

**Table 3. Plasmid details**

<i>pralF</i>	1179ON2 1179C3	<b>GGAATTCGAGCTCAGACTAAAGGAGCAGATTATG</b> <b>GCTCTAGATCAAACCTATCGAGGCCATTTTC</b>	pMMB207	<i>EcoRI XbaI</i>
pSB- <i>ralF</i> -M45	1179ON2 1179OC	<b>GGAATTCGAGCTCAGACTAAAGGAGCAGATTATG</b> <b>CATGCCATGGAAAATTTAATTGTCTACCTTTTTC</b>	pSB616	<i>EcoRI NcoI*</i>
<i>pralF</i> -M45	Subcloned from pSB- <i>ralF</i> -M45		pMMB207	<i>EcoRI XbaI</i>
pM45- <i>ralF</i>	1179ON 1179C3	<b>CGGGATCCTTATGCATCCAGAAATTGAAAAAG</b> <b>GCTCTAGATCAAACCTATCGAGGCCATTTTC</b>	pMMB207- N45NT	<i>BamHI XbaI</i>
<i>pcya-ralF</i>	CyaNN CyaNCNotI	<b>GAAGATCTTACAGCAATCGCATCAGGCTGG</b> <b>CGGGATCCGCGGCCGCTGTCATAGCCGGAATCCTGG</b>	pM45- <i>ralF</i>	<i>BglIII BamHI<sup>†,‡</sup></i>
<i>pcya</i>	CyaNN CyaNCfs	<b>GAAGATCTTACAGCAATCGCATCAGGCTGG</b> <b>CGGGATCCCTGTCATAGCCGGAATCCTGG</b>	pM45- <i>ralF</i>	<i>BglIII BamHI<sup>†,‡,§</sup></i>
<i>pralF</i> <sub>1-305</sub> - <i>cya</i>	ClaXbaCya CyaPst	<b>GCATCGATTCTAGAATGCAGCAATCGCATCAGGC</b> <b>GCGCTGCAGTTAGTCATAGCCGGAATCCTGGC</b>	pM45- <i>ralF</i>	<i>ClaI PstI<sup>*,†</sup></i>
<i>pralF-cya</i>	Seq1485 RalFXbaNew	<b>GGACAAACCAAAGAGTCAATCG</b> <b>GCTCTAGAAAATTTAATTGTCTACCTTTTTC</b>	<i>pralF</i> <sub>1-305</sub> - <i>cya</i>	<i>ClaI XbaI</i>
<i>pralF</i> <sub>1-194</sub>	1174ON2 S7-194r	<b>GGAATTCGAGCTCAGACTAAAGGAGCAGATTATG</b> <b>CCGAGCTCTAGATTACAACCTCAAAGGTTTGGCTTTAATC</b>	pMMB207	<i>EcoRI XbaI*</i>
<i>pcya-ralF</i> <sub>1-194</sub>	1179ON1 pMMB2073	<b>CGGGATCCTTATGCATCCAGAAATTGAAAAAG</b> <b>CAGACCGCTTCTGCGTTCTG</b>	<i>pcya-ralF</i>	<i>BamHI PstI<sup>¶</sup></i>
<i>pralF</i> <sub>1-339</sub>	1179ON2 ralF+340	<b>GGAATTCGAGCTCAGACTAAAGGAGCAGATTATG</b> <b>GATCGTCTGACTTAGTTATCATATGCTTTTATTAAATC</b>	pMMB207	<i>EcoRI SalI</i>
<i>pcya-ralF</i> <sub>1-339</sub>	1179ON1 pMMB2073	<b>CGGGATCCTTATGCATCCAGAAATTGAAAAAG</b> <b>CAGACCGCTTCTGCGTTCTG</b>	<i>pcya-ralF</i>	<i>BamHI PstI<sup>¶</sup></i>
	Seq1485	<b>GGACAAACCAAAGAGTCAATCG</b>	<i>pcya-ralF</i>	<i>ClaI PstI</i>

<i>pcya-ralF</i> <sub>1-344</sub>	and 345C	GCGCTGCAGTTAAATGAGTTTTTCTGGTTATC		
<i>pcya-ralF</i> <sub>1-349</sub>	350C	GCGCTGCAGTTAGTTTCTCTCAATCGTAATGAG		
<i>pcya-ralF</i> <sub>1-354</sub>	355C	GCGCTGCAGTTACTCCTTAAGTGCCAGGTTTC		
<i>pcya-ralF</i> <sub>1-359</sub>	360C	GCGCTGCAGTTAATCTTTGGGAACGCCCTCC		
<i>pcya-ralF</i> <sub>1-364</sub>	365C	GCGCTGCAGTTACATTTACAGCGTCTGGATCTTTG		
<i>pcya-ralF</i> <sub>1-369</sub>	370C	GCGCTGCAGTTAACCTTTTTCTTTTTGCATTTTC		
<i>pcya-ralF</i> <sub>1-370</sub>	371C	GCGCTGCAGTTATCTACCTTTTTCTTTTTGCATTTTC		
<i>pcya-ralF</i> <sub>1-371</sub>	372C	GCGCTGCAGTTATTGTCTACCTTTTTCTTTTTG		
<i>pcya-ralF</i> <sub>1-372</sub>	373C	GCGCTGCAGTTATAATTGTCTACCTTTTTCTTTTTG		
<i>pcya-ralF</i> <sub>1-373</sub>	374C	GCGCTGCAGTTATTTTAATTGTCTACCTTTTTTC		
<i>pcya-ralF</i> <sub>16-374</sub>	16N	CGGGATCCTTTTCAATGCCAAGCCAAAAAATG	<i>pcya-ralF</i>	<i>Bam</i> HI <i>Pst</i> I
<i>pcya-ralF</i> <sub>222-374</sub>	222N	CGGGATCCTTTTCTTGCAATTCAACGGATGTAAAC		
<i>pcya-ralF</i> <sub>301-374</sub>	301N	CGGGATCCTTACCAAAGAGTCAATCGATTTG		
<i>pcya-ralF</i> <sub>340-374</sub>	340N	CGGGATCCTTCAGAAAAACTCATTACGATTG		
<i>pcya-ralF</i> <sub>345-374</sub>	345N	CGGGATCCTTACGATTGAGAGAAACCTGGC		
<i>pcya-ralF</i> <sub>350-374</sub>	350N	CGGGATCCTTCTGGCACTTAAGGAGGGCG		
<i>pcya-ralF</i> <sub>355-374</sub>	355N	CGGGATCCTTGGCGTTCCCAAAGATCCAGAC		
<i>pcya-ralF</i> <sub>360-374</sub>	360N	CGGGATCCTTCCAGACGCTGAAATGCAAAAAG		
<i>pcya-ralF</i> <sub>365-374</sub>	365N	CGGGATCCTTCAAAAAGAAAAAGGTAGAC		
<i>pcya-ralF</i> <sub>370-374</sub>	370N	CGGGATCCTTAGACAATTAATAATTTAATCC		
	and RalFC4	GCGCTGCAGTCTAGACAAACTATCGAGGCCATTTC		
	Seq1485	GGACAAACCAAAGAGTCAATCG	<i>pcya-ralF</i>	<i>Cla</i> I <i>Pst</i> I
<i>pcya-ralF</i> <sub>L372V</sub>	and 372V	GCGCTGCAGTTAAAATTTTACTTGTCTACCTTTTTCTTTTTG		

<i>pcya-ralF</i> <sub>L372F</sub>	372F	<b>GCGCTGCAGTTAAAATTTAAATTGTCTACCTTTTTCTTTT</b> GG		
<i>pcya-ralF</i> <sub>L372P</sub>	372P	<b>GCGCTGCAGTTAAAATTTGGTTGTCTACCTTTTTCTTTT</b> GG		
<i>pcya-ralF</i> <sub>L372S</sub>	372S	<b>GCGCTGCAGTTAAAATTTGATTGTCTACCTTTTTCTTTT</b> GG		
<i>pcya-ralF</i> <sub>L372T</sub>	372T	<b>GCGCTGCAGTTAAAATTTGTTTGTCTACCTTTTTCTTTT</b> GG		
<i>pcya-ralF</i> <sub>L372A</sub>	372A	<b>GCGCTGCAGTTAAAATTTGCTTGTCTACCTTTTTCTTTT</b> GG		
<i>pcya-ralF</i> <sub>K373E</sub>	373E	<b>GCGCTGCAGTTAAAATTCTAATTGTCTACCTTTTTCTTTT</b> GG		
<i>pcya-ralF</i> <sub>K373A</sub>	373A	<b>GCGCTGCAGTTAAAATGCTAATTGTCTACCTTTTTCTTTT</b> GG		
<i>pcya-ralF</i> <sub>K373R</sub>	373R	<b>GCGCTGCAGTTAAAATCTTAATTGTCTACCTTTTTCTTTT</b> GG		
pJV450- <i>cya-ralF</i>	Subcloned from <i>pcya-ralF</i>		pJV450	<i>EcoRI SacI/PstI</i> (blunted)

Forward primers are listed first followed by the reverse primer. Restriction enzyme sites used in plasmid construction are bold.

\*These plasmids represent intermediate products and were not used in the experiments described in the text.

†pMS107 was used as a PCR template.

‡PCR fragments were digested with *Bgl*III and *Bam*HI. The resulting fragments were ligated into *Bam*HI-digested pM45-*ralF*. The resulting plasmids in the right orientation have unique *Bam*HI sites at the junction of the *cya* and the *ralF* genes.

§*pcya* has a frameshift at the junction of the *cya* and the *ralF* genes. The resulting gene expresses a Cya protein with an 8-aa extension on the C terminus as shown in Fig. 1A.

¶*pralF*<sub>1-194</sub> and *pralF*<sub>1-339</sub> was used as a PCR template, respectively. The reverse primer (pMMB2073) anneals to DNA downstream of the cloning sites in these plasmids.