

Resequenced amplicon	category	length (bp)	# SNPs	π	Θ_w	Tajima's D	* P<0.05
40S-RPS3a	CT	500	12	0.00601	0.00617	-0.336	
4CL1	WQ	628	8	0.00268	0.00316	-0.46	
4CL2	WQ	629	10	0.00237	0.00378	-1.128	
60S-RPL31a	CT	609	21	0.01011	0.00891	0.479	
ABA-WDS	CT	344	9	0.00662	0.00672	-0.048	
ABC_trnpr_53181_53184	CT	961	41	0.01182	0.0164	-1.49641	
adfb-F1R1	CT	603	4	0.00153	0.00184	-0.48187	
adf-FR	CT	631	10	0.00297	0.00492	-1.6699	
adprf-f1r1	CT	305	25	0.04122	0.03161	1.5989	
ads2_58648_58650	CT	159	3	0.00719	0.0077	-0.30187	
AFP1519	WQ	277	6	0.00739	0.00611	0.67111	
AGP4	WQ	379	8	0.0083	0.00791	-0.5575	
APX	CT	867	26	0.00636	0.00789	-0.7	
AT	WQ	2578	93	0.00936	0.00935	-0.037	
at1g10470_59575_59576	CT	470	44	0.01743	0.02759	-1.20897	
athb12_57781_57784	CT	411	24	0.01377	0.01836	-1.05904	
C3H	WQ	366	5	0.02183	0.01862	-0.17754	
ca-bp_17578069_17578072	CT	308	17	0.01098	0.01676	-1.45212	
CAD	WQ	142	3	0.00292	0.00575	-0.89659	
ca-d-pk_17796050_17796049	CT	365	2	0.00142	0.00147	-1.07169	
CCR1	WQ	436	7	0.00483	0.00405	0.57039	
ccr4_assoc_59599_59600	CT	371	8	0.00431	0.00592	-0.89329	
CESA	WQ	234	6	0.0568	0.00687	-0.521143	
chl-cor15-like_57801_57803	CT	372	29	0.01691	0.02349	-1.15639	
chr-remod_59584_59588	CT	500	25	0.01068	0.1395	-0.90872	
CL1400contig1_01	CT	358	9	0.01023	0.00688	2.15058	*
CL1432contig1_02	CT	585	13	0.00392	0.00602	-1.22346	
CL1692contig1_01	CT	336	13	0.01248	0.01144	-0.05584	
CL1811contig1_01	CT	716	31	0.00893	0.01203	-1.00173	
CL1814contig1_02	CT	342	3	0.00312	0.00238	0.79783	
CL1868contig1_02	CT	358	5	0.00455	0.00388	0.51794	
CL1994contig1_01	CT	481	8	0.00395	0.00469	-0.53053	
CL1997contig1_03	CT	415	6	0.00207	0.00397	-1.48513	
CL2089contig1_02	CT	398	18	0.01043	0.01225	-0.54269	
CL2133contig1_02	CT	308	4	0.0032	0.0036	0.30861	
CL2282contig1_01	CT	304	1	0.00172	0.00089	1.58255	
CL908contig1_01	CT	423	12	0.00423	0.00825	-1.79726	
CL919contig1_02	CT	374	15	0.01028	0.011	-0.23717	
CL922contig1_01	CT	409	4	0.00109	0.00268	-1.66713	
CL939contig1_03	CT	400	35	0.01577	0.02377	-1.25553	
CL969contig1_01	CT	379	6	0.00349	0.00449	-0.5551	

CN634677.1-01	CT	275	1	0.00086	0.00099	-0.21321
CN635596.1-01	CT	291	13	0.00906	0.01197	-0.79265
CN635674.1-02	CT	180	11	0.01378	0.01776	-1.17522
CN635691.1-01	CT	612	3	0.00053	0.00131	-1.49431
CN636134.1-01	CT	388	17	0.01536	0.01253	0.74323
CN636471.1-01	CT	332	10	0.01042	0.00841	0.66091
CN637306.1-01	CT	683	16	0.00491	0.0066	-0.95375
CN637339.1-01	CT	517	19	0.01104	0.0104	0.08198
CN637910.1-01	CT	266	4	0.00321	0.00407	-0.58793
CN637944.1-01	CT	239	7	0.00513	0.00951	-1.71874
CN638489.1-01	CT	418	11	0.00545	0.00713	-0.80801
CN638556.1-01	CT	363	3	0.00323	0.00234	0.40524
CN639087.1-01	CT	378	1	0.00023	0.00072	-1.16097
CN639480.1-02	CT	389	8	0.00656	0.00572	0.49156
CN639782.1-01	CT	443	12	0.01105	0.01077	-0.02523
CN640010.1-01	CT	345	19	0.01494	0.01516	-0.05866
CN640155.1-02	CT	299	3	0.0012	0.00299	-1.50358
CN640361.1-02	CT	715	21	0.01103	0.00819	1.1224
CN640485.1-02	CT	504	11	0.00442	0.00594	-1.24688
CN640521.1-01	CT	400	13	0.00897	0.00945	-0.18882
CN641217.1-01	CT	681	5	0.20251	0.18176	-1.78003
CN641226.1-01	CT	315	6	0.0067	0.00696	0.83674
COMT1	WQ	482	8	0.00514	0.00451	0.459141
dehydrin-like_57357_57358	CT	408	6	0.00256	0.00428	-1.31869
dehydrin-like_573587_57389	CT	517	11	0.00531	0.00647	-1.02757
EF1A	CT	1072	14	0.00274	0.00339	-0.656
ERD15-like	CT	646	14	0.00438	0.00563	-0.757
F3H1	CT	365	14	0.00528	0.00988	-1.576
F3H2	CT	647	14	0.00629	0.00562	0.15
FORMIN-like	CT	337	3	0.0048	0.00229	1.498
glut-S-trnsfr_58615_58616	CT	754	16	0.00412	0.00628	-1.33506
GLYHMT	WQ	143	4	0.00586	0.00758	-0.62834
LEA2	CT	504	18	0.00647	0.00878	-0.862
LEA-EMB11	CT	545	33	0.01378	0.01594	-0.593
LIM	WQ	281	14	0.1673	0.01367	0.80143
LP3-like	CT	481	16	0.00662	0.00848	-0.713
MT-LIKE	CT	579	20	0.01334	0.00911	1.639
PAL	WQ	394	16	0.01143	0.011	0.8924
pip_53159_530160	CT	260	8	0.00705	0.00869	-0.70064
pk19_58638_58641	CT	925	23	0.00609	0.00775	-0.83723
POLY-UBQ	CT	898	17	0.00544	0.00494	0.357
rc12a_58631_58630	CT	417	18	0.01313	0.01391	-0.23854

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SAM	WQ	512	7	0.00286	0.00368	-0.69695	
sSPcDFD005F06506_01	CT	455	9	0.00352	0.0055	-1.22302	
sSPcDFD009F11511_01	CT	647	24	0.00797	0.01039	-0.90266	
sSPcDFD021C11211_01	CT	381	14	0.0088	0.01074	-0.40672	
sSPcDFD024D11311_01	CT	364	4	0.00169	0.00298	-1.1941	
sSPcDFD040B03103_01	CT	347	4	0.00197	0.00316	-1.05777	
sSPcDFE002A03003_01	CT	462	17	0.00829	0.01054	-1.13215	
sSPcDFE003F04504_01	CT	342	5	0.00289	0.00409	-0.6767	
sSPcDFE025G02602_01	CT	328	16	0.00901	0.01338	-1.18548	
sSPcDFE028B10110_01	CT	414	9	0.00462	0.00604	-0.79668	
sSPcDFE038D06304_01	CT	360	7	0.00334	0.00529	-0.97687	
sSPcDFE044F10510_01	CT	342	7	0.00568	0.00569	-0.00468	
sSPcDFE049B06106_01	CT	322	1	0.0003	0.00086	-1.16356	
sSPcDFE049E11411_01	CT	387	8	0.00834	0.00567	1.54803	
sSPcDFF014F08508_01	CT	462	3	0.00155	0.00209	-0.8137	
sSPcDFF015H05705_01	CT	372	18	0.01399	0.01379	-0.43957	
sSPcDFF020H04704_01	CT	412	13	0.0066	0.00855	-0.79877	
sSPcDFF044H10710_01	CT	567	9	0.00444	0.00435	0.06868	
sSPcDFF053H03703_01	CT	598	15	0.00309	0.0068	-1.9476	
sSPcDFF06D104304_01	CT	266	57	0.04915	0.0608	-0.88965	*
TBE	CT	2954	58	0.00516	0.00626	-0.723	
xtr2_57342_57344	CT	684	19	0.01046	0.00895	0.71766	
mean		492.35	13.85	0.0114	0.0115	-0.4547	