

Table S1. *Arabidopsis* flowering time related genes

Locus id	chr	start	end	annotation	reference
AT1G01060	1	33379	37840	LHY	[1-3]
AT1G02340	1	465718	467844	HFR1	[4]
AT1G03970	1	1018100	1019247	GBF4	[2]
AT1G04400	1	1185549	1188516	CRY2	[1-3]
AT1G04440	1	1202254	1205802	CKL13	[1]
AT1G06040	1	1828412	1829889	STO	[1]
AT1G07980	1	2473165	2474927	NF-YC10	[5]
AT1G08620	1	2737031	2743724	PKDM7D	[6]
AT1G08970	1	2882543	2884337	NF-YC9	[5]
AT1G09030	1	2908611	2909030	NF-YB4	[5]
AT1G09530	1	3076584	3079541	PAP3; PIF3	[1,7]
AT1G09570	1	3095258	3100359	PHYA	[1-3]
AT1G09700	1	3137769	3140355	HYL1	[1]
AT1G10588	1	3501146	3501905	AT1G10588	[1]
AT1G12190	1	4394895	4396289	LWD1	[1,8]
AT1G12610	1	4289942	4291015	DDF1	[1,9]
AT1G12910	1	4394897	4396291	LWD1	[8]
AT1G13260	1	4542165	4543739	RAV1	[1,2]
AT1G14920	1	5149221	5151349	GAI	[1,10]
AT1G15550	1	5344473	5346161	GA4	[1]
AT1G17455	1	5997180	5998475	ELF4-L4	[2]
AT1G17590	1	6050222	6052541	NF-YA8	[5]
AT1G18100	1	6227217	6230188	MFT	[10]
AT1G18450	1	6348100	6351968	ATARP4	[1]
AT1G21970	1	7727577	7729617	NF-YB9	[5]
AT1G22690	1	8027287	8028114	AT1G22690	[1]
AT1G22770	1	8061833	8067705	GI	[1-3]
AT1G24260	1	8593631	8596087	SEPALLATA3	[1,11]
AT1G25540	1	8969052	8974647	PFT1	[1,2]
AT1G25560	1	8981677	8983041	TEM1	[2]
AT1G26310	1	9100140	9103590	CAL	[1,2]
AT1G26790	1	9273857	9275312	AtDOF1.3	[2]
AT1G29160	1	10183797	10184324	AtDOF1.5	[2]
AT1G30040	1	10537632	10539815	ATGA2OX2	[1]
AT1G30500	1	10804450	10806319	NF-YA7	[5]
AT1G30950	1	11036161	11037489	UFO	[1]
AT1G30960	1	11037593	11040014	AT1G30960	[1]
AT1G30970	1	11040262	11043732	SUF4	[1,2]
AT1G31814	1	11412589	11414483	FRL2	[1,10]
AT1G35160	1	12867181	12868793	GRF4	[2]
AT1G43700	1	16486671	16488681	VIP1	[1]
AT1G44090	1	16763117	16764926	ATGA20OX5	[1]
AT1G44446	1	16850799	16853664	CH1	[1]
AT1G45249	1	17165127	17167866	ABF2	[2]
AT1G47250	1	17319177	17321074	PAF2	[2]
AT1G48270	1	17831621	17834088	GCR1	[1]
AT1G49480	1	18317846	18320313	RTV1	[1]

AT1G49720	1	18400092	18402628	ABF1	[2]
AT1G50370	1	18658647	18661974	FYPP3	[12]
AT1G50680	1	18777601	18778614	RAVL2	[2]
AT1G50960	1	18893217	18895387	ATGA2OX7	[1]
AT1G51120	1	18938091	18939149	RAVL3	[2]
AT1G52800	1	19667712	19669030	AT1G52800	[1]
AT1G53090	1	19787020	19790570	SPA4	[1-3]
AT1G53160	1	19810087	19811276	SPL4	[1,3]
AT1G54160	1	20217336	20219452	NF-YA5	[5]
AT1G54830	1	20451609	20452671	NF-YC3	[5]
AT1G55080	1	20556678	20557921	AT1G55080	[1]
AT1G56170	1	21024764	21025883	NF-YC2	[5]
AT1G57820	1	21417835	21421611	ORTH2	[1]
AT1G59940	1	22069282	22070638	ARR3	[1]
AT1G60980	1	22456238	22457805	ATGA20OX4	[1]
AT1G61040	1	22486872	22489634	VIP5	[1]
AT1G62750	1	23237099	23240112	ATSCO1	[1]
AT1G62830	1	23368155	23270867	LDL1	[1,13]
AT1G63030	1	23371072	23372075	DDF2	[1,9]
AT1G65480	1	24335091	24337597	FT	[1-3]
AT1G66350	1	24751858	24753706	RGL1	[1,10]
AT1G66410	1	24774217	24775900	CAM4	[14]
AT1G68050	1	25512339	25514552	FKF1	[1-3]
AT1G68840	1	25880327	25881736	TEM2	[2]
AT1G69120	1	25985993	25989976	AP1	[1-3]
AT1G69570	1	26161528	26163496	AtDOF1.10	[2,15]
AT1G69935	1	26345464	26346789	AT1G69935	[1]
AT1G70170	1	26427537	26429019	MMP	[1]
AT1G71692	1	26956307	26958789	AGL12	[1]
AT1G72050	1	27118686	27121132	AT1G72050	[1]
AT1G72630	1	27344270	27345576	ELF4-L2	[2]
AT1G72830	1	27409118	27411630	HAP2C; NF-YA3	[1,5]
AT1G74660	1	28051237	28051788	MIF1	[1]
AT1G74670	1	28056947	28057810	AT1G74670	[1]
AT1G76710	1	28794623	28797570	ASHH1; SDG26	[1,16]
AT1G77080	1	28960531	28964990	FLM; MAF1	[1,17]
AT1G77300	1	29044816	29053704	EFS	[1,2]
AT1G78300	1	29461671	29463385	GRF2	[2]
AT1G78440	1	29516492	29517944	ATGA2OX1	[1]
AT1G79280	1	29824069	29837871	NUA	[1]
AT1G79460	1	29895285	29899480	GA2	[1]
AT1G79730	1	30005431	30008898	ELF7	[1,18]
AT1G80330	1	30202953	30204429	ATGA3OX4	[1]
AT1G80340	1	30205585	30207092	GA4H	[1]
AT1G80680	1	30328900	30333661	MOS3	[1]
AT2G01570	2	255248	257549	RGA1	[1,10]
AT2G02560	2	689787	697595	CAND1	[1]
AT2G02950	2	854946	856537	PKS1	[1]
AT2G03710	2	1129265	1131835	SEPALLATA4	[1]

AT2G04030	2	1281838	1286101	CR88	[1]
AT2G06210	2	2428900	2436684	ELF8	[1,18]
AT2G06255	2	2457573	2459811	ELF4-L3	[2]
AT2G13540	2	5636884	5642957	ABH1	[19]
AT2G13570	2	5655842	5656489	NF-YB7	[5]
AT2G14900	2	6411257	6412412	AT2G14900	[1]
AT2G17770	2	7723103	7724161	FDP	[2]
AT2G18040	2	7842164	7843870	PIN1AT	[20]
AT2G18790	2	8146963	8151512	PHYB	[1-3]
AT2G18870	2	8167471	8168668	VEL3	[2]
AT2G18880	2	8172189	8178995	VEL2	[2]
AT2G18915	2	8201654	8204656	LKP2	[1,2]
AT2G19520	2	8455936	8459525	FVE	[2]
AT2G20180	2	8711105	8713973	PIL5	[1]
AT2G21070	2	9047944	9050609	FIONA1	[1,21]
AT2G22540	2	9586954	9590973	SVP	[1-3]
AT2G22630	2	9625542	9629037	AGL17	[1,22]
AT2G23070	2	9823965	9826898	CK2alphaCp	[2]
AT2G23080	2	9827113	9829537	CK2A /CK2alphaC	[2]
AT2G23380	2	99626650	9967439	CLF	[1,2]
AT2G24790	2	10573977	10575224	COL3	[1,23]
AT2G25930	2	11066113	11070402	ELF3	[1,2]
AT2G26710	2	11387570	11390690	BAS1	[1]
AT2G27470	2	11744981	11746411	NF-YB11	[5]
AT2G27550	2	11780328	11781758	ATC	[1,10]
AT2G28190	2	12014498	12016569	CZS	[24]
AT2G28290	2	12063290	12080160	SYD	[1]
AT2G28550	2	12233028	12235620	TOE1	[1,2]
AT2G29950	2	12767585	12768435	ELF4-L1	[2]
AT2G30120	2	12860541	12861910	FLX	[25]
AT2G30810	2	13134903	13135743	AT2G30810	[1]
AT2G31650	2	13455272	13462181	ATX1	[26]
AT2G32950	2	13985010	13990612	COP1	[1-3]
AT2G33810	2	14312077	14313148	SPL3	[1,3]
AT2G33835	2	14318863	14321776	FES1	[1][[10]
AT2G34140	2	14413758	14414765	AtDOF2.3	[2]
AT2G34555	2	14564067	14565776	ATGA2OX3	[1]
AT2G34720	2	14649767	14651627	NF-YA4	[5]
AT2G34880	2	14711880	14716634	PKDM7C	[6]
AT2G35670	2	14992565	14996756	FIE2	[2]
AT2G36270	2	15204764	15207528	ABI5	[2]
AT2G37060	2	15576063	15577877	NF-YB8	[5]
AT2G37678	2	15808543	15809871	FHY1	[1]
AT2G38880	2	16238492	16240476	NF-YB1	[5]
AT2G39250	2	16395964	16398151	SNZ	[1,2]
AT2G39540	2	16507944	16508319	AT2G39540	[1]
AT2G39810	2	16619878	16625135	HOS1	[1,10]
AT2G40080	2	16741372	16741990	ELF4	[1,2]
AT2G41070	2	17130788	17132370	EEL	[2]

AT2G42200	2	17587407	17589630	SPL9	[3]
AT2G42590	2	17731855	17733967	GRF9	[27]
AT2G42830	2	17827332	17831090	SHP2	[1]
AT2G43010	2	17893504	17896127	PIF4	[1,28]
AT2G43410	2	18032324	18038320	FPA	[1,2]
AT2G44680	2	18433621	18435432	CKB4	[1,29]
AT2G45650	2	18811424	18813596	AGL6	[1,11,30,31]
AT2G45660	2	18814612	18818121	SOC1	[1-3]
AT2G46340	2	19029246	19034486	SPA1	[1-3]
AT2G46790	2	19239718	19242156	APRR9	[1,2]
AT2G46830	2	19252741	19255983	CCA1	[1,2]
AT2G47310	2	19430754	19434347	AT2G47310	[1]
AT2G47700	2	19559384	19561654	RF12	[1,32]
AT2G47810	2	19582750	19583618	NF-YB5	[5]
AT3G01460	3	173323	182461	MBD9	[1]
AT3G02310	3	464286	467081	SEPALLATA2	[1]
AT3G02380	3	487243	488700	COL2	[1,2]
AT3G02520	3	526437	528313	GRF7	[2]
AT3G02885	3	638028	639062	GASA5	[1]
AT3G03090	3	700463	704776	ATVGT1	[1]
AT3G03450	3	819337	821406	RGL2	[10]
AT3G04510	3	1215642	1216964	AT3G04510	[1]
AT3G04610	3	1250559	1254879	FLK	[1,2]
AT3G05040	3	1401277	1408203	HST	[1]
AT3G05120	3	1430477	1432784	ATGID1A; GID1/GAr2	[1,10,33,34]
AT3G05690	3	1676552	1678938	ATHAP2B; NF-YA2	[1,5]
AT3G06910	3	2178636	2181203	AT3G06910	[1]
AT3G06930	3	2185149	2189393	AtPRMT4b	[1]
AT3G07650	3	2441663	2444538	COL9	[1,35]
AT3G10185	3	3145584	3146204	AT3G10185	[1]
AT3G10390	3	3229298	3231824	FLD	[1,2]
AT3G11440	3	3602099	3605110	ATMYB65	[1]
AT3G11540	3	3631887	3637955	SPY	[2]
AT3G12480	3	3957777	3960472	NF-YC11	[5]
AT3G12810	3	4065049	4074085	PIE1	[1,2]
AT3G13550	3	4423349	4424883	FUS9	[36]
AT3G13682	3	4479200	4481516	LDL2	[1,13]
AT3G14020	3	4642968	4644301	NF-YA6	[5]
AT3G15270	3	5140372	5141355	SPL5	[1,3]
AT3G15354	3	5169102	5172844	SPA3	[1-3]
AT3G18990	3	6548875	6551859	VRN1	[1,2]
AT3G19040	3	6567163	6575288	HAF2	[1]
AT3G19140	3	6614910	6615335	DNF	[37]
AT3G19290	3	6687140	6690123	ABF4	[2]
AT3G19980	3	6961742	6965114	ATFYPP3	[1]
AT3G20550	3	7174470	7177948	DDL	[1]
AT3G20740	3	7248815	7252458	FIE	[1,2]
AT3G20910	3	7326382	7328576	NF-YA9	[5]
AT3G21320	3	7499059	7501847	putative EEC	[1,2]

AT3G22380	3	7912912	7919517	TIC	[1,38]
AT3G22590	3	8003934	8005579	PHP; CDC73	[39]
AT3G24440	3	8876034	8878178	VRN5	[1,2]
AT3G25730	3	9396420	9397747	ARF14	[2]
AT3G26120	3	9547635	9550423	TEL1	[1]
AT3G26640	3	9794457	9795694	LWD2	[1,8]
AT3G26790	3	9855065	9857226	FUS3	[1]
AT3G30180	3	11813216	11816244	BR6OX2	[1]
AT3G30260	3	11909119	11912880	AGL79	[2]
AT3G33520	3	14104642	14105635	ATARP6	[1,2]
AT3G44460	3	16079976	16081906	AtbZIP67	[2]
AT3G46640	3	17194075	17196203	PCL1; LUX	[1,2]
AT3G47500	3	17514985	17517043	CDF3	[1-3]
AT3G48430	3	17946594	17951731	REF6	[1,2]
AT3G48590	3	18008657	18009982	NF-YC1	[5]
AT3G50000	3	18534459	18536970	CKA2 / CK2alphaB	[2]
AT3G53340	3	19774331	19776205	NF-YB10	[5]
AT3G54220	3	20070158	20072780	SCR	[40]
AT3G54720	3	20265703	20268826	AMP1	[1]
AT3G54990	3	20384695	20387499	SMZ	[1,2]
AT3G56850	3	21046341	21048516	AREB3	[2]
AT3G57230	3	21188689	21191911	AGL16	[1]
AT3G57300	3	21210467	21218864	AT3G57300	[1]
AT3G57390	3	21244678	21236888	AGL18	[1,2]
AT3G58070	3	21517590	21518631	GIS	[1]
AT3G58780	3	21749437	21752884	SHP1	[1,41]
AT3G59060	3	21838955	21841484	PIL6	[1,4]
AT3G60250	3	22281312	22283088	CKB3	[1,42]
AT3G61120	3	22629234	22631466	AGL13	[1]
AT3G62090	3	22999522	23001684	PIL2; PIF6	[1,4]
AT3G63010	3	23300400	23302461	ATGID1B; GID1L2/GAr3	[1,10,33,34]
AT4G00450	4	203471	211003	CRP	[1]
AT4G00650	4	269026	271503	FRI	[1-3]
AT4G00690	4	281645	283129	AT4G00690	[1]
AT4G01060	4	460472	461085	ETC3	[1]
AT4G02020	4	886600	891955	EZA1; SWN1	[1,43]
AT4G02560	4	1123490	1128421	LD	[1,2]
AT4G02780	4	1237767	1244813	GA1	[1,10]
AT4G03400	4	1497535	1499864	DFL2	[1]
AT4G04885	4	2471972	2475981	PCFS4	[44]
AT4G04890	4	2476487	2482343	PDF2	[1]
AT4G05420	4	2746121	2752943	DDB1A	[45]
AT4G08920	4	5724100	5727250	CRY1	[1-3]
AT4G09000	4	5775210	5777481	GRF1	[2]
AT4G09610	4	6074767	6075642	GASA2	[1]
AT4G09960	4	6236473	6240681	STK	[41]
AT4G10180	4	6346463	6349276	DET1	[45,46]
AT4G11110	4	6771601	6777221	SPA2	[1-3]
AT4G11880	4	7143109	7147216	AGL14	[1,2]

AT4G12480	4	7406102	7406934	pEARLI1	[1]
AT4G14110	4	8132881	8134915	COP9	[1]
AT4G14540	4	8344616	8345218	NF-YB3	[5]
AT4G14690	4	8418278	8419258	ELIP2	[1]
AT4G15090	4	8614063	8618142	FAR1	[1,47]
AT4G15180	4	8651406	8662587	AT4G15180	[1]
AT4G15880	4	9012660	9016131	ESD4	[1,10]
AT4G16250	4	9195617	9199501	PHYD	[1,2]
AT4G16280	4	9206613	9214841	FCA	[1,2]
AT4G16780	4	9449133	9450762	HAT4	[1]
AT4G16810	4	9459889	9462272	B3H7I2	[1,2]
AT4G16845	4	9476162	9479897	VRN2	[1,2]
AT4G17640	4	9825210	9827285	CKB2	[1,42]
AT4G18130	4	10042149	10046094	PHYE	[1,2]
AT4G18960	4	10382856	10388539	AG	[48,49]
AT4G20370	4	11000782	11003007	TSF	[1-3]
AT4G20400	4	11008666	11013842	PKDM7B	[6]
AT4G21100	4	11258761	11265463	DDB1B	[45]
AT4G21200	4	11302695	11306611	ATGA2OX8	[1]
AT4G21690	4	11527241	11529072	ATGA3OX3	[1]
AT4G22140	4	11727738	11730521	EBS	[1,10]
AT4G22950	4	12023926	12027432	AGL19	[1,2]
AT4G23340	4	12195463	12196803	AT4G23340	[1]
AT4G24210	4	12563553	12564482	SLY1	[10]
AT4G24520	4	12990894	12992468	GA5	[1]
AT4G24540	4	12670975	12674082	AGL24	[1-3]
AT4G24620	4	12708762	12712835	PGI1	[1]
AT4G25530	4	13038369	13042452	FWA	[1,13]
AT4G26000	4	13197246	13199836	PEP	[2]
AT4G27060	4	13581407	13585161	TOR1	[1]
AT4G27430	4	13718685	13723330	CIP7	[1]
AT4G29130	4	14352043	14355109	ATHXK1	[1]
AT4G29830	4	14597667	14599306	VIP3	[1,50]
AT4G30200	4	14786639	14790509	VEL1	[1,2]
AT4G31120	4	15132017	15136645	ATPRMT5; SKB1	[1,51]
AT4G31160	4	15145853	15153075	DCAF1	[45]
AT4G31380	4	15229791	15230724	FLP1	[1]
AT4G31500	4	15273477	15275316	CYP83B1	[1]
AT4G32040	4	15494071	15493262	KNAT5	[1]
AT4G32980	4	15914725	15918047	ATH1	[1,52]
AT4G33280	4	16047358	16049359	AT4G33280	[1]
AT4G34000	4	16295558	16298255	ABF3	[2]
AT4G34530	4	16498391	16500174	CIB1	[2]
AT4G35900	4	17004598	17006290	FD	[1-3]
AT4G36920	4	17400844	17403329	AP2	[1,2]
AT4G37580	4	17658606	17660872	HLS1	[1]
AT4G37940	4	17835689	17838615	AGL21	[1,11]
AT4G39400	4	18324655	18328820	BRI1	[1]
AT5G01400	5	162550	171072	ESP4	[53]

AT5G02200	5	437458	438892	FHL	[1]
AT5G02810	5	637895	641975	PRR7	[1,54]
AT5G02840	5	648702	651970	LCL1	[1]
AT5G03790	5	1004982	1006372	ATHB51	[1]
AT5G03840	5	1024640	1025811	TFL1	[1,2]
AT5G04240	5	1169545	1174879	ELF6	[1,2]
AT5G05690	5	1702689	1706788	CPD	[1]
AT5G06100	5	1837908	1840728	MYB33	[1,10]
AT5G06510	5	1984991	1987044	NF-YA10	[5]
AT5G07200	5	2243554	2245340	YAP169	[1]
AT5G08190	5	2635902	2637433	NF-YB12	[5]
AT5G08330	5	2680745	2681814	CHE	[1,2]
AT5G09230	5	2871474	2873779	SIR2; SRT2	[2,55]
AT5G10140	5	3173498	3179449	FLC	[1-3]
AT5G10450	5	3283853	3286317	GRF6	[2]
AT5G10625	5	3358788	3359782	AT5G10625	[1]
AT5G11260	5	3593381	3594993	HY5	[1]
AT5G11530	5	3696863	3701549	EMF1	[1]
AT5G12840	5	4050694	4053609	HAP2A; NF-YA1	[1,5]
AT5G13480	5	4326531	4331702	FY	[1,2]
AT5G13790	5	4449017	4450846	AGL15	[1,11]
AT5G14920	5	4826482	4827983	AT5G14920	[1]
AT5G15230	5	4944903	4946219	GASA4	[1]
AT5G15800	5	5151334	5154154	SEP_1	[11,30]
AT5G15840	5	5171185	5172761	CO	[1,3,10]
AT5G15850	5	5176094	5177900	COL1	[1,2]
AT5G15960	5	5209901	5210730	KIN1	[1]
AT5G15970	5	5211914	5212668	KIN2	[1]
AT5G16050	5	5243745	5245820	GRF5	[2]
AT5G16260	5	5311207	5315767	ELF9	[2]
AT5G16320	5	5344505	5346022	FRL1	[1,10]
AT5G17690	5	5827173	5829684	TFL2	[1,10,56]
AT5G19550	5	6598019	6601821	ASP2	[1]
AT5G20570	5	6956660	6958311	RBX1	[32]
AT5G20730	5	7016447	7022115	NPH4	[1]
AT5G23090	5	7749070	7750544	NF-YB13	[5]
AT5G23150	5	7785838	7792492	HUA2	[1,2]
AT5G23260	5	7836096	7838505	AGL32	[2]
AT5G24470	5	8355954	8358876	APRR5	[1,2]
AT5G24860	5	8541781	8542452	PPF1	[1,10]
AT5G24930	5	8589234	8591236	AtCOL4	[2]
AT5G25810	5	8986774	8987790	TNY	[1]
AT5G25900	5	9036021	9038409	GA3	[1]
AT5G27220	5	9578757	9582752	A_TM021B04.7	[2]
AT5G27230	5	9584095	9588052	AT5G27230	[1]
AT5G27320	5	9629090	9631213	ATGID1C; GID1L3/GAr1	[1,10,33,34]
AT5G27910	5	9940736	9941299	NF-YC8	[5]
AT5G28450	5	10372942	10374194	AT5G28450	[1]
AT5G28490	5	10454397	10455200	LSH1	[1]

AT5G35840	5	14025056	14028994	PHYC	[1,2]
AT5G37055	5	14641551	14642440	SEF1	[57]
AT5G37770	5	15016084	15016849	TCH2	[1]
AT5G38140	5	15220208	15222272	NF-YC12	[5]
AT5G38150	5	15240346	15242177	PMI15	[1]
AT5G38480	5	15409697	15411516	GRF3	[2]
AT5G39660	5	15895927	15898272	CDF2	[1-3]
AT5G41360	5	16561568	16566508	XPB2	[1]
AT5G41790	5	16744758	16750075	CIP1	[1]
AT5G41990	5	16794967	16798006	WNK8	[58]
AT5G42400	5	16954469	16960671	SDG25	[59]
AT5G42790	5	17159044	17161070	PAF1	[2]
AT5G42910	5	17203742	17205380	AtbZIP15	[2]
AT5G43250	5	17356174	17356566	NF-YC13	[5]
AT5G44080	5	17738652	17739873	bzip13	[2]
AT5G46210	5	18748645	18754037	CUL4	[1,60]
AT5G46910	5	19065007	19068107	AT5G46910	[1]
AT5G47010	5	19089236	19096561	LBA1	[1]
AT5G47080	5	19141839	19143838	CKB1	[1,42]
AT5G47640	5	19309227	19310272	NF-YB2	[5]
AT5G47670	5	19315023	19316169	NF-YB6	[5]
AT5G48150	5	19539481	19541924	PAT1	[1]
AT5G49020	5	19888477	19892146	AtPRMT4a	[1]
AT5G49230	5	19958873	19960544	HRB1	[61]
AT5G50470	5	20555120	20555758	NF-YC7	[5]
AT5G50480	5	20557574	20558487	NF-YC6	[5]
AT5G50490	5	20560434	20561187	NF-YC5	[5]
AT5G51230	5	20840962	20846790	EMF2	[1,2]
AT5G51310	5	20870080	20871944	AT5G51310	[1]
AT5G51810	5	21072414	21074034	AT2353	[1]
AT5G51820	5	21080594	21085283	PGM	[1]
AT5G53360	5	21647902	21649567	SINAT5	[62,63]
AT5G54270	5	22038273	22039568	CAB1A; LHCB3	[64]
AT5G54510	5	22148319	22150904	DFL1	[1]
AT5G55390	5	22447966	22454414	EDM2	[65]
AT5G57360	5	23258653	23261816	ZTL	[1,2]
AT5G57380	5	23263621	23266730	VIN3	[1,2]
AT5G57660	5	23355464	23356989	ATCOL5	[2]
AT5G58230	5	23573238	23575471	MSI1	[1,2]
AT5G59560	5	24017782	24019183	SRR1	[1,66]
AT5G59570	5	24021114	24022738	AT5G59570	[1]
AT5G59710	5	24074633	24079144	VIP2	[1]
AT5G59845	5	24128550	24129246	AT5G59845	[1]
AT5G60100	5	24215225	24218590	APRR3	[1,2]
AT5G60120	5	24225012	24228950	TOE2	[1,2]
AT5G60910	5	24519708	24523369	AGL8; FUL	[1,2,67]
AT5G61150	5	24620882	24624951	VIP4	[1,68]
AT5G61270	5	24638775	24640600	PIF7	[4]
AT5G61380	5	24692290	24695776	TOC1	[1,2]

AT5G61850	5	24861521	24864159	LFY	[1,2]
AT5G62040	5	24940036	24940935	BFT	[1,2]
AT5G62430	5	25086319	25088160	CDF1	[1-3]
AT5G62640	5	25166659	25169767	ELF5	[1,10]
AT5G63310	5	25371904	25373861	NDPK2	[69]
AT5G63470	5	25415701	25417098	NF-YC4	[5]
AT5G64813	5	25927505	25930122	LIP1	[1]
AT5G65050	5	25999480	26003552	AGL31; MAF2	[1,2]
AT5G65060	5	26004655	26008541	MAF3	[1,2]
AT5G65070	5	26009486	26013360	MAF4	[1,2]
AT5G65080	5	26014730	26019691	AGL68	[1,2]
AT5G65430	5	26148201	26150342	GRF8	[2]
AT5G65540	5	26212915	26215548	AT5G65540	[1]
AT5G67100	5	26794220	26802330	ICU2	[1]
AT5G67180	5	26801949	26804249	TOE3	[2]
AT5G67380	5	26880966	26883430	CKA1	[2]

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