



S. No.	VARIETY	PATHOTYPES																			Gene postulation
		11	11A	15-1	21	21A-2	24A	34-1	40A	40-2	40-3	122	117-1	117-3	117-4	117-5	117-6	184	184-1	295	
		1	2	3	20	4	5	6	7	8	9	10	12	13	14	15	16	17	18	19	
28	WH1105 (C)	R	R	R	R	R	R	R	MR	R	R	R	S	R	R	R	R	R	S	R	Sr2+11+
29	WH1124 (C)	S	S	S	R	R	MS	R	R	S	MR	MS	MS	MR	R	R	R	R	R	S	Sr2+7b+
30	WH1142 (C)	R	R	S	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	Sr2+31+
<b>North Eastern Plain Zone</b>																					
31	C306 (C)	S	R	R	S	R	S	S	R	S	R	R	S	R	S	R	R	R	MR	R	-
32	HD2888 (C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	Sr2+24+
33	K8027 (C)	S	R	R	R	R	S	R	MS	R	R	S	S	S	S	MS	S	S	S	S	Sr2+11+
<b>Central Zone</b>																					
34	HD4728 (D)	MS	R	R	R	R	S	MS	R	R	R	R	R	S	S	MR	S	S	MR	R	Sr2+11+
35	HD4730 (D)	R	R	R	R	R	S	MR	MS	R	R	R	S	S	S	MR	S	S	MR	R	Sr2+11+
36	GW322 (C)	MS	R	R	R	R	R	R	R	R	R	R	R	MR	R	R	R	R	R	S	Sr2+11+
37	HD2864 (C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	-*
38	HD2932 (C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	-
39	HI1544 (C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	Sr2+24+R+
40	HI8498 (D) (C)	MR	R	R	MS	R	S	R	R	R	S	S	R	S	MR	MR	MS	S	MS	S	Sr2+11+
41	HI8737 (D) (C)	MR	R	MS	R	R	MS	R	MS	R	S	S	R	S	R	S	MS	R	MS	S	Sr2+9e+
42	MP3336 (C)	S	MS	MS	R	R	R	R	MR	R	R	R	R	MR	R	R	R	R	S	MR	Sr2+
43	MP4010 (C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	Sr2+24+
44	MPO1215 (D) (C)	R	R	R	R	R	R	R	R	R	S	S	MR	R	-	R	S	R	R	R	Sr11+
<b>Peninsular Zone</b>																					
45	MACS3927 (D)	MS	R	R	R	R	MR	R	R	R	S	S	MS	S	R	R	MS	S	S	MR	Sr2+11+
46	NAW2030	R	R	R	R	R	R	S	R	R	R	R	R	R	R	R	R	R	R	R	Sr2+24+
47	AKDW2997-16 (D) (C)	R	MR	S	S	S	MR	R	R	S	MS	S	MR	S	S	R	S	S	S	S	Sr2+7b+
48	DBW93 (I) (C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	Sr2+31+
49	MACS6222 (C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	Sr2+31+
50	MACS6478 (C)	S	R	S	R	R	R	R	R	R	R	R	S	R	R	R	R	R	R	MS	-
51	NI5439 (C)	MS	R	R	R	R	S	R	R	R	R	R	S	R	R	R	R	R	R	R	Sr11+
52	NIAW1415(C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	Sr2+31+
53	UAS347 (I) (C)	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	S	R	Sr2+7b+11+
54	UAS428 (D) (C)	R	R	R	R	R	S	R	R	R	S	R	R	S	R	MR	S	S	R	R	Sr2+11+

