Appendix B. The information of candidate TFs associated with yield based on the RNA-seq library

Code	Gene family	Gene ID	Function annotation
1	MADS-box	PB.58836.1	Pistillata MADS-box protein [Malus domestica]
2		PB.59945.1	pistillata MADS-box protein [Paeonia suffruticosa]
3		PB.59889.2	pistillata MADS-box protein [Paeonia suffruticosa]
4		PB.63552.1	MADS-box protein TM3 subfamily [Coffea arabica]
5		PB.58716.1	MADS-box protein 4 [Vitis vinifera]
6		PB.5375.1	MADS-box protein 5 [Paeonia suffruticosa]
7		PB.53794.2	MADS-box protein 3 [Vitis vinifera]
8		PB.50081.1	MADS-box protein 3 [Vitis vinifera]
9		PB.55834.2	MADS-box protein 5 [Paeonia suffruticosa]
10		PB.57703.1	MADS box transcription factor SEPALLATA1 [Paeonia lactiflora]
11		PB.53579.2	PREDICTED: MADS-box transcription factor 23-like isoform X2 [Vitis vinifera]
12		PB.59842.1	PREDICTED: MADS-box protein FBP24 [Vitis vinifera]
13		PB.44987.1	PREDICTED: MADS-box protein CMB1-like isoform X4 [Populus euphratica]
14		PB.58162.1	PREDICTED: MADS-box protein AGL72-like [Tarenaya hassleriana]
15		PB.32845.1	PREDICTED: MADS-box transcription factor 23 isoform X5 [Theobroma cacao]
16		PB.49089.1	agamous [Paeonia suffruticosa]
17		PB.41954.3	PREDICTED: agamous-like MADS-box protein AGL82 [Prunus mume]
18		PB.42396.2	PREDICTED: agamous-like MADS-box protein AGL104 isoform X4 [Citrus sinensis]
19		PB.59950.1	PREDICTED: agamous-like MADS-box protein AGL12 [Vitis vinifera]
20		PB.64552.1	PREDICTED: agamous-like MADS-box protein AGL65 [Vitis vinifera]
21		PB.51016.1	PREDICTED: agamous-like MADS-box protein AGL15 isoform X1 [Ricinus communis]
22		PB.45186.1	PREDICTED: agamous-like MADS-box protein AGL30 isoform X3 [Citrus sinensis]
23		PB.42133.1	AP1/FUL-like protein [Paeonia suffruticosa]

24		PB.36542.2	AP1/FUL-like protein [Paeonia suffruticosa]
25		PB.53877.2	AP1/FUL-like protein [Paeonia suffruticosa]
26		PB.54789.1	euAP1 APETALA1-like MADS-box [Paeonia suffruticosa]
27	MYB	PB.34252.1	myb domain protein 4a [Camellia sinensis]。
28		PB.33363.1	myb domain protein 4a [Camellia sinensis]。
29		PB.13528.5	myb domain protein 4a [Camellia sinensis]。
30		PB.26194.10	isoform 3 of cyclin-d-binding myb-like transcription factor 1 [Quercus suber]
31		PB.52697.1	MYB transcription factor [Paeonia suffruticosa]
32		PB.56012.1	myb transcription factor 2 [Silene littorea]
33		PB.55405.1	PREDICTED: myb-like protein J [Juglans regia]
34		PB.35855.1	MYB transcription factor MYB79 [Glycine max]
35		PB.59960.1	PREDICTED: transcription factor MYB114 [Ricinus communis]
36		PB.52649.2	transcription factor WER-like [Prunus avium]
37		PB.50929.1	PREDICTED: transcription factor MYB12 [Ziziphus jujuba]
38		PB.49899.2	PREDICTED: transcription factor MYB12 [Ziziphus jujuba]
39		PB.61016.1	PREDICTED: myb-related protein 305-like [Juglans regia]。
40		PB.54617.1	PREDICTED: transcription factor MYB24 [Vitis vinifera]
41		PB.55484.1	PREDICTED: transcription factor MYB26 [Nelumbo nucifera]
42		PB.57113.1	PREDICTED: transcription factor MYB44 [Jatropha curcas]
43		PB.54818.1	PREDICTED: transcription factor MYB44 [Jatropha curcas]
44		PB.48075.9	PREDICTED: transcription factor MYB44-like [Vitis vinifera]
45		PB.36428.1	PREDICTED: transcription factor MYB48 [Vitis vinifera]
46		PB.45912.1	PREDICTED: transcription factor MYB82-like [Camelina sativa]
47		PB.48069.2	transcription factor MYB61-like [Manihot esculenta]
48		PB.56343.4	PREDICTED: protein PHR1-LIKE 2 [Vitis vinifera]
49		PB.41467.1	PREDICTED: protein PHR1-LIKE 2 [Vitis vinifera]
50		PB.51245.3	PREDICTED: protein PHR1-LIKE 3-like isoform X1 [Nelumbo nucifera]

51	PB.48769.1	myb family transcription factor phl11 [Quercus suber]
52	PB.57971.1	PREDICTED: transcription factor MYB1R1 [Vitis vinifera]
53	PB.48245.2	PREDICTED: transcription factor MYB1R1 [Vitis vinifera]
54	PB.56535.1	PREDICTED: transcription repressor MYB6-like [Juglans regia]
55	PB.15637.2	PREDICTED: L10-interacting MYB domain-containing protein-like [Arachis duranensis]
56	PB.5429.5	PREDICTED: myb-like protein X isoform X3 [Populus euphratica]
57	PB.59485.1	PREDICTED: transcription repressor MYB6 [Amborella trichopoda]
58	PB.59400.1	myb-related protein 308-like [Cajanus cajan]
59	PB.59901.1	PREDICTED: myb-related protein 308-like [Populus euphratica]
60	PB.53387.3	PREDICTED: myb-related protein 306 [Vitis vinifera]
61	PB.52130.1	mixta-like2 MYB transcription factor [Thalictrum dioicum]
62	PB.50356.1	PREDICTED: L10-interacting MYB domain-containing protein-like isoform X1 [Ziziphus jujuba]
63	PB.54373.1	PREDICTED: myb-related protein 308 [Theobroma cacao]
64	PB.52122.2	Myb domain protein 106 [Theobroma cacao]
65	PB.52852.2	myb-related protein 306-like [Prunus avium]
66	PB.53756.4	transcription factor MYB41 [Herrania umbratica]
67	PB.54887.1	PREDICTED: L10-interacting MYB domain-containing protein-like isoform X1 [Ziziphus jujuba]
68	PB.45662.5	transcription factor MYB6 [Jatropha curcas]
69	PB.45821.1	PREDICTED: myb-related protein B-like [Juglans regia]
70	PB.42979.1	PREDICTED: target of Myb protein 1-like [Vitis vinifera]
71	PB.36093.1	PREDICTED: myb-related protein B-like [Nicotiana sylvestris]
72	PB.34237.4	Transcriptional activator Myb, partial [Capsicum baccatum]
73	PB.28062.1	PREDICTED: target of Myb protein 1 [Fragaria vesca subsp. vesca]
74	PB.18933.5	PREDICTED: myb-related protein Myb4-like [Populus euphratica]
75	PB.20226.2	PREDICTED: myb-related protein Myb4-like [Populus euphratica]
76	PB.16044.1	PREDICTED: myb-related protein 3R-1-like isoform X1 [Vitis vinifera]
77	PB.9600.12	Myb_DNA-bind_4 domain-containing protein [Cephalotus follicularis]

78		PB.32979.2	Myb_DNA-bind_6 domain-containing protein [Cephalotus follicularis]
79		PB.7084.14	PREDICTED: myb-like protein X [Nelumbo nucifera]
80		PB.4491.1	PREDICTED: myb-related protein 3R-1-like isoform X1 [Vitis vinifera]
81		PB.249.23	PREDICTED: myb-related protein 3R-1-like isoform X1 [Vitis vinifera]
82		PB.35529.1	PREDICTED: transcription factor GAMYB [Vitis vinifera]
83		PB.58199.1	myb domain protein 4a [Camellia sinensis]
84		PB.56618.1	Myb-like domain [Macleaya cordata]
85		PB.50160.1	Myb_DNA-bind_4 domain-containing protein [Cephalotus follicularis]
86	AP2	PB.40151.1	PREDICTED: AP2/ERF and B3 domain-containing transcription factor At1g50680-like [Nelumbo nucifera]
87		PB.50898.1	PREDICTED: AP2/ERF and B3 domain-containing transcription factor RAV1-like [Vitis vinifera]
88		PB.49031.3	PREDICTED: AP2/ERF and B3 domain-containing transcription factor RAV1-like [Vitis vinifera]
89		PB.57145.1	AP2 domain class transcription factor [Medicago truncatula]
90		PB.36607.2	AP2 transcriptional activator variant, partial [Triticum turgidum subsp. durum]
91		PB.57344.1	AP2/ERF domain-containing protein, partial [Cynara cardunculus var. scolymus]
92		PB.33003.1	AP2/ERF transcription factor [Trema orientalis]
93		PB.32595.6	AP2-like ethylene-responsive transcription factor AIL1 [Herrania umbratica]
94		PB.6086.1	AP2-like ethylene-responsive transcription factor ANT [Carica papaya]
95		PB.16895.1	AP2-like ethylene-responsive transcription factor AIL6 [Theobroma cacao]
96		PB.38745.1	AP2-like ethylene-responsive transcription factor AIL6 [Theobroma cacao]
97		PB.35214.4	AP2-like ethylene-responsive transcription factor AIL6 [Theobroma cacao]
98		PB.51298.1	PREDICTED: AP2-like ethylene-responsive transcription factor AIL5 [Nelumbo nucifera]
99		PB.39379.5	PREDICTED: AP2-like ethylene-responsive transcription factor AIL5 [Nelumbo nucifera]
100		PB.40354.1	PREDICTED: AP2-like ethylene-responsive transcription factor AIL5 [Vitis vinifera]
101		PB.42627.1	PREDICTED: AP2-like ethylene-responsive transcription factor AIL6 [Vitis vinifera]
102		PB.27348.1	PREDICTED: AP2-like ethylene-responsive transcription factor ANT [Vitis vinifera]
103		PB.39671.1	PREDICTED: AP2-like ethylene-responsive transcription factor ANT [Vitis vinifera]
104		PB.18023.10	PREDICTED: AP2-like ethylene-responsive transcription factor ANT [Vitis vinifera]

105	PB.6753.1	PREDICTED: AP2-like ethylene-responsive transcription factor ANT [Vitis vinifera]
106	PB.5853.33	PREDICTED: AP2-like ethylene-responsive transcription factor ANT [Vitis vinifera]
107	PB.56852.1	PREDICTED: AP2-like ethylene-responsive transcription factor ANT [Ziziphus jujuba]
108	PB.28809.3	PREDICTED: AP2-like ethylene-responsive transcription factor ANT [Ziziphus jujuba]
109	PB.28967.1	PREDICTED: AP2-like ethylene-responsive transcription factor PLT1 [Populus euphratica]
110	PB.7065.1	PREDICTED: AP2-like ethylene-responsive transcription factor PLT1 [Populus euphratica]
111	PB.9687.1	PREDICTED: AP2-like ethylene-responsive transcription factor PLT2 [Juglans regia]
112	PB.8190.1	PREDICTED: AP2-like ethylene-responsive transcription factor PLT1 [Populus euphratica]
113	PB.3987.3	PREDICTED: AP2-like ethylene-responsive transcription factor PLT1 [Populus euphratica]
114	PB.3749.4	PREDICTED: AP2-like ethylene-responsive transcription factor PLT2 [Vitis vinifera]
115	PB.3727.9	PREDICTED: AP2-like ethylene-responsive transcription factor PLT2 [Vitis vinifera]
116	PB.4787.1	PREDICTED: AP2-like ethylene-responsive transcription factor PLT2 [Ziziphus jujuba]
117	PB.9193.1	PREDICTED: AP2-like ethylene-responsive transcription factor PLT2 isoform X1 [Fragaria vesca subsp. vesca]
118	PB.40948.1	PREDICTED: AP2-like ethylene-responsive transcription factor TOE3 isoform X1 [Prunus mume]
119	PB.34858.1	PREDICTED: ethylene-responsive transcription factor WRI1-like isoform X2 [Juglans regia]
120	PB.48645.2	PREDICTED: ethylene-responsive transcription factor RAP2-4-like isoform X1 [Vitis vinifera]
121	PB.39602.1	PREDICTED: ethylene-responsive transcription factor RAP2-7 isoform X2 [Cucumis sativus]
122	PB.35869.3	PREDICTED: ethylene-responsive transcription factor RAP2-7 isoform X1 [Vitis vinifera]
123	PB.61307.2	PREDICTED: ethylene-responsive transcription factor 12 [Vitis vinifera]
124	PB.60194.1	PREDICTED: ethylene-responsive transcription factor 3-like [Vitis vinifera]
125	PB.45546.6	PREDICTED: ethylene-responsive transcription factor 3-like [Vitis vinifera]
126	PB.57609.2	PREDICTED: ethylene-responsive transcription factor 4 [Vitis vinifera]
127	PB.53173.1	PREDICTED: ethylene-responsive transcription factor 5 [Vitis vinifera]
128	PB.45149.2	PREDICTED: ethylene-responsive transcription factor CRF2 [Vitis vinifera]
129	PB.1195.1	PREDICTED: ethylene-responsive transcription factor ERF003-like [Nelumbo nucifera]
130	PB.63627.1	PREDICTED: ethylene-responsive transcription factor RAP2-1-like [Juglans regia]
131	PB.65493.1	PREDICTED: ethylene-responsive transcription factor ERF016 [Vitis vinifera]

132		PB.61382.1	PREDICTED: ethylene-responsive transcription factor ERF026-like [Nelumbo nucifera]
133		PB.53098.1	PREDICTED: ethylene-responsive transcription factor ERF061-like [Populus euphratica]
134		PB.41803.9	PREDICTED: ethylene-responsive transcription factor ERF073 [Jatropha curcas]
135		PB.55385.1	PREDICTED: ethylene-responsive transcription factor ERF105-like [Nelumbo nucifera]
136		PB.39155.1	PREDICTED: ethylene-responsive transcription factor ERF118 [Vitis vinifera]
137		PB.33711.4	PREDICTED: ethylene-responsive transcription factor ERF118 [Vitis vinifera]
138		PB.23012.1	PREDICTED: ethylene-responsive transcription factor RAP2-2-like [Vitis vinifera]
139		PB.40358.2	PREDICTED: ethylene-responsive transcription factor RAP2-4 [Vitis vinifera]
140		PB.41377.2	PREDICTED: ethylene-responsive transcription factor RAP2-4 [Vitis vinifera]
141		PB.41538.1	PREDICTED: ethylene-responsive transcription factor RAP2-7 isoform X2 [Vitis vinifera]
142		PB.38710.6	PREDICTED: ethylene-responsive transcription factor RAP2-7-like isoform X1 [Nelumbo nucifera]
143		PB.54694.1	PREDICTED: ethylene-responsive transcription factor-like protein At4g13040 isoform X4 [Cucumis sativus]
144		PB.38966.1	PREDICTED: ethylene-responsive transcription factor RAP2-7 isoform X2 [Theobroma cacao]
145		PB.49968.2	ethylene-responsive transcription factor RAP2-1-like [Carica papaya]，
146		PB.30760.1	ethylene-responsive transcription factor [Quercus suber]
147		PB.55158.1	ethylene-responsive transcription factor RAP2-12-like [Hevea brasiliensis]
148		PB.45610.2	ethylene-responsive transcription factor RAP2-12-like [Hevea brasiliensis]
149		PB.56734.1	ethylene-responsive transcription factor 1A [Paeonia lactiflora]
150		PB.46556.4	ethylene-responsive transcription factor 3-like [Jatropha curcas]
151		PB.44222.1	ethylene-responsive transcription factor CRF4-like [Durio zibethinus]
152		PB.48385.1	AP2 domain-containing protein [Cephalotus follicularis]
153		PB.29891.1	PREDICTED: AP2-associated protein kinase 1 isoform X2 [Prunus mume]
154		PB.44479.2	ERF transcription factor [Paeonia suffruticosa]
155		PB.24110.1	ERF transcription factor [Paeonia suffruticosa]
156		PB.4643.15	ERF transcription factor [Paeonia suffruticosa]
157	BHLH	PB.28156.2	bHLH transcription factor [Paeonia suffruticosa]
158		PB.12882.2	bHLH transcription factor [Paeonia suffruticosa]

159	PB.27982.1	bHLH transcription factor [Paeonia suffruticosa]
160	PB.25717.3	bHLH transcription factor [Paeonia suffruticosa]
161	PB.23564.2	bHLH transcription factor [Paeonia suffruticosa]
162	PB.26400.1	bHLH transcription factor [Paeonia suffruticosa]
163	PB.21845.1	bHLH transcription factor [Paeonia suffruticosa]
164	PB.21690.1	bHLH transcription factor [Paeonia suffruticosa]
165	PB.20681.15	bHLH transcription factor [Paeonia suffruticosa]
166	PB.14125.1	bHLH transcription factor [Paeonia suffruticosa]
167	PB.9004.1	bHLH transcription factor [Paeonia suffruticosa]
168	PB.9881.1	bHLH transcription factor [Paeonia suffruticosa]
169	PB.8370.1	bHLH transcription factor [Paeonia suffruticosa]
170	PB.5991.1	bHLH transcription factor [Paeonia suffruticosa]
171	PB.36620.1	transcription factor bhlh91 [Quercus suber]
172	PB.30711.2	transcription factor bhlh91 [Quercus suber]
173	PB.60004.1	PREDICTED: transcription factor bHLH104 isoform X1 [Citrus sinensis]
174	PB.61860.1	PREDICTED: transcription factor bHLH104-like [Nelumbo nucifera]
175	PB.62187.1	PREDICTED: transcription factor bHLH104-like [Vigna radiata var. radiata]
176	PB.57453.1	PREDICTED: transcription factor bHLH106-like [Fragaria vesca subsp. vesca]
177	PB.25180.1	PREDICTED: transcription factor bHLH112 isoform X2 [Vitis vinifera]
178	PB.58085.1	PREDICTED: transcription factor bHLH113 isoform X2 [Prunus mume]
179	PB.57537.1	PREDICTED: transcription factor bHLH113 isoform X3 [Vitis vinifera]
180	PB.54763.1	PREDICTED: transcription factor bHLH121 [Pyrus x bretschneideri]
181	PB.42377.1	LOW QUALITY PROTEIN: transcription factor bHLH122-like [Durio zibethinus]
182	PB.39795.1	PREDICTED: transcription factor bHLH122 isoform X2 [Jatropha curcas]
183	PB.38848.1	PREDICTED: transcription factor bHLH123 isoform X1 [Vitis vinifera]
184	PB.49637.2	PREDICTED: transcription factor bHLH128 isoform X2 [Vitis vinifera]
185	PB.40560.1	PREDICTED: transcription factor bHLH122-like isoform X1 [Populus euphratica]

186	PB.50241.1	PREDICTED: transcription factor bHLH130-like [Vitis vinifera]
187	PB.33949.1	PREDICTED: transcription factor bHLH140 isoform X1 [Vitis vinifera]
188	PB.19291.8	PREDICTED: transcription factor bHLH140 isoform X2 [Vitis vinifera]
189	PB.18829.2	PREDICTED: transcription factor bHLH143-like isoform X1 [Vitis vinifera]
190	PB.13140.4	transcription factor bHLH145 [Herrania umbratica]
191	PB.63393.1	PREDICTED: transcription factor bHLH147 [Nelumbo nucifera]
192	PB.53994.4	PREDICTED: transcription factor bHLH147 isoform X1 [Vitis vinifera]
193	PB.52481.3	PREDICTED: transcription factor bHLH147 isoform X1 [Vitis vinifera]
194	PB.61150.1	PREDICTED: transcription factor bHLH25-like [Juglans regia]
195	PB.12182.1	PREDICTED: transcription factor bHLH25-like [Vitis vinifera]
196	PB.15089.1	PREDICTED: transcription factor bHLH30 [Ricinus communis]
197	PB.41344.7	PREDICTED: transcription factor bHLH30-like [Populus euphratica]
198	PB.59514.2	PREDICTED: transcription factor bHLH35-like [Pyrus x bretschneideri]
199	PB.44353.4	PREDICTED: transcription factor bHLH48 [Vitis vinifera]
200	PB.36412.1	PREDICTED: transcription factor bHLH49 isoform X1 [Vitis vinifera]
201	PB.16398.11	PREDICTED: transcription factor bHLH49 isoform X1 [Vitis vinifera]
202	PB.23497.1	Basic helix-loop-helix transcription factor [Parasponia andersonii]
203	PB.64769.1	PREDICTED: transcription factor bHLH51 [Vitis vinifera]
204	PB.61918.2	PREDICTED: transcription factor ICE1-like isoform X1 [Prunus mume]
205	PB.45594.2	PREDICTED: transcription factor bHLH63 [Vitis vinifera]
206	PB.20659.1	PREDICTED: transcription factor bHLH63 [Vitis vinifera]
207	PB.61331.1	Basic helix-loop-helix transcription factor [Parasponia andersonii]
208	PB.45722.1	PREDICTED: transcription factor bHLH68 [Ricinus communis]
209	PB.45322.6	PREDICTED: transcription factor bHLH68 [Vitis vinifera]
210	PB.38882.2	PREDICTED: transcription factor bHLH68 [Vitis vinifera]
211	PB.51146.1	PREDICTED: transcription factor bHLH68-like [Vitis vinifera]
212	PB.48928.1	PREDICTED: transcription factor bHLH69-like isoform X2 [Vitis vinifera]

213	PB.54762.1	transcription factor bHLH66-like isoform X2 [Olea europaea var. sylvestris]
214	PB.51309.1	PREDICTED: transcription factor bHLH69-like isoform X2 [Vitis vinifera]
215	PB.29949.1	PREDICTED: transcription factor bHLH69-like isoform X2 [Vitis vinifera]
216	PB.53640.1	transcription factor bhlh96 [Quercus suber]
217	PB.40828.7	transcription factor bhlh96 [Quercus suber]
218	PB.33822.1	PREDICTED: transcription factor bHLH74 isoform X2 [Prunus mume]
219	PB.23724.8	PREDICTED: transcription factor bHLH74 isoform X2 [Prunus mume]
220	PB.17214.4	PREDICTED: transcription factor bHLH74 isoform X2 [Prunus mume]
221	PB.55387.1	transcription factor BHLH094 isoform X2 [Momordica charantia]
222	PB.58855.1	PREDICTED: transcription factor bHLH79 isoform X1 [Ziziphus jujuba]
223	PB.54828.1	PREDICTED: transcription factor bHLH79 isoform X1 [Ziziphus jujuba]
224	PB.49472.4	PREDICTED: transcription factor bHLH79 isoform X1 [Ziziphus jujuba]
225	PB.46142.8	PREDICTED: transcription factor bHLH79-like isoform X2 [Nelumbo nucifera]
226	PB.43399.1	PREDICTED: transcription factor bHLH87 [Populus euphratica]
227	PB.11210.4	PREDICTED: transcription factor bHLH87 [Vitis vinifera]
228	PB.46999.1	PREDICTED: transcription factor bHLH91 [Ricinus communis]
229	PB.49165.1	PREDICTED: transcription factor bHLH91-like [Nelumbo nucifera]
230	PB.44849.2	Basic helix-loop-helix transcription factor [Parasponia andersonii]
231	PB.31016.1	Basic helix-loop-helix transcription factor [Parasponia andersonii]
232	PB.48873.1	PREDICTED: transcription factor bHLH93 [Nelumbo nucifera]
233	PB.48506.7	PREDICTED: transcription factor bHLH93 [Nelumbo nucifera]
234	PB.51785.1	PREDICTED: transcription factor bHLH93 [Vitis vinifera]
235	PB.63030.1	PREDICTED: transcription factor bHLH95-like [Citrus sinensis]
236	PB.36910.10	PREDICTED: transcription factor bHLH96-like [Nelumbo nucifera]
237	PB.45004.1	Transcription factor bHLH140 family, partial [Cajanus cajan]
238	PB.19628.1	PREDICTED: transcription factor SPATULA isoform X1 [Vitis vinifera]
239	PB.49835.3	PREDICTED: transcription factor SPATULA isoform X2 [Vitis vinifera]

240		PB.40753.2	PREDICTED: transcription factor SPATULA isoform X2 [Vitis vinifera]
241		PB.26331.4	PREDICTED: transcription factor MYC2-like [Vitis vinifera]
242		PB.61985.1	PREDICTED: transcription factor MYC2-like [Vitis vinifera]
243		PB.6104.1	Myc-type [Macleaya cordata]
244		PB.63551.1	Myc-type [Macleaya cordata]
245		PB.18398.7	Basic helix-loop-helix transcription factor [Parasponia andersonii]
246		PB.34254.7	PREDICTED: transcription factor PIF1 isoform X2 [Vitis vinifera]
247		PB.1385.1	PREDICTED: transcription factor PIF3 isoform X3 [Ricinus communis]
248		PB.6454.21	PREDICTED: transcription factor PIF3-like isoform X1 [Vitis vinifera]
249		PB.31359.3	PREDICTED: transcription factor PIF4 [Vitis vinifera]
250		PB.35177.3	PREDICTED: transcription factor PIF4-like [Fragaria vesca subsp. vesca]
251		PB.18984.1	PREDICTED: transcription factor ILR3 [Brachypodium distachyon]
252		PB.58867.1	PREDICTED: transcription factor ILR3 [Ziziphus jujuba]
253		PB.58094.2	PREDICTED: transcription factor ILR3 [Ziziphus jujuba]
254		PB.55581.5	PREDICTED: transcription factor ILR3 [Ziziphus jujuba]
255		PB.57643.1	PREDICTED: transcription factor ILR3-like [Prunus mume]
256	TCP	PB.61740.1	PREDICTED: transcription factor TCP11 [Vitis vinifera]
257		PB.59930.1	PREDICTED: transcription factor TCP11-like [Ziziphus jujuba]
258		PB.48125.5	PREDICTED: transcription factor TCP15 [Vitis vinifera]
259		PB.30344.6	PREDICTED: transcription factor TCP15-like [Vitis vinifera]
260		PB.50380.4	PREDICTED: transcription factor TCP20 [Vitis vinifera]
261		PB.32065.1	PREDICTED: transcription factor TCP2-like [Vitis vinifera]
262		PB.2028.10	PREDICTED: transcription factor TCP2-like [Vitis vinifera]
263		PB.42993.2	PREDICTED: transcription factor TCP4 [Ricinus communis]
264		PB.19925.5	PREDICTED: transcription factor TCP4 [Vitis vinifera]
265		PB.45037.5	PREDICTED: transcription factor TCP7 [Nelumbo nucifera]
266		PB.57717.1	PREDICTED: transcription factor TCP9 [Nelumbo nucifera]

267		PB.41437.2	PREDICTED: transcription factor TCP9 [Nelumbo nucifera]
268		PB.42683.2	PREDICTED: transcription factor TCP9 [Vitis vinifera]
269		PB.60868.1	PREDICTED: transcription factor TCP9-like [Vitis vinifera]
270		PB.42060.4	PREDICTED: transcription factor CYCLOIDEA [Vitis vinifera]
271		PB.52476.1	Transcription factor, TCP [Corchorus olitorius]
272	GRAS	PB.54807.1	GRAS family transcription factor [Quercus suber]
273		PB.16537.2	GRAS family transcription factor [Theobroma cacao]
274		PB.22311.1	GRAS family transcription factor [Theobroma cacao]
275		PB.13169.1	GRAS family transcription factor [Theobroma cacao]
276		PB.14691.1	GRAS transcription factor [Trema orientalis]
277		PB.22143.1	GRAS transcription factor [Trema orientalis]
278		PB.24124.9	GRAS domain-containing protein, partial [Cephalotus follicularis]
279		PB.7809.1	GRAS transcription factor [Trema orientalis]
280		PB.5948.1	GRAS transcription factor [Trema orientalis]
281		PB.2355.5	GRAS transcription factor [Trema orientalis]
282		PB.46584.3	scarecrow-like protein 32 [Hevea brasiliensis]
283		PB.42766.1	scarecrow-like protein 18 [Momordica charantia]
284		PB.20960.1	PREDICTED: scarecrow-like protein 28 [Theobroma cacao]
285		PB.26780.1	scarecrow-like protein 14 [Prunus avium]
286		PB.25724.1	scarecrow-like protein 4 [Jatropha curcas]
287		PB.20904.22	scarecrow-like protein 4 [Jatropha curcas]
288		PB.1246.5	scarecrow-like protein 14 [Quercus suber]
289		PB.28367.1	scarecrow-like protein 21 isoform X1 [Hevea brasiliensis]
290		PB.29999.1	scarecrow-like protein 21 isoform X1 [Hevea brasiliensis]
291		PB.32203.3	PREDICTED: scarecrow-like transcription factor PAT1 [Vitis vinifera]
292		PB.34698.1	PREDICTED: scarecrow-like transcription factor PAT1 [Vitis vinifera]
293		PB.26001.1	PREDICTED: scarecrow-like transcription factor PAT1 [Vitis vinifera]

294	PB.48992.2	PREDICTED: scarecrow-like protein 23 [Ziziphus jujuba]
295	PB.32125.2	Scarecrow-like 3 [Theobroma cacao]
296	PB.19055.5	PREDICTED: scarecrow-like protein 27 [Vitis vinifera]
297	PB.49322.1	scarecrow-like protein 14 [Hevea brasiliensis]
298	PB.56345.1	PREDICTED: scarecrow-like protein 8 [Nelumbo nucifera]
299	PB.47737.1	PREDICTED: scarecrow-like protein 4 [Vitis vinifera]
300	PB.43588.2	PREDICTED: scarecrow-like protein 15 [Vitis vinifera]
301	PB.44401.1	PREDICTED: scarecrow-like protein 3 [Ziziphus jujuba]
302	PB.42936.1	PREDICTED: scarecrow-like protein 3 isoform X1 [Vitis vinifera]
303	PB.31852.7	PREDICTED: scarecrow-like protein 8 [Vitis vinifera]
304	PB.29144.1	PREDICTED: scarecrow-like protein 1 [Vitis vinifera]
305	PB.26713.1	PREDICTED: scarecrow-like protein 33 [Juglans regia]
306	PB.25844.7	PREDICTED: scarecrow-like protein 14 [Ricinus communis]
307	PB.27533.1	PREDICTED: scarecrow-like protein 9 [Nelumbo nucifera]
308	PB.25439.2	PREDICTED: scarecrow-like protein 27 [Vitis vinifera]
309	PB.25392.1	PREDICTED: scarecrow-like protein 21 [Vitis vinifera]
310	PB.23858.1	PREDICTED: scarecrow-like protein 18 [Populus euphratica]
311	PB.20931.1	PREDICTED: scarecrow-like protein 28 [Citrus sinensis]
312	PB.17929.14	PREDICTED: scarecrow-like protein 9 [Vitis vinifera]
313	PB.18353.5	PREDICTED: scarecrow-like protein 1 [Vitis vinifera]
314	PB.15462.3	PREDICTED: scarecrow-like protein 28 [Ziziphus jujuba]
315	PB.15977.17	PREDICTED: scarecrow-like protein 21 [Vitis vinifera]
316	PB.13855.1	PREDICTED: protein SCARECROW [Jatropha curcas]
317	PB.2823.1	PREDICTED: scarecrow-like protein 6 [Pyrus x bretschneideri]
318	PB.3793.54	PREDICTED: scarecrow-like protein 22 isoform X2 [Vitis vinifera]
319	PB.396.26	PREDICTED: protein SCARECROW [Jatropha curcas]
320	PB.58646.1	PREDICTED: protein SCARECROW-like, partial [Ziziphus jujuba]

321	HSF	PB.24178.1	Heat shock transcription factor [Trema orientalis]
322		PB.20399.5	Heat shock factor (HSF)-type [Macleaya cordata]
323		PB.44942.1	Heat shock factor (HSF)-type [Macleaya cordata]
324		PB.40630.1	Heat shock factor (HSF)-type [Macleaya cordata]
325		PB.46361.1	PREDICTED: heat stress transcription factor A-5-like [Ziziphus jujuba]
326		PB.26336.2	heat stress transcription factor A-2d-like isoform X3 [Durio zibethinus]
327		PB.23304.3	heat stress transcription factor A-7a [Prunus persica]
328		PB.36595.1	heat stress transcription factor A-7a [Prunus persica]
329		PB.34020.1	heat stress transcription factor A-7a [Prunus persica]
330		PB.51070.1	PREDICTED: heat stress transcription factor B-2b [Vitis vinifera]
331		PB.47891.2	PREDICTED: heat stress transcription factor B-4 [Jatropha curcas]
332		PB.52864.1	PREDICTED: heat stress transcription factor B-4 [Jatropha curcas]
333		PB.52165.4	HSF_DNA-bind domain-containing protein [Cephalotus follicularis]
334		PB.53862.1	PREDICTED: LOW QUALITY PROTEIN: heat stress transcription factor B-2a [Vitis vinifera]
335		PB.51448.2	PREDICTED: activator of 90 kDa heat shock protein ATPase homolog 1 [Ricinus communis]
336		PB.48536.1	PREDICTED: heat shock factor protein HSF24 [Vitis vinifera]
337		PB.37643.3	PREDICTED: heat shock cognate 70 kDa protein 2-like [Elaeis guineensis]
338		PB.29638.2	PREDICTED: heat shock protein 83 [Cucumis sativus]
339		PB.29004.2	PREDICTED: heat shock protein 90-2 [Pyrus x bretschneideri]
340		PB.17708.1	PREDICTED: heat shock 70 kDa protein 16 [Fragaria vesca subsp. vesca]
341		PB.16477.1	PREDICTED: heat shock 70 kDa protein 15-like [Ziziphus jujuba]
342		PB.61812.1	PREDICTED: heat shock 22 kDa protein, mitochondrial isoform X1 [Cicer arietinum]
343		PB.59868.1	23.6 kDa heat shock protein, mitochondrial-like [Durio zibethinus]
344		PB.42852.1	small heat shock protein, chloroplastic-like isoform X2 [Cucurbita moschata]
345		PB.41740.5	small heat shock protein, chloroplastic-like isoform X2 [Cucurbita moschata]
346		PB.40031.1	PREDICTED: heat shock cognate 70 kDa protein 2 [Ricinus communis]
347		PB.38939.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]

348	PB.45710.1	PREDICTED: heat shock factor protein HSF8 [Ricinus communis]
349	PB.45975.1	PREDICTED: heat shock protein 83-like isoform X1 [Gossypium raimondii]
350	PB.45475.1	PREDICTED: heat shock factor protein HSF24 [Vitis vinifera]
351	PB.41893.2	PREDICTED: LOW QUALITY PROTEIN: heat shock protein 83-like [Nicotiana sylvestris]
352	PB.37741.1	PREDICTED: heat shock 70 kDa protein [Gossypium raimondii]
353	PB.36145.1	Heat shock protein 70 family [Trema orientalis]
354	PB.31370.2	PREDICTED: heat shock cognate 70 kDa protein 2 [Vitis vinifera]
355	PB.36035.5	PREDICTED: heat shock cognate 70 kDa protein 2-like [Cicer arietinum]
356	PB.26141.3	PREDICTED: heat shock 70 kDa protein, mitochondrial [Vitis vinifera]
357	PB.32985.1	PREDICTED: heat shock protein 90-2-like [Daucus carota subsp. sativus]
358	PB.28205.9	PREDICTED: heat shock 70 kDa protein, mitochondrial isoform X2 [Nelumbo nucifera]
359	PB.29772.6	PREDICTED: heat shock protein 83 [Ricinus communis]
360	PB.33995.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Gossypium raimondii]
361	PB.33390.1	PREDICTED: heat shock 70 kDa protein [Gossypium raimondii]
362	PB.36765.1	PREDICTED: heat shock cognate 70 kDa protein 2 [Nicotiana sylvestris]
363	PB.25991.1	heat shock cognate 70 kDa protein 2-like [Herrania umbratica]
364	PB.24581.10	PREDICTED: heat shock protein 83 [Cucumis sativus]
365	PB.28517.6	PREDICTED: heat shock cognate 70 kDa protein 2 [Nicotiana tomentosiformis]
366	PB.31018.1	PREDICTED: heat shock cognate protein 80 [Ricinus communis]
367	PB.26797.1	PREDICTED: heat shock 70 kDa protein 6, chloroplastic-like [Gossypium raimondii]
368	PB.26294.5	PREDICTED: heat shock cognate 70 kDa protein 2 [Nicotiana sylvestris]
369	PB.26606.1	heat shock 70 kDa protein 14-like [Hevea brasiliensis]
370	PB.11203.1	Heat shock 70 kDa protein 15 [Capsicum baccatum]
371	PB.23654.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nelumbo nucifera]
372	PB.23450.1	PREDICTED: heat shock cognate 70 kDa protein 2 [Nicotiana tomentosiformis]
373	PB.24168.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nelumbo nucifera]
374	PB.23516.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Sesamum indicum]

375	PB.16655.2	PREDICTED: heat shock protein 83-like [Fragaria vesca subsp. vesca]
376	PB.20527.21	heat shock protein 90-6, mitochondrial [Quercus suber]
377	PB.23410.6	PREDICTED: heat shock protein 83 [Prunus mume]
378	PB.14866.15	PREDICTED: heat shock 70 kDa protein, mitochondrial [Vitis vinifera]
379	PB.19070.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nelumbo nucifera]
380	PB.20946.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]
381	PB.21471.15	PREDICTED: heat shock 70 kDa protein 16 [Ziziphus jujuba]
382	PB.14850.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]
383	PB.15988.1	PREDICTED: heat shock 70 kDa protein, mitochondrial [Vitis vinifera]
384	PB.10348.1	PREDICTED: heat shock 70 kDa protein 15-like [Ziziphus jujuba]
385	PB.11285.1	PREDICTED: heat shock cognate 70 kDa protein 2 [Prunus mume]
386	PB.12752.8	PREDICTED: heat shock cognate 70 kDa protein 2-like isoform X1 [Ziziphus jujuba]
387	PB.9020.16	PREDICTED: heat shock cognate 70 kDa protein 2 [Sesamum indicum]
388	PB.8783.13	PREDICTED: heat shock protein 83 [Cucumis melo]
389	PB.23815.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]
390	PB.1346.15	PREDICTED: heat shock cognate 70 kDa protein 2 [Prunus mume]
391	PB.1348.21	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]
392	PB.1947.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]
393	PB.1208.11	PREDICTED: heat shock 70 kDa protein 15-like [Ziziphus jujuba]
394	PB.1061.2	PREDICTED: heat shock cognate 70 kDa protein 2 [Cucumis melo]
395	PB.1449.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana sylvestris]
396	PB.163.44	PREDICTED: heat shock cognate 70 kDa protein 2 [Prunus mume]
397	PB.752.1	PREDICTED: heat shock 70 kDa protein 15-like [Ziziphus jujuba]
398	PB.472.1	PREDICTED: heat shock 70 kDa protein 15-like [Ziziphus jujuba]
399	PB.1477.1	PREDICTED: heat shock 70 kDa protein 15-like [Ziziphus jujuba]
400	PB.450.42	PREDICTED: heat shock 70 kDa protein 15-like [Ziziphus jujuba]
401	PB.1199.1	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]

402	PB.639.127	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]
403	PB.904.16	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]
404	PB.1100.16	PREDICTED: heat shock cognate 70 kDa protein 2-like [Nicotiana tomentosiformis]
405	PB.65291.1	PREDICTED: heat shock 22 kDa protein, mitochondrial-like [Erythranthe guttata]
406	PB.64541.1	PREDICTED: heat shock protein 83 [Prunus mume]
407	PB.61850.1	PREDICTED: heat shock cognate protein 80 [Vitis vinifera]
408	PB.62241.3	PREDICTED: heat shock 22 kDa protein, mitochondrial-like [Erythranthe guttata]
409	PB.26144.1	PREDICTED: heat shock cognate protein 80-like [Cicer arietinum]
410	PB.62533.1	PREDICTED: 22.0 kDa class IV heat shock protein [Ziziphus jujuba]
411	PB.55062.2	PREDICTED: 17.1 kDa class II heat shock protein-like [Juglans regia]
412	PB.55174.1	PREDICTED: 17.9 kDa class II heat shock protein-like [Vigna radiata var. radiata]
413	PB.55233.1	PREDICTED: 17.1 kDa class II heat shock protein-like [Fragaria vesca subsp. vesca]
414	PB.56064.1	PREDICTED: 17.1 kDa class II heat shock protein [Ricinus communis]
415	PB.48392.1	PREDICTED: 18.1 kDa class I heat shock protein [Vitis vinifera]
416	PB.60028.1	PREDICTED: 22.0 kDa class IV heat shock protein [Ziziphus jujuba]
417	PB.57873.1	PREDICTED: 17.4 kDa class III heat shock protein-like [Pyrus x bretschneideri]
418	PB.58069.3	PREDICTED: 17.1 kDa class II heat shock protein-like [Juglans regia]
419	PB.42914.1	PREDICTED: 18.1 kDa class I heat shock protein [Vitis vinifera]
420	PB.34002.6	PREDICTED: 18.1 kDa class I heat shock protein [Vitis vinifera]
421	PB.30756.1	PREDICTED: stromal 70 kDa heat shock-related protein, chloroplastic [Vitis vinifera]
422	PB.25386.1	PREDICTED: stromal 70 kDa heat shock-related protein, chloroplastic [Populus euphratica]
423	PB.63825.1	PREDICTED: 17.4 kDa class III heat shock protein-like [Pyrus x bretschneideri]
424	PB.64717.1	PREDICTED: 17.4 kDa class III heat shock protein [Nicotiana tomentosiformis]
425	PB.64424.1	PREDICTED: 17.8 kDa class I heat shock protein-like [Cucumis melo]
426	PB.63931.1	PREDICTED: 17.1 kDa class II heat shock protein-like [Juglans regia]
427	PB.63208.4	PREDICTED: 18.1 kDa class I heat shock protein [Vitis vinifera]
428	PB.61931.3	PREDICTED: 17.4 kDa class III heat shock protein-like [Pyrus x bretschneideri]

429	PB.64660.2	17.8 kDa class I heat shock protein-like [Cucurbita maxima]
430	PB.64391.2	PREDICTED: 17.4 kDa class III heat shock protein [Vitis vinifera]
431	PB.64795.1	17.8 kDa class I heat shock protein-like isoform X1 [Momordica charantia]
432	PB.13232.2	17.4 kDa class III heat shock protein [Glycine soja]
433	PB.28818.1	PREDICTED: stromal 70 kDa heat shock-related protein, chloroplastic [Vitis vinifera]
434	PB.10662.2	PREDICTED: stromal 70 kDa heat shock-related protein, chloroplastic [Vitis vinifera]
435	PB.736.21	PREDICTED: stromal 70 kDa heat shock-related protein, chloroplastic [Vitis vinifera]
436	PB.64165.1	PREDICTED: 26.5 kDa heat shock protein, mitochondrial-like [Nicotiana sylvestris]
437	PB.64975.1	PREDICTED: 17.4 kDa class III heat shock protein-like [Pyrus x bretschneideri]
438	PB.65141.1	PREDICTED: 18.1 kDa class I heat shock protein-like [Cucumis melo]
439	PB.66066.1	PREDICTED: 18.1 kDa class I heat shock protein-like [Pyrus x bretschneideri]
440	PB.53615.4	PREDICTED: small heat shock protein, chloroplastic-like [Prunus mume]
441	PB.1041.24	PREDICTED: stromal 70 kDa heat shock-related protein, chloroplastic [Vitis vinifera]
442	PB.1006.14	PREDICTED: stromal 70 kDa heat shock-related protein, chloroplastic [Cucumis melo]
443	PB.54679.1	Small heat shock protein isoform 2 [Theobroma cacao]
444	PB.63130.2	17.3 kDa class I heat shock protein-like [Momordica charantia]
445	PB.65636.1	23.6 kDa heat shock protein, mitochondrial [Herrania umbratica]
446	PB.49929.1	Heat shock protein Hsp90, partial [Cynara cardunculus var. scolymus]
447	PB.31368.1	Heat shock cognate protein 70-1 [Theobroma cacao]
448	PB.30998.3	Heat shock cognate 70 kDa protein [Cajanus cajan]
449	PB.66136.1	heat shock factor-binding protein 1-like isoform X2 [Manihot esculenta]
450	PB.45204.1	Heat shock transcription factor A2 isoform 1 [Theobroma cacao]
451	PB.33030.1	Heat shock protein Hsp90, partial [Cynara cardunculus var. scolymus]
452	PB.30233.1	heat shock protein 90-5, chloroplastic-like isoform X1 [Durio zibethinus]
453	PB.31254.3	PREDICTED: heat shock cognate 70 kDa protein 2 [Theobroma cacao]
454	PB.26510.1	heat shock cognate protein 80 [Herrania umbratica]
455	PB.31862.1	Heat shock cognate protein 70-1 isoform 1 [Theobroma cacao]

456	PB.854.2	Heat shock protein Hsp90 [Corchorus olitorius]
457	PB.2103.2	heat shock cognate protein 80 [Durio zibethinus]
458	PB.23865.1	heat shock cognate 70 kDa protein 2-like [Olea europaea var. sylvestris]
459	PB.7600.1	heat shock cognate 70 kDa protein 2-like [Olea europaea var. sylvestris]
460	PB.17754.1	Heat shock protein 70 family [Corchorus olitorius]
461	PB.66203.1	Heat shock cognate 70 kDa protein 1 [Aegilops tauschii]
462	PB.62225.1	Heat shock protein Hsp90 [Cynara cardunculus var. scolymus]
463	PB.25437.3	DNAJ heat shock N-terminal domain-containing family protein [Populus trichocarpa]
464	PB.47601.1	heat stress transcription factor A-1b-like [Durio zibethinus]
465	PB.19624.1	PREDICTED: heat shock factor protein HSF8-like [Lupinus angustifolius]
466	PB.57559.1	heat shock protein 90, partial [Paeonia lactiflora]
467	PB.60692.1	heat shock protein 90-1-like protein, partial [Betula luminifera]
468	PB.60261.1	heat shock protein 60, partial [Paeonia lactiflora]
469	PB.47637.2	heat shock protein 70 [Paeonia suffruticosa]
470	PB.43833.1	heat shock protein 90, partial [Paeonia lactiflora]
471	PB.38017.1	heat shock protein 70 [Cyclamen persicum]
472	PB.39802.1	heat shock protein cognate 70 [Oryza sativa Japonica Group]
473	PB.38035.1	heat shock cognate protein 80 [Herrania umbratica]
474	PB.38603.1	Heat shock protein 90-5, chloroplastic [Ananas comosus]
475	PB.34948.1	heat shock cognate 70 kDa protein [Medicago truncatula]
476	PB.34613.1	heat shock protein [Picea mariana]
477	PB.37078.1	heat shock protein 70 family protein [Klebsormidium flaccidum]
478	PB.30626.1	heat shock protein 83 [Quercus suber]
479	PB.26045.1	heat shock protein 70 [Paeonia lactiflora]
480	PB.24980.1	heat shock protein 70 [Paeonia lactiflora]
481	PB.63663.1	heat shock protein 90, partial [Capparis spinosa]
482	PB.23135.10	heat shock cognate 70 kDa protein 2-like protein [Solanum melongena]

483		PB.22220.30	heat shock protein 90 [Lactuca sativa]
484		PB.24845.1	heat shock protein 70 [Paeonia lactiflora]
485		PB.19672.1	PREDICTED: heat shock cognate protein 80 [Ipomoea nil]
486		PB.19781.1	heat shock protein 70 [Paeonia lactiflora]
487		PB.10300.1	heat shock protein 70 [Gossypium hirsutum]
488		PB.11638.1	heat shock protein 70 [Paeonia lactiflora]
489		PB.15296.13	heat shock protein 70 [Chlorella variabilis]
490		PB.1160.1	heat shock protein 70 [Paeonia lactiflora]
491		PB.2774.1	heat shock protein 90, partial [Paeonia lactiflora]
492		PB.313.71	heat shock protein 70 [Paeonia lactiflora]
493		PB.65479.1	heat shock protein 17.8 [Oryza sativa Japonica Group]
494		PB.31563.1	PREDICTED: heat shock cognate 70 kDa protein 1-like [Nicotiana attenuata]
495		PB.43678.1	PREDICTED: heat stress transcription factor A-4a [Vitis vinifera]
496	NAC	PB.54542.3	PREDICTED: NAC domain-containing protein 72-like [Juglans regia]
497		PB.48215.4	NAC domain-containing protein 83-like isoform X1 [Hevea brasiliensis]
498		PB.10529.1	NAC transcription factors 88 [Manihot esculenta]
499		PB.16045.3	NAC transcription factors 88 [Manihot esculenta]
500		PB.35537.2	NAC domain-containing protein 78-like [Hevea brasiliensis]
501		PB.56895.1	NAC domain-containing protein [Chrysanthemum x morifolium]
502		PB.45142.1	PREDICTED: NAC domain-containing protein 8 isoform X1 [Theobroma cacao]
503		PB.50066.2	NAC domain-containing 72 -like protein [Gossypium arboreum]
504		PB.40347.5	PREDICTED: NAC domain-containing protein 8 isoform X1 [Theobroma cacao]
505		PB.15658.1	PREDICTED: NAC domain-containing protein 17-like [Gossypium hirsutum]
506		PB.5970.1	NAC domain-containing protein [Boehmeria nivea]
507		PB.52641.1	PREDICTED: NAC domain-containing protein 78 [Vitis vinifera]
508		PB.27448.1	NAC domain-containing protein 62-like [Carica papaya]
509		PB.53716.1	PREDICTED: NAC domain-containing protein 21/22 [Vitis vinifera]

510	PB.52085.1	PREDICTED: NAC domain-containing protein 30 isoform X2 [Vitis vinifera]
511	PB.62951.1	PREDICTED: NAC domain-containing protein 83 [Vitis vinifera]
512	PB.53226.1	PREDICTED: NAC domain-containing protein 83 [Ricinus communis]
513	PB.55663.1	PREDICTED: NAC domain-containing protein 2-like [Beta vulgaris subsp. vulgaris]
514	PB.56100.1	PREDICTED: NAC domain-containing protein 73 [Citrus sinensis]
515	PB.51766.1	NAC domain-containing protein 53 isoform X1 [Prunus avium]
516	PB.54070.1	PREDICTED: NAC domain-containing protein 2 [Ziziphus jujuba]
517	PB.51862.2	PREDICTED: NAC domain-containing protein 8 isoform X1 [Ziziphus jujuba]
518	PB.49051.4	PREDICTED: NAC domain-containing protein 72 isoform X1 [Vitis vinifera]
519	PB.47777.1	NAC domain containing protein 50-like [Olea europaea var. sylvestris]
520	PB.43761.6	PREDICTED: NAC domain-containing protein 62-like isoform X1 [Juglans regia]
521	PB.45240.1	PREDICTED: NAC domain-containing protein 86 [Vitis vinifera]
522	PB.38926.1	PREDICTED: NAC domain-containing protein 8 isoform X2 [Vitis vinifera]
523	PB.41351.2	PREDICTED: NAC domain-containing protein 8 isoform X1 [Ziziphus jujuba]
524	PB.37366.1	PREDICTED: NAC domain-containing protein 82 [Vitis vinifera]
525	PB.36005.6	PREDICTED: NAC domain-containing protein 82 [Vitis vinifera]
526	PB.28598.3	PREDICTED: NAC domain-containing protein 82 [Vitis vinifera]
527	PB.27224.1	PREDICTED: NAC domain-containing protein 82 [Vitis vinifera]
528	PB.26620.2	PREDICTED: NAC domain-containing protein 82 [Vitis vinifera]
529	PB.7861.1	PREDICTED: NAC domain-containing protein 78-like [Nicotiana tomentosiformis]
530	PB.800.4	PREDICTED: NAC domain-containing protein 53 [Vitis vinifera]
531	PB.64178.1	PREDICTED: NAC domain-containing protein 67-like [Vitis vinifera]
532	PB.60139.4	PREDICTED: basic transcription factor 3 [Eucalyptus grandis]
533	PB.39017.3	PREDICTED: transcription factor VOZ1 [Nelumbo nucifera]
534	PB.42253.1	PREDICTED: transcription factor VOZ1 [Vitis vinifera]
535	PB.28186.1	PREDICTED: transcription factor VOZ1 isoform X2 [Ricinus communis]
536	PB.25251.6	PREDICTED: transcription factor VOZ1-like isoform X3 [Sesamum indicum]

537	Nuclear TF Y	PB.63798.1	Nuclear transcription factor Y subunit B-8, partial [Glycine soja]
538		PB.55866.1	Nuclear transcription factor Y subunit B-8, partial [Glycine soja]
539		PB.55193.1	PREDICTED: nuclear transcription factor Y subunit C-9-like isoform X1 [Vitis vinifera]
540		PB.58127.1	PREDICTED: nuclear transcription factor Y subunit B-1-like isoform X1 [Vitis vinifera]
541		PB.51789.3	PREDICTED: nuclear transcription factor Y subunit A-7 [Vitis vinifera]
542		PB.54671.2	PREDICTED: nuclear transcription factor Y subunit C-9-like isoform X1 [Vitis vinifera]
543		PB.59218.2	PREDICTED: nuclear transcription factor Y subunit B-3 [Jatropha curcas]
544		PB.53495.1	PREDICTED: nuclear transcription factor Y subunit C-1 [Cucumis melo]
545		PB.51415.1	PREDICTED: nuclear transcription factor Y subunit C-2 [Ziziphus jujuba]
546		PB.48540.3	PREDICTED: nuclear transcription factor Y subunit A-1 [Vitis vinifera]
547		PB.48987.2	PREDICTED: nuclear transcription factor Y subunit B-10 [Beta vulgaris subsp. vulgaris]
548		PB.40592.8	PREDICTED: nuclear transcription factor Y subunit B-1-like isoform X1 [Vitis vinifera]
549		PB.27755.3	PREDICTED: nuclear transcription factor Y subunit A-3-like isoform X1 [Malus domestica]
550		PB.20229.1	PREDICTED: nuclear transcription factor Y subunit C-4-like [Gossypium raimondii]
551		PB.62215.1	PREDICTED: nuclear transcription factor Y subunit B-3-like [Malus domestica]
552	YABBY	PB.57254.2	PREDICTED: axial regulator YABBY 1 [Theobroma cacao]，
553		PB.57368.1	PREDICTED: axial regulator YABBY 5 isoform X1 [Vitis vinifera]
554		PB.52416.1	PREDICTED: protein YABBY 4 [Vitis vinifera]
555		PB.49311.9	PREDICTED: protein YABBY 4 [Vitis vinifera]
556		PB.4027.3	Protein YABBY 6, partial [Capsicum baccatum]
557		PB.64261.1	PREDICTED: protein YABBY 4 [Vitis vinifera]
558		PB.64312.1	PREDICTED: protein YABBY 4 [Vitis vinifera]
559	B3	PB.33333.5	PREDICTED: B3 domain-containing transcription factor ABI3 isoform X1 [Vitis vinifera]
560		PB.59284.1	PREDICTED: B3 domain-containing transcription factor VRN1-like [Fragaria vesca subsp. vesca]
561		PB.56751.1	PREDICTED: B3 domain-containing protein REM8-like [Jatropha curcas]
562		PB.56233.1	PREDICTED: B3 domain-containing protein REM20-like, partial [Vitis vinifera]
563		PB.49963.1	PREDICTED: B3 domain-containing protein REM20-like, partial [Vitis vinifera]

564	PB.50653.1	B3 domain-containing protein REM14-like isoform X4 [Manihot esculenta]
565	PB.48032.1	PREDICTED: B3 domain-containing protein REM20-like, partial [Vitis vinifera]
566	PB.42450.1	PREDICTED: B3 domain-containing protein REM10-like [Nicotiana sylvestris]
567	PB.35930.1	PREDICTED: expansin-B3-like [Nelumbo nucifera]
568	PB.51851.1	PREDICTED: expansin-B3-like [Nelumbo nucifera]
569	PB.46753.2	PREDICTED: B3 domain-containing protein Os01g0723500-like [Vitis vinifera]
570	PB.10969.1	PREDICTED: B3 domain-containing transcription repressor VAL1-like [Ziziphus jujuba]
571	PB.5082.2	B3 DNA binding domain containing protein [Parasponia andersonii]
572	PB.60058.1	PREDICTED: B3 domain-containing protein At2g33720-like [Populus euphratica]
573	PB.58726.2	PREDICTED: B3 domain-containing protein At3g19184 isoform X1 [Vitis vinifera]
574	PB.60526.1	PREDICTED: B3 domain-containing protein At2g33720-like [Nelumbo nucifera]
575	PB.52206.1	PREDICTED: B3 domain-containing protein Os01g0723500-like [Vitis vinifera]
576	PB.54752.2	PREDICTED: B3 domain-containing protein At2g36080-like [Vitis vinifera]
577	PB.50567.1	PREDICTED: B3 domain-containing protein Os01g0723500-like [Vitis vinifera]
578	PB.49339.1	PREDICTED: B3 domain-containing protein At3g19184 isoform X1 [Vitis vinifera]
579	PB.43814.1	PREDICTED: B3 domain-containing protein Os01g0234100-like [Prunus mume]
580	PB.45261.1	PREDICTED: B3 domain-containing protein At3g19184 [Solanum tuberosum]
581	PB.43556.3	PREDICTED: B3 domain-containing protein Os01g0234100-like [Jatropha curcas]
582	PB.42155.1	PREDICTED: B3 domain-containing protein Os03g0120900-like [Vitis vinifera]
583	PB.37256.1	B3 domain-containing protein/DUF724 domain-containing protein [Cephalotus follicularis]
584	PB.42137.2	PREDICTED: B3 domain-containing protein Os03g0120900-like [Malus domestica]
585	PB.29685.1	PREDICTED: B3 domain-containing transcription repressor VAL1-like [Ziziphus jujuba]
586	PB.12409.2	B3 DNA binding domain containing protein [Parasponia andersonii]
587	PB.12352.1	PREDICTED: B3 domain-containing transcription repressor VAL1-like [Populus euphratica]
588	PB.14758.1	B3 domain-containing protein Os07g0679700 [Herrania umbratica]
589	PB.12104.2	PREDICTED: B3 domain-containing protein Os01g0723500-like [Vitis vinifera]
590	PB.9496.3	b3 domain-containing protein [Quercus suber]

591		PB.9233.3	PREDICTED: B3 domain-containing transcription repressor VAL1-like [Ziziphus jujuba]
592		PB.6705.1	PREDICTED: B3 domain-containing protein At5g42700 [Vitis vinifera]
593		PB.37740.1	PREDICTED: LOW QUALITY PROTEIN: B3 domain-containing protein Os07g0563300-like [Cucumis melo]
594		PB.31902.1	B3 domain-containing protein Os07g0563300-like isoform X2 [Hevea brasiliensis]
595		PB.14299.1	B3 domain-containing protein [Morus notabilis]
596		PB.10539.24	B3 domain-containing protein Os07g0563300-like isoform X2 [Hevea brasiliensis]
597		PB.12637.1	B3 domain-containing protein Os07g0563300-like isoform X2 [Hevea brasiliensis]
598		PB.6470.1	Transcriptional factor B3 family protein / auxin-responsive factor AUX/IAA-related isoform 1 [Theobroma cacao]
599	BZIP	PB.49578.3	PREDICTED: bZIP transcription factor 53 [Ricinus communis]
600		PB.60155.1	PREDICTED: bZIP transcription factor TRAB1-like isoform X2 [Ziziphus jujuba]
601		PB.52867.3	PREDICTED: bZIP transcription factor 11-like [Juglans regia]
602		PB.48185.3	bZIP transcription factor bZIP7 [Camellia sinensis]
603		PB.48852.3	bZIP transcription factor 60 [Jatropha curcas]
604		PB.44778.7	bZIP transcription factor family protein 2 [Camellia sinensis]
605		PB.44308.2	bZIP transcription factor bZIP7 [Camellia sinensis]
606		PB.40196.1	bZIP transcription factor bZIP7 [Camellia sinensis]
607		PB.37211.1	bZIP transcription factor bZIP7 [Camellia sinensis]
608		PB.35331.4	bZIP transcription factor bZIP8 [Camellia sinensis]
609		PB.19310.2	bZIP transcription factor bZIP7 [Camellia sinensis]
610		PB.4262.8	bZIP transcription factor bZIP8 [Camellia sinensis]
611		PB.51852.1	PREDICTED: basic leucine zipper and W2 domain-containing protein 2-like [Elaeis guineensis]
612		PB.60821.1	PREDICTED: basic leucine zipper 9 [Vitis vinifera]
613		PB.42554.1	PREDICTED: basic leucine zipper 61-like [Nelumbo nucifera]
614		PB.28368.4	PREDICTED: probable basic-leucine zipper transcription factor F isoform X1 [Ziziphus jujuba]
615		PB.66239.1	PREDICTED: transcription factor HY5 [Ricinus communis]
616		PB.13000.1	PREDICTED: transcription factor HY5-like [Ricinus communis]
617		PB.11815.1	PREDICTED: transcription factor HY5-like [Ricinus communis]

618		PB.63493.1	PREDICTED: transcription factor HY5-like [Ricinus communis]
619		PB.61919.1	PREDICTED: transcription factor TGA6-like [Juglans regia]
620		PB.58209.1	PREDICTED: transcription factor TGA2.2 isoform X1 [Vitis vinifera]
621		PB.48850.1	PREDICTED: transcription factor TGA2.2 isoform X1 [Vitis vinifera]
622		PB.419.8	PREDICTED: transcription factor TGA2.2 isoform X1 [Vitis vinifera]
623		PB.27534.1	PREDICTED: transcription factor TGA2.2 isoform X1 [Vitis vinifera]
624		PB.41773.1	PREDICTED: transcription factor TGA2.2 isoform X1 [Vitis vinifera]
625		PB.37460.1	PREDICTED: transcription factor TGA2.2 isoform X1 [Vitis vinifera]
626		PB.29970.1	PREDICTED: transcription factor TGA2.2 isoform X1 [Vitis vinifera]
627		PB.29920.1	PREDICTED: transcription factor TGA2.2 isoform X1 [Vitis vinifera]
628		PB.47997.1	PREDICTED: probable transcription factor PosF21 [Cicer arietinum]
629		PB.65076.1	PREDICTED: probable transcription factor PosF21 isoform X1 [Nelumbo nucifera]
630		PB.63039.1	PREDICTED: transcription factor TGA2-like [Populus euphratica]
631		PB.33186.4	PREDICTED: transcription factor TGA2.3 [Vitis vinifera]
632		PB.53403.1	PREDICTED: transcription factor TGA2-like [Ziziphus jujuba]
633		PB.12479.1	PREDICTED: transcription factor TGA2-like [Ziziphus jujuba]
634		PB.59367.1	bZIP_1 domain-containing protein/DOG1 domain-containing protein [Cephalotus follicularis]
635		PB.11347.4	PREDICTED: transcription factor TGA4 [Vitis vinifera]
636		PB.36647.2	Transcription factor TGA like domain [Macleaya cordata]
637		PB.43180.4	PREDICTED: transcription factor TGA2.3-like [Nelumbo nucifera]
638		PB.53233.2	PREDICTED: transcription factor TGA5-like [Ziziphus jujuba]
639		PB.60731.1	PREDICTED: transcription factor TGA6-like isoform X2 [Ziziphus jujuba]
640		PB.62707.1	TGA transcription factor [Trema orientalis]
641	E2F	PB.47255.4	PREDICTED: E2F transcription factor-like E2FE isoform X1 [Vitis vinifera]
642		PB.46529.2	PREDICTED: transcription factor E2FC isoform X1 [Vitis vinifera]
643		PB.45642.4	PREDICTED: transcription factor E2FA [Nelumbo nucifera]
644		PB.34649.2	PREDICTED: transcription factor E2FA isoform X2 [Vitis vinifera]

645		PB.65380.1	PREDICTED: E2F-associated phosphoprotein [Vitis vinifera]
646	GATA	PB.61728.1	GATA transcription factor 8-like [Durio zibethinus]
647		PB.51687.1	PREDICTED: GATA transcription factor 12 [Nelumbo nucifera]
648		PB.52574.1	PREDICTED: GATA transcription factor 12-like [Nelumbo nucifera]
649		PB.58132.1	PREDICTED: GATA transcription factor 15 [Malus domestica]
650		PB.62654.1	PREDICTED: GATA transcription factor 15-like [Vitis vinifera]
651		PB.59474.1	PREDICTED: GATA transcription factor 16 [Vitis vinifera]
652		PB.51661.1	PREDICTED: GATA transcription factor 16-like [Nelumbo nucifera]
653		PB.61870.2	GATA transcription factor 15-like [Durio zibethinus]
654		PB.39506.1	PREDICTED: GATA transcription factor 21 isoform X2 [Vigna radiata var. radiata]
655		PB.51383.1	PREDICTED: GATA transcription factor 21 isoform X2 [Vitis vinifera]
656		PB.51808.1	PREDICTED: GATA transcription factor 24 [Vitis vinifera]
657		PB.46835.1	PREDICTED: GATA transcription factor 24 [Vitis vinifera]
658		PB.47228.7	PREDICTED: GATA transcription factor 24-like isoform X2 [Vitis vinifera]
659		PB.25862.1	PREDICTED: GATA transcription factor 24-like isoform X2 [Vitis vinifera]
660		PB.11646.11	PREDICTED: GATA transcription factor 25 [Vitis vinifera]
661		PB.66326.1	gata transcription factor 5 [Quercus suber]
662		PB.51634.2	PREDICTED: GATA transcription factor 5 [Vitis vinifera]
663		PB.48336.1	PREDICTED: GATA transcription factor 5 [Vitis vinifera]
664		PB.47826.2	PREDICTED: GATA transcription factor 5-like [Populus euphratica]
665		PB.57979.1	GATA transcription factor [Trema orientalis]
666		PB.51400.1	GATA transcription factor [Parasponia andersonii]
667		PB.37168.2	PREDICTED: GATA transcription factor 7-like [Vitis vinifera]
668		PB.45875.1	PREDICTED: GATA transcription factor 8-like [Vitis vinifera]
669		PB.52258.1	PREDICTED: GATA transcription factor 9 [Vitis vinifera]
670		PB.52240.1	GATA transcription factor 5-like [Malus domestica]
671	Membrane-Bound	PB.35288.5	PREDICTED: membrane-bound transcription factor site-2 protease homolog isoform X1 [Jatropha curcas]

672		PB.37755.1	PREDICTED: membrane-bound transcription factor site-2 protease homolog isoform X2 [Ricinus communis]
673		PB.43585.1	PREDICTED: membrane-bound transcription factor site-2 protease isoform X1 [Nelumbo nucifera]
674	RING	PB.12206.1	putative E3 ubiquitin-protein ligase RING1a [Durio zibethinus]
675		PB.52745.1	PREDICTED: putative E3 ubiquitin-protein ligase RING1a [Vitis vinifera]
676		PB.43809.1	PREDICTED: putative E3 ubiquitin-protein ligase RING1a isoform X1 [Ziziphus jujuba]
677		PB.34078.1	PREDICTED: putative E3 ubiquitin-protein ligase RING1a isoform X1 [Ziziphus jujuba]
678		PB.59966.1	PREDICTED: E3 ubiquitin-protein ligase RING1-like [Cucumis melo]
679	WRKY	PB.55374.1	PREDICTED: probable WRKY transcription factor 13 [Beta vulgaris subsp. vulgaris]
680		PB.7710.1	DNA-binding WRKY [Corchorus olitorius]
681		PB.17256.4	PREDICTED: probable WRKY transcription factor 2 [Vitis vinifera]
682		PB.5613.1	PREDICTED: probable WRKY transcription factor 2 [Vitis vinifera]
683		PB.39959.2	PREDICTED: probable WRKY transcription factor 20 [Nelumbo nucifera]
684		PB.31133.4	DNA-binding WRKY [Macleaya cordata]
685		PB.19800.1	PREDICTED: probable WRKY transcription factor 20 [Prunus mume]
686		PB.48211.1	PREDICTED: probable WRKY transcription factor 20 [Prunus mume]
687		PB.21154.1	PREDICTED: probable WRKY transcription factor 20 [Prunus mume]
688		PB.17147.2	PREDICTED: probable WRKY transcription factor 20 [Prunus mume]
689		PB.44475.4	PREDICTED: probable WRKY transcription factor 20 isoform X1 [Vitis vinifera]
690		PB.28010.1	PREDICTED: probable WRKY transcription factor 20 isoform X1 [Vitis vinifera]
691		PB.28522.2	PREDICTED: probable WRKY transcription factor 20 isoform X1 [Vitis vinifera]
692		PB.5093.2	PREDICTED: probable WRKY transcription factor 20 isoform X1 [Vitis vinifera]
693		PB.32099.6	PREDICTED: probable WRKY transcription factor 21 [Prunus mume]
694		PB.56690.1	PREDICTED: probable WRKY transcription factor 29 isoform X2 [Pyrus x bretschneideri]
695		PB.48767.1	PREDICTED: probable WRKY transcription factor 31 [Vitis vinifera]
696		PB.43811.1	PREDICTED: probable WRKY transcription factor 32 isoform X2 [Vitis vinifera]
697		PB.8236.3	PREDICTED: probable WRKY transcription factor 32 isoform X2 [Vitis vinifera]
698		PB.53405.1	PREDICTED: probable WRKY transcription factor 48 [Populus euphratica]

699	PB.57400.1	probable WRKY transcription factor 57 [Olea europaea var. sylvestris]
700	PB.54824.1	PREDICTED: probable WRKY transcription factor 65 [Vitis vinifera]
701	PB.54448.2	PREDICTED: probable WRKY transcription factor 65 [Vitis vinifera]
702	PB.52477.6	DNA-binding WRKY [Macleaya cordata]
703	PB.48811.5	PREDICTED: probable WRKY transcription factor 7 [Nicotiana tomentosiformis]
704	PB.35782.4	PREDICTED: probable WRKY transcription factor 74 [Vitis vinifera]
705	PB.6758.1	PREDICTED: probable WRKY transcription factor 74 [Vitis vinifera]
706	PB.41795.1	PREDICTED: probable WRKY transcription factor 33 [Daucus carota subsp. sativus]
707	PB.14083.1	PREDICTED: probable WRKY transcription factor 32 isoform X3 [Vitis vinifera]
708	PB.28248.1	probable WRKY transcription factor 13 [Manihot esculenta]
709	PB.46613.2	PREDICTED: probable WRKY transcription factor 7 [Vitis vinifera]
710	PB.49343.1	WRKY transcription factor 15 family protein [Populus trichocarpa]
711	PB.60580.1	probable WRKY transcription factor 23 [Manihot esculenta]
712	PB.28626.1	WRKY transcription factor 2 [Osmanthus fragrans]
713	PB.51998.2	WRKY transcription factor 2 [Solanum tuberosum]
714	PB.31707.1	WRKY transcription factor 29-4 [Dimocarpus longan]
715	PB.52783.1	WRKY transcription factor 22 [Manihot esculenta]
716	PB.53629.1	probable WRKY transcription factor 13 [Manihot esculenta]
717	PB.54972.1	probable WRKY transcription factor 13 [Jatropha curcas]
718	PB.31608.2	WRKY13 transcription factor [Vitis amurensis]
719	PB.50498.1	WRKY3 transcription factor [Vitis pseudoreticulata]
720	PB.31270.1	PREDICTED: WRKY transcription factor 44 [Vitis vinifera]
721	PB.18733.6	PREDICTED: WRKY transcription factor 44 [Vitis vinifera]
722	PB.32353.1	PREDICTED: WRKY transcription factor SUSIBA2-like [Ziziphus jujuba]
723	PB.5438.1	PREDICTED: WRKY transcription factor SUSIBA2-like [Ziziphus jujuba]
724	PB.43929.1	DNA binding protein WRKY2 [Vitis vinifera]
725	PB.39483.1	DNA binding protein WRKY2 [Vitis vinifera]

726		PB.26559.3	DNA binding protein WRKY2 [Vitis vinifera]
727	Trihelix	PB.57063.1	PREDICTED: trihelix transcription factor ASIL1 [Gossypium raimondii]
728		PB.47605.4	PREDICTED: trihelix transcription factor ASIL1 [Ricinus communis]
729		PB.24652.4	PREDICTED: trihelix transcription factor ASIL1-like [Jatropha curcas]
730		PB.35632.1	PREDICTED: trihelix transcription factor ASIL2-like [Juglans regia]
731		PB.53565.1	PREDICTED: trihelix transcription factor ASIL1-like [Vitis vinifera]
732		PB.13372.1	PREDICTED: trihelix transcription factor ASIL1-like [Vitis vinifera]
733		PB.49561.7	trihelix transcription factor ASIL2-like isoform X3 [Carica papaya]
734		PB.38208.6	PREDICTED: trihelix transcription factor ASIL1-like [Gossypium hirsutum]
735		PB.58798.1	PREDICTED: trihelix transcription factor ASIL2 [Nelumbo nucifera]
736		PB.47534.1	PREDICTED: trihelix transcription factor ASIL2 [Vitis vinifera]
737		PB.57826.1	PREDICTED: trihelix transcription factor ASIL2 [Ziziphus jujuba]
738		PB.48419.1	PREDICTED: trihelix transcription factor ASIL2-like [Vitis vinifera]
739		PB.48212.5	PREDICTED: trihelix transcription factor ASIL2-like [Vitis vinifera]
740		PB.49009.1	PREDICTED: trihelix transcription factor ASIL2-like [Vitis vinifera]
741		PB.63479.1	PREDICTED: trihelix transcription factor ASIL2-like [Vitis vinifera]
742		PB.13922.2	PREDICTED: trihelix transcription factor GT-1 [Nelumbo nucifera]
743		PB.49514.1	PREDICTED: trihelix transcription factor GT-1-like [Nelumbo nucifera]
744		PB.45699.1	PREDICTED: trihelix transcription factor GT-1-like [Nelumbo nucifera]
745		PB.46215.2	PREDICTED: trihelix transcription factor GT-1-like [Nelumbo nucifera]
746		PB.39157.1	PREDICTED: trihelix transcription factor GT-1-like [Phoenix dactylifera]
747		PB.45858.2	PREDICTED: trihelix transcription factor GT-2 [Cucumis sativus]
748		PB.57616.1	PREDICTED: trihelix transcription factor ASR3-like [Nelumbo nucifera]
749		PB.30312.1	PREDICTED: trihelix transcription factor GT-2-like [Nelumbo nucifera]
750		PB.63009.1	PREDICTED: trihelix transcription factor GT-3b-like [Nelumbo nucifera]
751		PB.59782.1	PREDICTED: trihelix transcription factor GT-3b-like [Nicotiana sylvestris]
752		PB.558.22	PREDICTED: trihelix transcription factor GTL1 isoform X1 [Vitis vinifera]

753		PB.35178.2	PREDICTED: trihelix transcription factor GTL2 [Vitis vinifera]
754		PB.42052.1	PREDICTED: trihelix transcription factor GTL2 [Vitis vinifera]
755		PB.43063.1	PREDICTED: trihelix transcription factor PTL-like [Vitis vinifera]
756	C2H2	PB.54877.3	C2H2-type zinc finger protein [Medicago truncatula]
757		PB.55520.1	Zinc finger, C2H2 [Cynara cardunculus var. scolymus]
758		PB.44073.2	C2H2-like zinc finger protein [Theobroma cacao]
759		PB.40650.4	C2H2-like zinc finger protein [Medicago truncatula]
760	WD	PB.50194.2	PREDICTED: WD repeat-containing protein 82 [Jatropha curcas]
761		PB.53789.1	PREDICTED: COMPASS-like H3K4 histone methylase component WDR5A [Vitis vinifera]
762		PB.50689.3	PREDICTED: WD repeat-containing protein VIP3 [Nelumbo nucifera]
763		PB.47780.1	PREDICTED: WD repeat-containing protein 74 isoform X1 [Vitis vinifera]
764		PB.47781.1	WD40 protein [Paeonia suffruticosa]
765		PB.42095.1	Transducin/WD40 repeat-like superfamily protein isoform 1 [Theobroma cacao]
766		PB.28569.2	F-box/WD-40 repeat-containing protein At5g21040 family [Cajanus cajan]
767		PB.28178.3	PREDICTED: WD repeat-containing protein 26-like [Prunus mume]
768		PB.24729.1	PREDICTED: WD repeat-containing protein 48 isoform X1 [Jatropha curcas]
769		PB.22654.3	PREDICTED: WD and tetratricopeptide repeats protein 1 isoform X1 [Vitis vinifera]
770		PB.18100.1	WD repeat-containing protein 75 [Manihot esculenta]
771		PB.20107.2	PREDICTED: WD repeat and HMG-box DNA-binding protein 1 isoform X2 [Jatropha curcas]
772		PB.18480.1	PREDICTED: katanin p80 WD40 repeat-containing subunit B1 homolog isoform X2 [Vitis vinifera]
773		PB.11301.4	PREDICTED: WD repeat-containing protein YMR102C [Vitis vinifera]
774		PB.59915.1	Transducin/WD40 repeat-like superfamily protein [Theobroma cacao]
775		PB.58526.1	Transducin family protein / WD-40 repeat family protein isoform 2 [Theobroma cacao]
776		PB.59918.1	PREDICTED: WD repeat-containing protein 55 homolog isoform X2 [Sesamum indicum]
777		PB.58473.1	PREDICTED: WD repeat-containing protein 25 [Erythranthe guttata]
778		PB.63586.1	WD repeat-containing protein 11 [Helianthus annuus]
779		PB.58906.1	WD-40 repeat protein/beige protein [Medicago truncatula]

780	PB.56609.1	PREDICTED: WD repeat-containing protein 89 homolog [Prunus mume]
781	PB.57613.1	PREDICTED: WD repeat-containing protein 55 homolog isoform X2 [Sesamum indicum]
782	PB.59502.1	PREDICTED: F-box/WD-40 repeat-containing protein At3g52030-like [Jatropha curcas]
783	PB.52892.1	PREDICTED: WD repeat-containing protein 76 isoform X1 [Jatropha curcas]
784	PB.49894.2	PREDICTED: WD repeat-containing protein 74 isoform X1 [Vitis vinifera]
785	PB.51821.1	PREDICTED: COMPASS-like H3K4 histone methylase component WDR5A [Vitis vinifera]
786	PB.50274.1	PREDICTED: ribosome biogenesis protein WDR12 homolog [Nicotiana tomentosiformis]
787	PB.48908.4	PREDICTED: WD repeat-containing protein DWA2 isoform X1 [Vitis vinifera]
788	PB.50189.1	PREDICTED: WD repeat-containing protein 89 homolog [Vitis vinifera]
789	PB.54754.1	PREDICTED: WD repeat-containing protein tag-125 [Vitis vinifera]
790	PB.53898.1	PREDICTED: WD repeat-containing protein VIP3 [Nelumbo nucifera]
791	PB.46379.1	PREDICTED: COMPASS-like H3K4 histone methylase component WDR5A [Lupinus angustifolius]
792	PB.47376.1	PREDICTED: WD repeat-containing protein 53 [Ziziphus jujuba]
793	PB.45714.1	wd repeat-containing protein lwd1 [Quercus suber]
794	PB.46069.1	PREDICTED: WD-40 repeat-containing protein MSI3 [Vitis vinifera]
795	PB.46023.1	PREDICTED: dystrophia myotonica WD repeat-containing protein isoform X2 [Jatropha curcas]
796	PB.48040.1	PREDICTED: COMPASS-like H3K4 histone methylase component WDR5B [Nelumbo nucifera]
797	PB.46107.3	PREDICTED: WD repeat domain-containing protein 83 [Nelumbo nucifera]
798	PB.46733.1	PREDICTED: WD repeat-containing protein 74-like [Eucalyptus grandis]
799	PB.45962.1	PREDICTED: COMPASS-like H3K4 histone methylase component WDR5B [Nelumbo nucifera]
800	PB.43719.1	Katanin p80 WD40 repeat-containing subunit B1 like 1 [Glycine soja]
801	PB.46058.8	PREDICTED: WD-40 repeat-containing protein MSI3 [Vitis vinifera]
802	PB.45115.1	PREDICTED: WD repeat-containing protein 89 homolog [Prunus mume]
803	PB.45422.1	PREDICTED: WD repeat-containing protein 76 isoform X1 [Jatropha curcas]
804	PB.43282.7	PREDICTED: COMPASS-like H3K4 histone methylase component WDR5B [Nelumbo nucifera]
805	PB.45695.6	PREDICTED: WD-40 repeat-containing protein MSI4-like [Vigna radiata var. radiata]
806	PB.42980.4	PREDICTED: ribosome biogenesis protein WDR12 homolog [Vitis vinifera]

807	PB.43263.1	PREDICTED: WD repeat-containing protein 43 isoform X2 [Vitis vinifera]
808	PB.39472.1	Transducin family protein / WD-40 repeat family protein isoform 4 [Theobroma cacao]
809	PB.36483.1	PREDICTED: WD-40 repeat-containing protein MSI4-like [Jatropha curcas]
810	PB.41134.1	wd repeat-containing protein lwd1 [Quercus suber]
811	PB.39763.1	PREDICTED: WD repeat-containing protein 11 [Prunus mume]
812	PB.29659.1	PREDICTED: F-box/WD-40 repeat-containing protein At3g52030 [Populus euphratica]
813	PB.38948.1	transducin/WD40 repeat protein [Medicago truncatula]
814	PB.39716.1	wd repeat-containing protein 53 [Quercus suber]
815	PB.38722.1	PREDICTED: WD repeat-containing protein 89 homolog [Arachis duranensis]
816	PB.29118.6	PREDICTED: WD repeat-containing protein 89 homolog [Vitis vinifera]
817	PB.35844.1	PREDICTED: glutamate-rich WD repeat-containing protein 1 [Pyrus x bretschneideri]
818	PB.35846.2	WD40 protein [Paeonia suffruticosa]
819	PB.36042.1	PREDICTED: WD repeat-containing protein YMR102C [Vitis vinifera]
820	PB.33081.1	WD repeat-containing 44 [Gossypium arboreum]
821	PB.34757.12	PREDICTED: WD repeat-containing protein 43 isoform X1 [Vitis vinifera]
822	PB.36021.4	PREDICTED: WD repeat-containing protein 26 isoform X1 [Juglans regia]
823	PB.34664.1	WD repeat-containing 26 [Gossypium arboreum]
824	PB.35830.3	PREDICTED: WD repeat-containing protein 76-like [Ziziphus jujuba]
825	PB.32832.3	WD repeat-containing protein WRAP73 [Hevea brasiliensis]
826	PB.31962.6	PREDICTED: WD repeat-containing protein 76-like [Ziziphus jujuba]
827	PB.35071.1	PREDICTED: WD repeat-containing protein 11 isoform X1 [Gossypium raimondii]
828	PB.39385.1	PREDICTED: WD repeat-containing protein 55 homolog [Beta vulgaris subsp. vulgaris]
829	PB.27538.1	PREDICTED: WD repeat-containing protein 55 homolog [Nelumbo nucifera]
830	PB.27270.1	PREDICTED: WD repeat-containing protein 44 [Nelumbo nucifera]
831	PB.31213.4	PREDICTED: WD repeat-containing protein 26 [Vitis vinifera]
832	PB.30575.1	PREDICTED: WD repeat-containing protein 25 [Nelumbo nucifera]
833	PB.27342.1	PREDICTED: LOW QUALITY PROTEIN: WD repeat and HMG-box DNA-binding protein 1 [Vitis vinifera]

834	PB.28669.1	PREDICTED: WD repeat-containing protein 43 isoform X2 [Vitis vinifera]
835	PB.30212.1	PREDICTED: WD repeat-containing protein 26 isoform X1 [Nelumbo nucifera]
836	PB.30748.11	PREDICTED: WD repeat-containing protein 43 isoform X2 [Vitis vinifera]
837	PB.29074.1	WD repeat-containing protein 70 [Glycine soja]
838	PB.26390.1	PREDICTED: WD repeat-containing protein 48 isoform X1 [Jatropha curcas]
839	PB.27622.3	PREDICTED: WD repeat-containing protein 43 isoform X2 [Vitis vinifera]
840	PB.26588.13	PREDICTED: WD repeat-containing protein 70 [Prunus mume]
841	PB.26520.1	PREDICTED: WD repeat-containing protein 26-like [Prunus mume]
842	PB.26784.2	PREDICTED: WD repeat-containing protein 43 isoform X2 [Vitis vinifera]
843	PB.26242.1	WD repeat-containing protein 44-like isoform X2 [Hevea brasiliensis]
844	PB.26069.1	PREDICTED: WD repeat-containing protein 70 [Vitis vinifera]
845	PB.24690.2	PREDICTED: WD repeat-containing protein 26-like [Prunus mume]
846	PB.25148.2	PREDICTED: WD repeat-containing protein DWA2 isoform X2 [Vitis vinifera]
847	PB.22774.1	Transducin/WD40 repeat-like superfamily protein [Theobroma cacao]
848	PB.43415.1	Transducin/WD40 repeat-like superfamily protein [Theobroma cacao]
849	PB.23077.7	PREDICTED: glutamate-rich WD repeat-containing protein 1 [Pyrus x bretschneideri]
850	PB.24885.1	PREDICTED: WD repeat-containing protein 70 [Ricinus communis]
851	PB.23930.1	PREDICTED: WD repeat-containing protein 75 [Citrus sinensis]
852	PB.23948.1	PREDICTED: glutamate-rich WD repeat-containing protein 1 [Pyrus x bretschneideri]
853	PB.17023.1	PREDICTED: WD repeat-containing protein 55 [Vitis vinifera]
854	PB.14517.2	Transducin family protein / WD-40 repeat family protein isoform 2 [Theobroma cacao]
855	PB.17580.1	Transducin/WD40 repeat-like superfamily protein [Theobroma cacao]
856	PB.20258.1	PREDICTED: WD repeat-containing protein YMR102C [Vitis vinifera]
857	PB.19664.1	WD repeat-containing protein 44 [Manihot esculenta]
858	PB.13551.1	PREDICTED: katanin p80 WD40 repeat-containing subunit B1 homolog isoform X3 [Populus euphratica]
859	PB.19552.1	PREDICTED: bromodomain and WD repeat-containing protein 3 isoform X2 [Vitis vinifera]
860	PB.21120.1	PREDICTED: WD repeat-containing protein 75 [Prunus mume]

861	PB.14820.7	PREDICTED: LOW QUALITY PROTEIN: WD repeat and HMG-box DNA-binding protein 1 [Vitis vinifera]
862	PB.12444.1	PREDICTED: bromodomain and WD repeat-containing protein 1 [Nelumbo nucifera]
863	PB.12026.1	PREDICTED: WD repeat-containing protein 44 isoform X1 [Cucumis melo]
864	PB.13640.8	WD repeat-containing protein 75 [Manihot esculenta]
865	PB.11398.1	PREDICTED: WD repeat-containing protein YMR102C [Vitis vinifera]
866	PB.15787.2	PREDICTED: WD repeat-containing protein 82-B [Sesamum indicum]
867	PB.15203.1	PREDICTED: WD repeat-containing protein 44 [Theobroma cacao]
868	PB.9989.2	PREDICTED: WD repeat-containing protein 76 isoform X2 [Jatropha curcas]
869	PB.12458.5	WD repeat-containing 44 [Gossypium arboreum]
870	PB.12121.1	PREDICTED: WD repeat-containing protein 44 isoform X1 [Cucumis melo]
871	PB.12627.1	PREDICTED: WD repeat-containing protein 44 isoform X1 [Cucumis melo]
872	PB.13887.1	PREDICTED: WD repeat-containing protein 55 homolog [Nelumbo nucifera]
873	PB.12982.3	PREDICTED: WD repeat-containing protein 89 homolog [Vitis vinifera]
874	PB.17087.1	PREDICTED: LOW QUALITY PROTEIN: WD repeat and HMG-box DNA-binding protein 1 [Vitis vinifera]
875	PB.20159.1	PREDICTED: WD repeat-containing protein 48-like [Populus euphratica]
876	PB.19799.1	PREDICTED: WD repeat-containing protein 55 homolog [Beta vulgaris subsp. vulgaris]
877	PB.15556.13	wd repeat-containing protein 3 [Quercus suber]
878	PB.8923.1	PREDICTED: WD repeat-containing protein 44 isoform X1 [Cucumis melo]
879	PB.11544.1	PREDICTED: WD repeat-containing protein 55 [Vitis vinifera]
880	PB.9347.9	PREDICTED: WD repeat-containing protein YMR102C [Vitis vinifera]
881	PB.8827.19	PREDICTED: LOW QUALITY PROTEIN: WD repeat-containing protein 44 [Prunus mume]
882	PB.11175.6	PREDICTED: WD repeat-containing protein 55 homolog [Nelumbo nucifera]
883	PB.10631.1	PREDICTED: WD repeat-containing protein 44-like [Juglans regia]
884	PB.10503.1	PREDICTED: WD repeat-containing protein 74 [Fragaria vesca subsp. vesca]
885	PB.8371.1	PREDICTED: WD repeat-containing protein 44 isoform X1 [Cucumis melo]
886	PB.6801.11	PREDICTED: WD repeat-containing protein 3 isoform X2 [Ricinus communis]
887	PB.6448.1	PREDICTED: WD repeat-containing protein 55 homolog [Nelumbo nucifera]

888		PB.5198.13	PREDICTED: katanin p80 WD40 repeat-containing subunit B1 homolog [Vitis vinifera]
889		PB.1946.1	PREDICTED: katanin p80 WD40 repeat-containing subunit B1 homolog isoform X1 [Vitis vinifera]
890		PB.6075.2	PREDICTED: katanin p80 WD40 repeat-containing subunit B1 homolog [Vitis vinifera]
891		PB.2849.2	PREDICTED: bromodomain and WD repeat-containing protein 3 [Jatropha curcas]
892		PB.4004.2	PREDICTED: WD repeat-containing protein 13 [Vitis vinifera]
893		PB.2032.27	WD repeat-containing protein LWD1 [Hevea brasiliensis]
894		PB.868.18	PREDICTED: bromodomain and WD repeat-containing protein 3 [Vitis vinifera]
895		PB.1993.17	PREDICTED: WD repeat-containing protein 11-like isoform X2 [Pyrus x bretschneideri]
896		PB.385.46	PREDICTED: katanin p80 WD40 repeat-containing subunit B1 homolog isoform X1 [Vitis vinifera]
897		PB.2789.1	WD repeat-containing 44 [Gossypium arboreum]
898		PB.65874.1	WD repeat-containing protein 48 isogeny [Cajanus cajan]
899		PB.66056.1	F-box-like/WD repeat-containing protein [Morus notabilis]
900		PB.66352.1	PREDICTED: glutamate-rich WD repeat-containing protein 1 [Erythranthe guttata]
901		PB.65432.1	Transducin/WD40 repeat-like superfamily protein [Theobroma cacao]
902	TFIID	PB.58150.1	PREDICTED: transcription initiation factor TFIID subunit 15 [Jatropha curcas]
903		PB.35353.3	PREDICTED: transcription initiation factor TFIID subunit 6-like isoform X2 [Vitis vinifera]
904		PB.65121.1	PREDICTED: transcription initiation factor TFIID subunit 10 [Ziziphus jujuba]
905		PB.62735.1	PREDICTED: transcription initiation factor TFIID subunit 7 [Nelumbo nucifera]
906		PB.56110.1	PREDICTED: transcription initiation factor TFIID subunit 15 [Jatropha curcas]
907		PB.56989.1	PREDICTED: transcription initiation factor TFIID subunit 11 [Ziziphus jujuba]
908		PB.56539.1	PREDICTED: transcription initiation factor TFIID subunit 15 [Jatropha curcas]
909		PB.56444.1	PREDICTED: transcription initiation factor TFIID subunit 8 [Vitis vinifera]
910		PB.56504.1	PREDICTED: transcription initiation factor TFIID subunit 14b-like [Jatropha curcas]
911		PB.49866.2	PREDICTED: transcription initiation factor TFIID subunit 14b-like [Jatropha curcas]
912		PB.52430.2	PREDICTED: transcription initiation factor TFIID subunit 11 [Ziziphus jujuba]
913		PB.45295.1	PREDICTED: transcription initiation factor TFIID subunit 15b [Vitis vinifera]
914		PB.40817.3	PREDICTED: transcription initiation factor TFIID subunit 6 [Eucalyptus grandis]

915	PB.36887.1	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X2 [Vitis vinifera]
916	PB.37948.1	PREDICTED: transcription initiation factor TFIID subunit 6-like isoform X2 [Vitis vinifera]
917	PB.35265.2	PREDICTED: transcription initiation factor TFIID subunit 8-like [Vitis vinifera]
918	PB.34176.1	PREDICTED: transcription initiation factor TFIID subunit 12, partial [Vitis vinifera]
919	PB.32512.4	PREDICTED: transcription initiation factor TFIID subunit 12b-like [Brassica rapa]
920	PB.27019.1	PREDICTED: transcription initiation factor TFIID subunit 11 [Ziziphus jujuba]
921	PB.30416.3	PREDICTED: transcription initiation factor TFIID subunit 12-like [Glycine max]
922	PB.34516.1	PREDICTED: transcription initiation factor TFIID subunit 5 [Vitis vinifera]
923	PB.27763.1	PREDICTED: transcription initiation factor TFIID subunit 11 [Ziziphus jujuba]
924	PB.27407.1	PREDICTED: transcription initiation factor TFIID subunit 11 isoform X2 [Vitis vinifera]
925	PB.31629.2	PREDICTED: transcription initiation factor TFIID subunit 15 [Jatropha curcas]
926	PB.24337.1	PREDICTED: transcription initiation factor TFIID subunit 5 [Vitis vinifera]
927	PB.26007.1	PREDICTED: transcription initiation factor TFIID subunit 11 [Ziziphus jujuba]
928	PB.24883.1	PREDICTED: transcription initiation factor TFIID subunit 5 [Vitis vinifera]
929	PB.21073.1	PREDICTED: transcription initiation factor TFIID subunit 4b-like [Vitis vinifera]
930	PB.23338.1	PREDICTED: transcription initiation factor TFIID subunit 5 [Vitis vinifera]
931	PB.24726.1	PREDICTED: transcription initiation factor TFIID subunit 5 [Vitis vinifera]
932	PB.16221.1	PREDICTED: transcription initiation factor TFIID subunit 2 isoform X1 [Vitis vinifera]
933	PB.21369.1	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X2 [Vitis vinifera]
934	PB.11644.1	PREDICTED: transcription initiation factor TFIID subunit 5 [Vitis vinifera]
935	PB.12717.1	PREDICTED: transcription initiation factor TFIID subunit 2 isoform X3 [Vitis vinifera]
936	PB.13783.1	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X1 [Vitis vinifera]
937	PB.14064.1	PREDICTED: transcription initiation factor TFIID subunit 15 [Jatropha curcas]
938	PB.12822.5	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X2 [Vitis vinifera]
939	PB.4486.27	PREDICTED: transcription initiation factor TFIID subunit 4b-like isoform X1 [Ziziphus jujuba]
940	PB.7570.4	PREDICTED: transcription initiation factor TFIID subunit 11 [Ziziphus jujuba]
941	PB.8787.1	PREDICTED: transcription initiation factor TFIID subunit 2 [Camelina sativa]

942	PB.8719.2	Transcription initiation factor TFIID subunit 10 [Cajanus cajan]
943	PB.10429.1	PREDICTED: transcription initiation factor TFIID subunit 5 [Vitis vinifera]
944	PB.7941.1	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X2 [Jatropha curcas]
945	PB.9266.3	PREDICTED: transcription initiation factor TFIID subunit 4b-like [Vitis vinifera]
946	PB.11793.1	PREDICTED: transcription initiation factor TFIID subunit 4b-like [Vitis vinifera]
947	PB.5157.9	PREDICTED: transcription initiation factor TFIID subunit 12, partial [Vitis vinifera]
948	PB.6557.1	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X1 [Vitis vinifera]
949	PB.5468.1	PREDICTED: transcription initiation factor TFIID subunit 2 isoform X2 [Vitis vinifera]
950	PB.4105.2	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X1 [Vitis vinifera]
951	PB.1432.27	PREDICTED: transcription initiation factor TFIID subunit 2 isoform X2 [Vitis vinifera]
952	PB.852.11	PREDICTED: transcription initiation factor TFIID subunit 5 [Vitis vinifera]
953	PB.1450.23	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X2 [Vitis vinifera]
954	PB.921.2	PREDICTED: transcription initiation factor TFIID subunit 4b-like isoform X2 [Ziziphus jujuba]
955	PB.801.1	PREDICTED: transcription initiation factor TFIID subunit 1 [Nelumbo nucifera]
956	PB.1704.4	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X1 [Vitis vinifera]
957	PB.2151.9	PREDICTED: transcription initiation factor TFIID subunit 1 isoform X1 [Eucalyptus grandis]
958	PB.3472.1	PREDICTED: transcription initiation factor TFIID subunit 2 isoform X2 [Vitis vinifera]
959	PB.2277.2	PREDICTED: transcription initiation factor TFIID subunit 11 [Ziziphus jujuba]