

Table S2 Analysis of a sample of *SRKb*-containing plants derived from the Col-0 *nripd1a-8[SRKb]* x C24 WT[*SRKb-SCRb*] cross.

F2 Plant	Pollination Tests ^a		<i>nripd1a</i>	<i>SRKb</i> Integration Number ^b	Genomic
			homozygote		Methylation ^c
1-7	+++	+++	yes	1	-
1-8	+++	+++	yes	2	-
1-9	0	0	yes	1	-
1-11	0	0	yes	2	-
2-1	0	0	no	2	+
2-3	0	3	yes	1	-
2-4	0	0	no	1	+
2-6	0	0	yes	1	-
2-8	0	0	no	2	+
2-10	0	0	no	2	+
3-1	0	0	no	2	+
3-2	0	0	no	2	+
3-3	0	0	no	2	+
3-4	0	0	no	2	+
3-5	0	0	no	2	+
3-7	0	0	no	2	+
3-8	0	8	yes	1	-
3-9	0	0	no	2	+
3-11	3	0	yes	1	-
3-12	+	20	yes	1	-
6-3	+	++	yes	1	-
6-4	0	0	no	2	+
6-5	0	0	no	2	+
6-6	0	0	no	2	+
6-7	0	0	no	2	+
6-8	0	0	no	2	+
6-10	0	X	no	2	+
6-11	0	0	no	1	+

^a Two replicate pollinations are shown. + indicates 20-50 pollen tubes per pollinated stigma, ++ indicates 50-75 pollen tubes per pollinated stigma, +++ indicates over 75 pollen tubes per pollinated stigma.

^b The number of *SRKb* transgene integrations

^c Determined by *AtSN1* methylation