

Table S2: binding sites in END-1 promoters				SKN-1	SKN-1core	POP-1	POP-1helper	PAL-1	GATA	MED-1	MED-1core
#	Name	Length	%AT	WWTRTCAT	RTCAT	CTTTGWWC	GCCRVNW	ATTATGAC	HGATAR	GTATACTYYY	RGTATAC
1	japonica	600bp	57.83	1	3	0	2	1**	3	1*	2**
2	afra	600bp	57.00	0	3	0	2	0	4*	0	0
3	sulstoni	600bp	54.17	0	4	0	2	0	4*	0	0
4	waitukubuli	600bp	57.83	0	2	0	2	0	3	0	3**
5	panamensis	600bp	56.00	0	2	0	2	0	2	0	0
6	nouraguensis	600bp	54.33	0	2	0	1	0	2	0	0
7	becei	600bp	54.17	1	2	0	3	0	2	0	0
8	kamaaina	600bp	63.50	1	4	0	2	0	3	1*	1
9	inopinata	600bp	58.67	0	0	0	6*	0	2	0	0
10	elegans	600bp	66.33	0	1	1	4*	0	3	2**	3**
11	doughertyi	600bp	63.17	0	3	0	3	0	4	0	0
12	tropicalis	600bp	63.50	0	1	0	0	0	5	0	2*
13	brenneri	600bp	65.83	0	2	0	1	0	4	0	0
14	remanei	600bp	62.17	0	2	0	0	0	1	0	0
15	latens	600bp	61.83	0	3	0	1	0	2	0	0
16	nigoni	600bp	63.50	0	2	0	1	0	2	0	4**
17	briggsae	600bp	61.67	0	1	0	2	0	1	0	1
18	sinica	600bp	61.50	0	1	0	1	0	6*	0	0
19	zanzibari	600bp	58.50	0	2	0	2	0	1	0	0
20	tribulationis	600bp	58.17	0	1	0	0	0	1	0	0

\*  $0.01 \leq p \leq 0.05$

\*\*  $p < 0.01$