



Heatmap visualization of protein-protein interaction scores between various proteins. The color scale ranges from -15 (cyan) to -5 (magenta).

	DD <sub>3</sub>	LL <sub>3</sub>	LD <sub>3</sub>	DD <sub>6</sub>	LL <sub>6</sub>	LD <sub>6</sub>	
	-12	-12	-12	-12	-12	-12	SAR116_0091 DNA recombination-mediator protein A (TIGR00730 family Rossman fold protein)
K03978 engB; GTP-binding protein	-12	-12	-12	-12	-12	-12	SAR116_1501 ribosome biogenesis GTP-binding protein YihA/YsxC
K03574 mutT; 8-oxo-dGTP diphosphatase [EC:3.6.1.55]	-12	-12	-12	-12	-12	-12	SAR116_1921 NUDIX hydrolase
	-12	-12	-12	-12	-12	-12	SAR116_2183 probable MutT/nudix family protein
K02316 dnaG; DNA primase [EC:2.7.7.101]	-12	-12	-12	-10	-12	-8	SAR116_2232 DNA primase
K06204 dksA; DnaK suppressor protein	-8	-8	-8	-12	-12	-12	SAR116_0086 dksA
K03060 rpoZ; DNA-directed RNA polymerase subunit omega [EC:2.7.7.6]	-8	-8	-8	-12	-8	-12	SAR116_0537 RpoZ/omega
K08311 nudH; putative (di)nucleoside polyphosphate hydrolase [EC:3.6.1.-]	-12	-12	-12	-12	-12	-12	SAR116_1846 RNA pyrophosphohydrolase
K03086 rpoD; RNA polymerase primary sigma factor	-8	-8	-8	-10	-8	-12	SAR116_2233 RpoD/sigma 70
K03043 rpoB; DNA-directed RNA polymerase subunit beta [EC:2.7.7.6]	-8	-8	-8	-8	-8	-8	SAR116_2384 RpoB
K03046 rpoC; DNA-directed RNA polymerase subunit beta' [EC:2.7.7.6]	-8	-8	-8	-8	-8	-8	SAR116_2385 RpoC
K03040 rpoA; DNA-directed RNA polymerase subunit alpha [EC:2.7.7.6]	-8	-8	-8	-12	-8	-12	SAR116_2415 RpoA