





logo

042	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYD	RHFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	ALMDEM	FAGMTLEEA	1180
11368	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYD	RHFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	ALMDEM	FAGMTLEEA	1180
APEC078	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYD	RHFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	ALMDEM	FAGMTLEEA	1180
BW2952	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYE	HFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	ALMDEM	FAGMTLEEA	1180
CE10	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	ATRPNAGLI	DLMDEM	FAGMTLEEA	1180
E2348/69	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	ATRPNAGLI	DLMDEM	FAGMTLEEA	1180
EDL933	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	DLMDEM	FAGMTLEEA	1180
IAI1	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYD	RHFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	ALMDEM	FAGMTLEEA	1180
IAI39	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	ATRPNAGLI	DLMDEM	FAGMTLEEA	1180
MG1655	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYE	HFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	ALMDEM	FAGMTLEEA	1180
REL606	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYD	RHFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	DLMDEM	FAGMTLEEA	1180
S88	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	ATRPNAGLI	DLMDEM	FASMTLEEA	1180
Sakai DNA	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	DLMDEM	FAGMTLEEA	1180
SMS-3-5	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	ATRPNAGLI	DLMDEM	FAGMTLEEA	1180
TW14359	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	DLMDEM	FAGMTLEEA	1180
UMN026	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYD	RHFGGVIYLF	FLRGVDKEHPQQGIY	TTRPNAGLI	DLMDEM	FAGMTLEEA	1180
UTI89	AMQAHRYDLQYQLYTLALHRYLRHRIADYDYER	HFGGVIYLF	FLRGVDKEHPQQGIY	ATRPNAGLI	DLMDEM	FASMTLEEA	1180
consensus	!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!	* * !!!!!!!!!!!!!!!!!!!!!!!	* !!!!!!!!!!!!!!!	* !!!!!!!!!!!!!!!	* !!!!!!!!!!!!!!!	* !!!!!!!!!!!!!!!	!