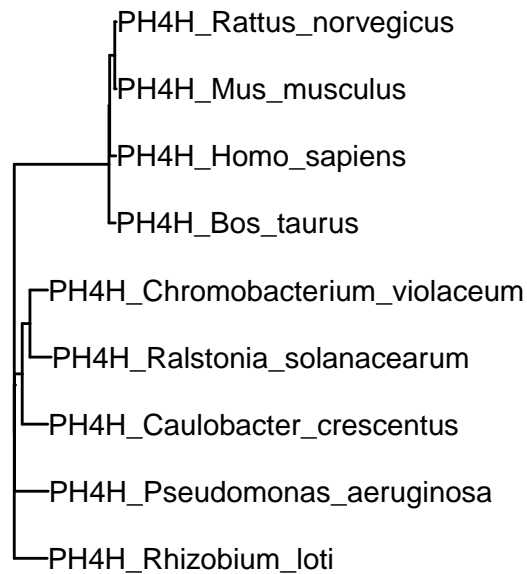


Tree

Multiple Sequence Alignment



QFADIAYNRYRHGQP	IPRVEY	TEEEKQTWGT	VFRTLKALYK	THACYE	HNHIFPLLE	EKYCGFR																																															
QFADIAYNRYRHGQP	IPRVEY	TEEERKTWGT	VFRTLKALYK	THACYE	HNHIFPLLE	EKYCGFR																																															
QFADIAYNRYRHGQP	IPRVEY	MEEEKK	TWGT	VFRTLKSLYK	THACYE	YNHIFPLLE																																															
QFADIAYNRYRHGQP	IPRVEY	TEEEKK	TWGT	VFRTLKSLYK	THACYE	HNHIFPLLE																																															
DAN----	DF	TL	PQPL	DRY	SAED	HA	TW	ATLY	QR	QCKL	LP	GRAC	DEF	MEGL	----	ER	LEV	D																																			
GL	DG	QTL	RP	DF	T	ME	QPV	HRY	T	A	AD	HA	TW	RTL	Y	DR	Q	E	ALL	P	GRAC	DEF	L	QGL	----	ST	L	GMS																									
GAR	----	PD	W	T	ID	QGW	E	T	Y	T	Q	A	E	H	D	V	W	I	T	LY	E	R	Q	T	D	M	L	H	GRAC	DEF	M	RGL	----	DA	L	D	L	H															
QPD	----	----	----	----	----	----	D	N	G	F	I	H	Y	P	E	T	E	H	Q	V	W	N	T	L	I	T	R	Q	L	K	V	I	E	G	RAC	Q	E	Y	L	D	G	I	----	----	EQ	L	G	L	P				
DYS	--	VC	R	A	D	F	T	V	A	Q	D	Y	D	----	Y	S	D	E	E	Q	A	V	W	R	T	L	C	D	R	Q	T	K	L	T	R	K	L	A	H	S	Y	L	D	G	V	----	----	E	K	L	G	L	----