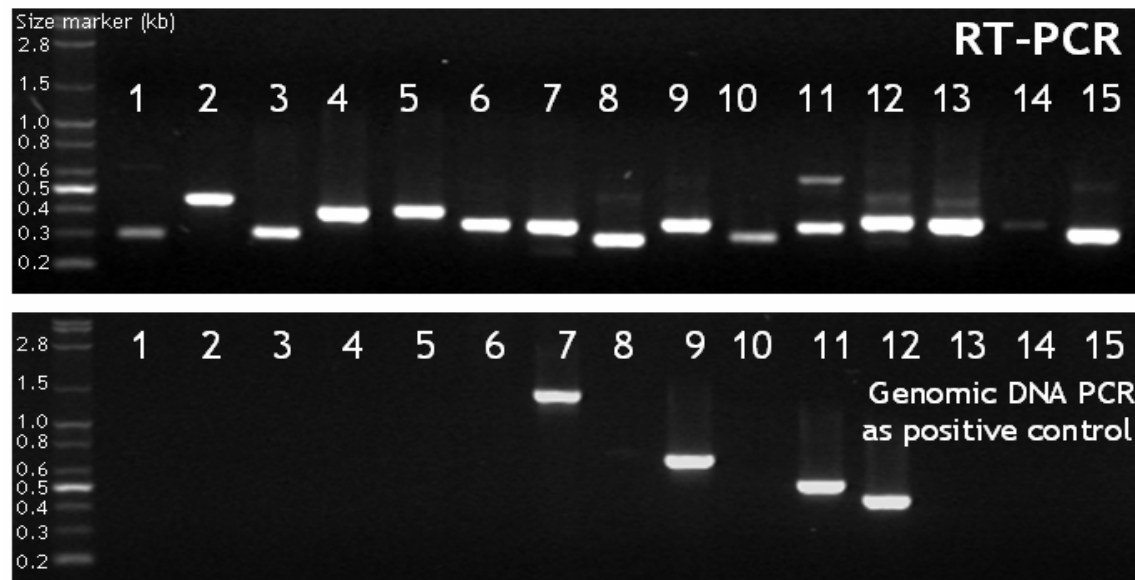


A. RT-PCR and genomic PCR



B. Information of primer pairs

No	AtRTPrimerID (gene name)	target cDNA	similar cDNA	gDNA
1	At4g16280.1f822r602 (FCA)	300	300,300	0
2	At5g13480.1f486r277 (FY)	448	0	0
3	At4g02560.1f891r51 (LD)	310	0	0
4	At2g19520.1f295r13 (FVE)	394	0	0
5	At4g00650.1f360r4 (FRI)	401	0	0
6	At5g10140.1f42r114 (FLC)	354	0	0
7	At3g18990.1f12r18 (VRN1)	350	0	1,531
8	At4g16845.1f200r290 (VRN2)	300	300	0
9	At1g09530.1f164r20 (PIF3)	355	355	753
10	At1g22770.1f1238r805 (GI)	304	0	0
11	At5g15840.1f178r94 (CO)	343	0	576
12	At1g65480.1f103r25 (FT)	366	0	490
13	At2g45660.1f57r30 (SOC1)	359	0	0
14	At5g61850.1f290r441 (LFY)	345	0	0
15	At1g69120.1f80r36 (AP1)	315	0	0

C. Evaluation of predictability

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
N*	G*	G	G	G	G	N	N	G	G	G	G	B*	G	N

* G: Good, N: Normal, B: Bad