Organisation	Mx	Mf	Cc	Sa	Ad1	AdC	AK	AF	Sc	Ho
TRR,RTR	0230	07550	07813	0289						
T,RR,T,R	redC/0459	06400		7971						
RTRR,T	4445	30360	03338	4801						
RT,R,RT			00131							
R,RT				0461						
T,RTR			04616							
RT				5100						
TRT,TRT			01454	8144						
TRTTRT				6958						
RTR,TRRR										2075
TRRR,RT,TRT				1169						
TRRR,RT	6735	12790	07311							
TRT			05180							
TR,R,T	6953	11650	7521	0871						
T,R,TRR,RHRR			07950							
R,TRR,RHRR	7364									
T,R,RT		03710	00946	7006						
T,R,RT		04825		7451						
RRT,R,T	6966	11575								
RT,//,R,T			07558/07539	0856/0859						
T,RR,RT	4246	29365			1329	2626	2530	2438		
TR,R,T					2773	1193	2678			
TR,R								2297		
R,RT,T,R,RT				08017						
R,TRRH									1345	
TRT,R									2653	
RT,TR									4956	
TR,RTR									6827	
TRRH,R									7776	
T,R,TRH,RT									7809	
TR,R,RT									8715	

R,T,T,RR,R								8796	
T,T,RR								5168	
TR,R,RRRT									2367
RTRR,T									1854
TR,RT									0803
TRT,TR									0539

Additional Table 1: Intricate multi-gene TCS foci. Each row shows all myxobacterial orthologues of intricate multi-gene foci, including non-intricate orthologues, arranged by organism. Foci in particular genomes are denoted by the number from the locus tag of the first protein within the focus. '//' refers to an unspecified number (2 or more) of non-TCS genes.