SOM clusters affected in SAS mutants

value

PAR1_RNAi09.sun		1					4	1	1			
pif3.sun							3	3	2		1	
mida9_4.sun	1					1	5	1	7			
aos.sun		3	2		1	1	6	9	16	3	10	
co_9.sun							2	1	1			
hy5.sun	2	1				10	5	2	3			
jar1.sun		1	1				3		1	1	1	
kat1_2.sun		1					2	2				
phyB.sun pif45.sun spt_11.sun yuc2589.sun PAR1_RNAi09.shade pif3.shade mida9_4.shade aos.shade	10	6	1		1	3	7	10	8	1	1	
o pif45.sun		1	1	1			6	4	8			
. <u>⊆</u> spt_11.sun					1	1	5	7	6			
yuc2589.sun		1	1			3	2	8	3		1	
PAR1_RNAi09.shade							3	1	2			
pif3.shade							2	1	1			
mida9_4.shade						1	3	2	1			
o aos.shade		5					5	8	12	3	10	
co_9.shade								1				
hy5.shade	18	8	2		1	23	6	10	5		3	
jar1.shade		2				2	2	4	6	3	7	
kat1_2.shade		2				1	2	1	1			
phyB.shade	3	1		1	1		4	4	5		1	
pif45.shade	6	1		1		4	8	2	7			
spt_11.shade							1					
yuc2589.shade	11	4				7	3	6	4		1	
	10	7	12	~	_	4	2	∞	2	0	9	9
	genotype specific genes											