea mays</i>	monocot
33	162461970	LOC541902	<i>Zea mays</i>	monocot
34	73747074	ADP-glucose pyrophosphorylase small subunit	<i>Zea mays</i>	monocot
35	23664353	Brittle 2	<i>Zea mays</i> subsp. <i>mays</i>	monocot
36	14582768	ADP-glucose pyrophosphorylase small subunit	<i>Zea mays</i>	monocot
37	115476014	Os08g0345800	<i>Oryza sativa</i> Japonica Group	monocot
38	115478426	Os09g0298200	<i>Oryza sativa</i> Japonica Group	monocot
39	125987830	GLGS_ORYSJ	<i>Oryza sativa</i> Japonica Group	monocot
40	357157910	PREDICTED: glucose-1-phosphate adenylyltransferase small subunit	<i>Brachypodium distachyon</i>	monocot
41	357145854	PREDICTED: glucose-1-phosphate adenylyltransferase small subunit, isoform 2	<i>Brachypodium distachyon</i>	monocot
42	357145851	PREDICTED: glucose-1-phosphate adenylyltransferase small subunit, isoform 1	<i>Brachypodium distachyon</i>	monocot
43	242048788	hypothetical protein SORBIDRAFT_02g020410	<i>Sorghum bicolor</i>	monocot
44	27464770	ADP-glucose pyrophosphorylase small subunit	<i>Hordeum vulgare</i> subsp. <i>vulgare</i>	monocot
45	51556842	ADP-glucose pyrophosphorylase small subunit a	<i>Hordeum vulgare</i>	monocot
46	1707940	GLGS_HORVU	<i>Hordeum vulgare</i>	monocot
47	52430025	ADP-glucose pyrophosphorylase small subunit	<i>Triticum aestivum</i>	monocot
48	21687	ADP-glucose pyrophosphorylase	<i>Triticum aestivum</i>	monocot
49	232166	GLGL1_SOLTU	<i>Solanum tuberosum</i>	dicot
50	1707929	GLGL2_SOLTU	<i>Solanum tuberosum</i>	dicot
51	1707932	GLGL3_SOLTU	<i>Solanum tuberosum</i>	dicot

52	14916987	GLGL1_ARATH	<i>Arabidopsis thaliana</i>	dicot
53	12644324	GLGL2_ARATH	<i>Arabidopsis thaliana</i>	dicot
54	17433716	GLGL3_ARATH	<i>Arabidopsis thaliana</i>	dicot
55	11386853	GLGL4_ARATH	<i>Arabidopsis thaliana</i>	dicot
56	1947084	ADP-glucose pyrophosphorylase large subunit agpl1	<i>Solanum lycopersicum</i>	dicot
57	1840114	ADP-glucose pyrophosphorylase large subunit agpl2	<i>Solanum lycopersicum</i>	dicot
58	1840116	ADP-glucose pyrophosphorylase large subunit agpl3	<i>Solanum lycopersicum</i>	dicot
59	1778434	ADP-glucose pyrophosphorylase large subunit agp-s1	<i>Solanum lycopersicum</i>	dicot
60	1778436	ADP-glucose pyrophosphorylase large subunit agp-s2	<i>Solanum lycopersicum</i>	dicot
61	297812109	hypothetical protein ARALYDRAFT_488807 agpl1	<i>Arabidopsis lyrata</i> subsp. <i>lyrata</i>	dicot
62	297845724	hypothetical protein ARALYDRAFT_472972 agpl2	<i>Arabidopsis lyrata</i> subsp. <i>lyrata</i>	dicot
63	297797902	hypothetical protein ARALYDRAFT_490693 agpl3	<i>Arabidopsis lyrata</i> subsp. <i>lyrata</i>	dicot
64	297821353	predicted protein agpl4	<i>Arabidopsis lyrata</i> subsp. <i>lyrata</i>	dicot
65	357467317	Glucose-1-phosphate adenylyltransferase large subunit	<i>Medicago truncatula</i>	dicot
66	356571037	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 2	<i>Glycine max</i>	dicot
67	356563435	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 1	<i>Glycine max</i>	dicot
68	356562361	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit	<i>Glycine max</i>	dicot
69	356553863	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit	<i>Glycine max</i>	dicot
70	356545193	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 1-like	<i>Glycine max</i>	dicot
71	356538761	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 1-like	<i>Glycine max</i>	dicot
72	356521967	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 1	<i>Glycine max</i>	dicot
73	356518710	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit	<i>Glycine max</i>	dicot
74	356517038	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 1-like	<i>Glycine max</i>	dicot
75	356509672	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit	<i>Glycine max</i>	dicot
76	356508352	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 1-like	<i>Glycine max</i>	dicot
77	356503982	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 2	<i>Glycine max</i>	dicot
78	357511621	Glucose-1-phosphate adenylyltransferase	<i>Medicago truncatula</i>	dicot

79	357480219	Glucose-1-phosphate adenylyltransferase	<i>Medicago truncatula</i>	dicot
80	357473317	Glucose-1-phosphate adenylyltransferase	<i>Medicago truncatula</i>	dicot
81	224128113	predicted protein	<i>Populus trichocarpa</i>	dicot
82	224103389	predicted protein	<i>Populus trichocarpa</i>	dicot
83	224117842	predicted protein	<i>Populus trichocarpa</i>	dicot
84	224100249	predicted protein	<i>Populus trichocarpa</i>	dicot
85	224095317	predicted protein	<i>Populus trichocarpa</i>	dicot
86	255585297	glucose-1-phosphate adenylyltransferase, putative	<i>Ricinus communis</i>	dicot
87	255552303	glucose-1-phosphate adenylyltransferase, putative	<i>Ricinus communis</i>	dicot
88	255548169	glucose-1-phosphate adenylyltransferase, putative	<i>Ricinus communis</i>	dicot
89	255543725	glucose-1-phosphate adenylyltransferase, putative	<i>Ricinus communis</i>	dicot
90	255538708	glucose-1-phosphate adenylyltransferase, putative	<i>Ricinus communis</i>	dicot
91	225458219	PREDICTED: hypothetical protein	<i>Vitis vinifera</i>	dicot
92	225437808	PREDICTED: hypothetical protein	<i>Vitis vinifera</i>	dicot
93	225434249	PREDICTED: hypothetical protein	<i>Vitis vinifera</i>	dicot
94	225432564	PREDICTED: hypothetical protein	<i>Vitis vinifera</i>	dicot
95	225428422	PREDICTED: hypothetical protein	<i>Vitis vinifera</i>	dicot
96	224085694	predicted protein	<i>Populus trichocarpa</i>	dicot
97	224080375	predicted protein	<i>Populus trichocarpa</i>	dicot
98	224062107	predicted protein	<i>Populus trichocarpa</i>	dicot
99	162460455	plastid ADP-glucose pyrophosphorylase large subunit	<i>Zea mays</i>	monocot
100	189027076	glucose-1-phosphate adenylyltransferase large subunit 1	<i>Zea mays</i>	monocot
101	162458350	glucose-1-phosphate adenylyltransferase large subunit 2	<i>Zea mays</i>	monocot
102	162463875	putative glucose-1-phosphate adenylyltransferase large subunit 3 precursor	<i>Zea mays</i>	monocot
103	357132398	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit	<i>Brachypodium distachyon</i>	monocot
104	357119087	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 2	<i>Brachypodium distachyon</i>	monocot
105	357116651	PREDICTED: glucose-1-phosphate adenylyltransferase large subunit 1	<i>Brachypodium distachyon</i>	monocot

106	115455167	Os03g0735000	<i>Oryza sativa</i> Japonica Group	monocot
107	297604962	Os05g0580000	<i>Oryza sativa</i> Japonica Group	monocot
108	115471355	Os07g0243200	<i>Oryza sativa</i> Japonica Group	monocot
109	115438749	Os01g0633100	<i>Oryza sativa</i> Japonica Group	monocot
110	242088961	hypothetical protein SORBIDRAFT_09g029610	<i>Sorghum bicolor</i>	monocot
111	242053733	hypothetical protein SORBIDRAFT_03g028850	<i>Sorghum bicolor</i>	monocot
112	242033053	hypothetical protein SORBIDRAFT_01g008940	<i>Sorghum bicolor</i>	monocot
113	1279513	glucose-1-phosphate adenyltransferase	<i>Hordeum vulgare</i> subsp. vulgare	monocot
114	2105137	ADP-glucose pyrophosphorylase large subunit	<i>Hordeum vulgare</i> subsp. vulgare	monocot
115	445623	ADP glucose pyrophosphorylase : SUBUNIT=L	<i>Hordeum vulgare</i>	monocot
116	22347636	ADP-glucose pyrophosphorylase large subunit	<i>Oncidium Goldiana</i>	monocot
117	32812836	ADP-glucose pyrophosphorylase large subunit	<i>Triticum aestivum</i>	monocot
118	89277026	plastid ADP-glucose pyrophosphorylase large subunit	<i>Triticum aestivum</i>	monocot
119	1707930	GLGL2_WHEAT	<i>Triticum aestivum</i>	monocot
120	121293	GLGL3_WHEAT	<i>Triticum aestivum</i>	monocot
121	87124328	ADP-glucose pyrophosphorylase	<i>Synechococcus</i> sp. RS9917	cyanobacteria
122	284051940	glucose-1-phosphate adenyltransferase	<i>Arthrospira platensis</i> str. Paraca	cyanobacteria
123	119509469	glucose-1-phosphate adenyltransferase	<i>Nodularia spumigena</i> CCY9414	cyanobacteria
124	126660345	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. CCY0110	cyanobacteria
125	126658160	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. CCY0110	cyanobacteria
126	318041355	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. CB0101	cyanobacteria
127	317969822	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. CB0205	cyanobacteria
128	87302823	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. WH 5701	cyanobacteria
129	119490200	glucose-1-phosphate adenyltransferase	<i>Lyngbya</i> sp. PCC 8106	cyanobacteria
130	116074714	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. RS9916	cyanobacteria
131	116070673	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. BL107	cyanobacteria
132	88808518	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. WH 7805	cyanobacteria

133	37523829	glucose-1-phosphate adenyltransferase	<i>Gloeobacter violaceus</i> PCC 7421	cyanobacteria
134	33861326	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. CCMP1986	cyanobacteria
135	33240292	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. CCMP1375	cyanobacteria
136	22298830	glucose-1-phosphate adenyltransferase	<i>Thermosynechococcus elongatus</i> BP-1	cyanobacteria
137	218248785	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. PCC 8801	cyanobacteria
138	186686123	glucose-1-phosphate adenyltransferase	<i>Nostoc punctiforme</i> PCC 73102	cyanobacteria
139	124023387	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. MIT 9303	cyanobacteria
140	86608545	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. JA-2-3B'a(2-13)	cyanobacteria
141	86606226	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. JA-3-3Ab	cyanobacteria
142	33862839	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. MIT 9313	cyanobacteria
143	33865652	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. WH 8102	cyanobacteria
144	17232137	glucose-1-phosphate adenyltransferase	<i>Nostoc</i> sp. PCC 7120	cyanobacteria
145	159903534	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. MIT 9211	cyanobacteria
146	126696167	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. MIT 9301	cyanobacteria
147	124025514	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. NATL1A	cyanobacteria
148	123968364	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. AS9601	cyanobacteria
149	123966049	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. MIT 9515	cyanobacteria
150	113954397	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. CC9311	cyanobacteria
151	218437477	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. PCC 7424	cyanobacteria
152	78779161	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. MIT 9312	cyanobacteria
153	16332282	glucose-1-phosphate adenyltransferase	<i>Synechocystis</i> sp. PCC 6803	cyanobacteria
154	257061844	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. PCC 8802	cyanobacteria
155	56750930	glucose-1-phosphate adenyltransferase	<i>Synechococcus elongatus</i> PCC 6301	cyanobacteria
156	220910118	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. PCC 7425	cyanobacteria
157	81299414	glucose-1-phosphate adenyltransferase	<i>Synechococcus elongatus</i> PCC 7942	cyanobacteria
158	78212786	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. CC9605	cyanobacteria
159	78184800	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. CC9902	cyanobacteria

160	158335435	glucose-1-phosphate adenyltransferase	<i>Acaryochloris marina</i> MBIC11017	cyanobacteria
161	158335089	glucose-1-phosphate adenyltransferase	<i>Acaryochloris marina</i> MBIC11017	cyanobacteria
162	113477795	glucose-1-phosphate adenyltransferase	<i>Trichodesmium erythraeum</i> IMS101	cyanobacteria
163	75908241	glucose-1-phosphate adenyltransferase	<i>Anabaena variabilis</i> ATCC 29413	cyanobacteria
164	172037571	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. ATCC 51142	cyanobacteria
165	172035903	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. ATCC 51142	cyanobacteria
166	170076729	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. PCC 7002	cyanobacteria
167	157413198	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. MIT 9215	cyanobacteria
168	72382015	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. NATL2A	cyanobacteria
169	148242352	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. RCC307	cyanobacteria
170	148239634	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. WH 7803	cyanobacteria
171	166365546	glucose-1-phosphate adenyltransferase	<i>Microcystis aeruginosa</i> NIES-843	cyanobacteria
172	254526910	glucose-1-phosphate adenyltransferase	<i>Prochlorococcus marinus</i> str. MIT 9202	cyanobacteria
173	254432095	glucose-1-phosphate adenyltransferase	<i>Cyanobium</i> sp. PCC 7001	cyanobacteria
174	307151922	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. PCC 7822	cyanobacteria
175	298492804	glucose-1-phosphate adenyltransferase	<i>Nostoc azollae</i> 0708	cyanobacteria
176	332709240	glucose-1-phosphate adenyltransferase	<i>Lyngbya majuscula</i> 3L	cyanobacteria
177	354565610	glucose-1-phosphate adenyltransferase	<i>Fischerella</i> sp. JSC-11	cyanobacteria
178	354556049	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. ATCC 51472	cyanobacteria
179	354553549	glucose-1-phosphate adenyltransferase	<i>Cyanothece</i> sp. ATCC 51472	cyanobacteria
180	352094384	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. WH 8016	cyanobacteria
181	284929352	glucose-1-phosphate adenyltransferase	cyanobacterium UCYN-A	cyanobacteria
182	334120832	glucose-1-phosphate adenyltransferase	<i>Microcoleus vaginatus</i> FGP-2	cyanobacteria
183	300865383	glucose-1-phosphate adenyltransferase	<i>Oscillatoria</i> sp. PCC 6506	cyanobacteria
184	282899378	glucose-1-phosphate adenyltransferase	<i>Cylindrospermopsis raciborskii</i> CS-505	cyanobacteria
185	282895605	glucose-1-phosphate adenyltransferase	<i>Raphidiopsis brookii</i> D9	cyanobacteria
186	209527099	glucose-1-phosphate adenyltransferase	<i>Arthrospira maxima</i> CS-328	cyanobacteria

187	67924676	glucose-1-phosphate adenyltransferase	<i>Crocospaera watsonii</i> WH 8501	cyanobacteria
188	260436638	glucose-1-phosphate adenyltransferase	<i>Synechococcus</i> sp. WH 8109	cyanobacteria
189	303273364	adp-glucose pyrophosphorylase	<i>Micromonas pusilla</i> CCMP1545	green alga
190	303271247	adp-glucose pyrophosphorylase	<i>Micromonas pusilla</i> CCMP1545	green alga
191	255070935	adp-glucose pyrophosphorylase	<i>Micromonas</i> sp. RCC299	green alga
192	255080070	adp-glucose pyrophosphorylase	<i>Micromonas</i> sp. RCC299	green alga
193	159470605	ADP-glucose pyrophosphorylase large subunit	<i>Chlamydomonas reinhardtii</i>	green alga
194	159467349	ADP-glucose pyrophosphorylase small subunit	<i>Chlamydomonas reinhardtii</i>	green alga
195	308814250	AGPLU2 (ISS)	<i>Ostreococcus tauri</i>	green alga
196	308806175	AGPSU1 (ISS)	<i>Ostreococcus tauri</i>	green alga
197	302849075	hypothetical protein VOLCADRAFT_76956	<i>Volvox carteri</i> f. nagariensis	green alga
198	302840808	hypothetical protein VOLCADRAFT_75183	<i>Volvox carteri</i> f. nagariensis	green alga
199	145356323	predicted protein	<i>Ostreococcus lucimarinus</i> CCE9901	green alga
200	145349062	predicted protein	<i>Ostreococcus lucimarinus</i> CCE9901	green alga
201	302825850	hypothetical protein SELMODRAFT_138695	<i>Selaginella moellendorffii</i>	moss
202	302815217	hypothetical protein SELMODRAFT_129625	<i>Selaginella moellendorffii</i>	moss
203	302802313	hypothetical protein SELMODRAFT_117069	<i>Selaginella moellendorffii</i>	moss
204	302800351	hypothetical protein SELMODRAFT_115472	<i>Selaginella moellendorffii</i>	moss
205	302798196	hypothetical protein SELMODRAFT_233627	<i>Selaginella moellendorffii</i>	moss
206	302788037	hypothetical protein SELMODRAFT_267891	<i>Selaginella moellendorffii</i>	moss
207	302783933	hypothetical protein SELMODRAFT_149205	<i>Selaginella moellendorffii</i>	moss
208	302773934	hypothetical protein SELMODRAFT_231637	<i>Selaginella moellendorffii</i>	moss
209	302769466	hypothetical protein SELMODRAFT_169778	<i>Selaginella moellendorffii</i>	moss