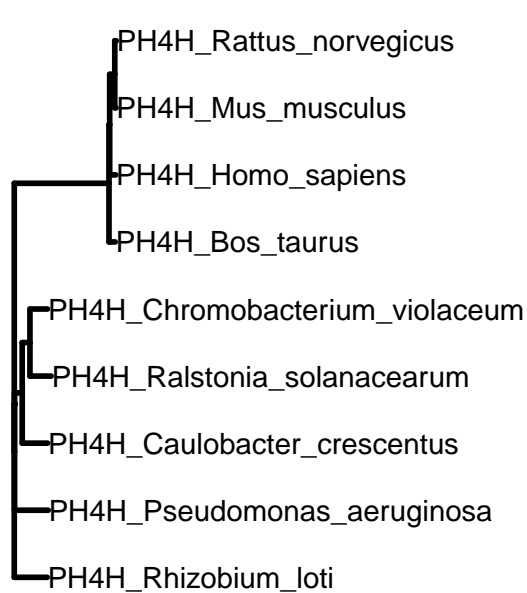


Tree

Multiple Sequence Alignment



QFADIAYNRYRHGQP	IPRVEY	TEEEKQ	TWGT	TVFRTL	KALYK	THACYE	HNHIF	PLLE	EKY	CGFR																																										
QFADIAYNRYRHGQP	IPRVEY	TEEERK	TWGT	TVFRTL	KALYK	THACYE	HNHIF	PLLE	EKY	CGFR																																										
QFADIAYNRYRHGQP	IPRVEY	MEEEKK	TWGT	TVFKTL	KSLYK	THACYE	YNHIF	PLLE	EKY	CGFH																																										
QFADIAYNRYRHGQP	IPRVEY	TEEEKK	TWGT	TVFRTL	KSLYK	THACYE	HNHIF	PLLE	EKY	CGFR																																										
DAN----	DF	TL	PQPL	DRY	SAED	HA	TW	ATLY	QRQCK	LLPGRAC	DEFMEGL	-----	ER	LEV	D																																					
GLDGQTL	RP	DF	TM	EQPV	HRYT	AAD	HA	TW	RTL	YDRQE	ALLPGRAC	DEF	LQGL	-----	ST	LGMS																																				
GAR-----	PD	WT	ID	QGW	ET	YT	QAE	HD	VW	ITLY	ERQ	TDML	HGRAC	DEF	MRGL	-----	DAL	DLH																																		
QPD-----	D	NGF	I	H	YP	E	TE	H	QVW	N	T	L	I	T	R	QLKV	IE	GRAC	QEY	LDGI	-----	E	Q	L	GLP																											
DYS--	V	C	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	H	S	Y	L	D	G	V	-----	E	K	L	GL	-