

Supplementary table 1. Sequence identity (%) at the amino acid level between different RaCIs

	RaCI2	RaCI3	RaCI6	RaCI7	RaCI4	RaCI1	RaCI5
RaCI2	100	41	38	42	40	42	38
RaCI3		100	71	73	44	35	36
RaCI6			100	76	47	36	38
RaCI7				100	45	35	37
RaCI4					100	38	35
RaCI1						100	40
RaCI5							100

Supplementary Table 2 Plasmids, primers and synthetic genes used in this study

Plasmid	Description	Restriction sites	Primers	Expression
pEpreS2-2	From Express2ion Biotechnologies, Denmark			<i>Drosophila melanogaster</i> S2
pMJ41	Modified pExpress2-2 vector; Kozak-BIP-His6-NgoMIV	Acc65I/EcoRI	MJ41 + MJ42 (annealed and inserted)	
pMJ80	pMJ41 His6-RaCl1	NgoMIV/NotI		
pMJ85	pExpress2-2 RaCl2 (native signal peptide)	EcoRI/NotI		
pMJ81	pExpress2-2 RaCl3 (native signal peptide)	EcoRI/NotI		
pMJ82	pExpress2-2 RaCl4 (native signal peptide)	EcoRI/NotI		
pMJ84	pExpress2-2 RaCl5 (native signal peptide)	EcoRI/NotI		
pMJ83	pExpress2-2 RaCl7 (native signal peptide)	EcoRI/NotI		
pMJ100	pETM14 RaCl1 (no signal peptide)	NcoI/NotI		
pMJ99	pETM14 RaCl2 (no signal peptide)	NcoI/NotI		
pMJ97	pETM14 RaCl3 (no signal peptide)	BspHI/NotI		
pMJ98	pETM14 RaCl4 (no signal peptide)	NcoI/NotI		
pMJ111	pETM14 - RaCl2 ΔC10 (for NMR experiments)		MJ127 + MJ128 (template = pMJ99)	
pMJ103	pETM14 - RaCl2 ΔN10		MJ87 + MJ88 (template = pMJ99)	
pMJ104	pETM14 - RaCl2 ΔC18		MJ149 + MJ150 (template pMJ99)	
pMJ15	pKLAC2-His10-OmCl wildtype ¹		MJ6 + MJ8	<i>Kluyveromyces lactis</i>
pMJ19	pKLAC2-His10-OmCl N78Q		MJ20 + MJ21 (template = pMJ15)	
pMJ27	pKLAC2-His10-OmCl N78Q/N102Q		MJ22 + MJ23 (template = pMJ19)	

Primer	Sequence	Description
MJ6	GCGCGGCGGCGCCCTAGCAGTCCTTGAGATGGGGAT	OmCl + stopcodon + NotI (rv)
MJ8	GGCCCCCTCGAGAAAAGAATGGCTAGCCATCACCATCACCATCACCATCACCATCACTCCGGAGACAGCGAAAAGCCGACTGCACTG	XhoI + His10 + OmCl (fw)
MJ20	CGGTGATGATGACGTTTAAGCAAGGCACAGACTGGGCTTCAACCG	OmCl - N78Q (fw)
MJ21	CGGTTGAAGCCAGTCTGTGCC TTGCT TAAACGTCATCATCACCG	OmCl - N78Q (rv)
MJ22	GGTAACGGCAACCCTTGGTCAACTAACCCAAATAGGGAAGTGG	OmCl - N102Q (fw)
MJ23	CCACTTCCCTATTTTGGGTAGTTGACCAAGGGTTGCCGTACC	OmCl - N102Q (rv)
MJ41	GTACCGCCACCATGAAGCTGTGCATCCTGCTGGCCGTGGTGGCCCTCGTGGGACTGAGCCTGGGACACCACCACCATCACCACGGCGCG	Acc65I-Kozak-BIP-His6-NgoMIV-EcoRI
MJ42	AATTCGCGGGCGTGGTGATGGTGGTGGTGTCCCAGGCTCAGTCCCAGGAAGGCCACCACGGCCAGCAGGATGCACAGTTCATGGTGGCG	Acc65I-Kozak-BIP-His6-NgoMIV-EcoRI

MJ87	CTGTTCCAGGGGCCCATGAAAGATCAGTGTGCAAATG	RaCl2 ΔN10 (fw)
MJ88	CATTTGCACACTGATCTTTATGGGCCCTGGAACAG	RaCl2 ΔN10 (rv)
MJ127	CACCGAAAAGCACCACCACCTAAACCGAACAGAGCTTTAATATG	RaCl2 ΔC10 (fw)
MJ128	CATATTAAGCTCTGTTCCGGTTAAGGTGGTGGTCTTTTCGGTG	RaCl2 ΔC10 (rv)
MJ149	CACCTGTTATCTGCTGGCATAAACACCGAAAAGCACCACC	RaCl2 ΔC18 (fw)
MJ150	GGTGGTCTTTTCGGTGTTATGCCAGCAGATAACAGGTG	RaCl2 ΔC18 (rv)

Synthetic gene ²	Sequence
RaCl1 (pMJ80)	CGAATTGGCGGAAGGCCGTC AAGGCCACGTGCTTTGTCCAGAGCTCGCCGGCGAGGAAGTGAAGACCACCCCATCCCCAACCC ACCAGTGCCTGAACGCCACGTGCGAGCGCAAGCTGGATGCCCTGGGCAATGCCGTGATCACC AAGTGCCACAGGGCTGCCTG TGCGTCTGTCGCGGAGCCAGCAATATCGTGCCCGCAACGGAACTGCTTCCAGCTGGCCACCACCAAGCCACCAATGGCCCC AGGGGATAACAAGGACAACAAGGAGGAGGAGTCCAAGTGA <u>CGCGCCGCGGT</u> ACCTGGAGCACAAGACTGGCCTCATGGGCCT CCGCTCACTGC
RaCl2 (pMJ85)	CGAATTGAAGGAAGGCCGTC AAGGCCACGTGCTTTGTCCAGAGCTCGAATTCGCCACCATGAACGCCGTGACCGTGTGGCCTT TACCGCCTTCGCCCTGATTGTGCACGATTGCTACAGCGAGGAGGCCAACACCACCCCATCAGCGTGAAGGATCAGTGCGCCAA CGTGACCTGCCGCGCACCGTGGATAATCGCGGCAAGCGCCACATCGATGGCTGCCACCAGGATGCCCTGTGCGTGTGAAGG GCCCGATAGCAAGGATAACCTGGATGGCACGTGCTACCTGCTGGCCACCACCCCAAAGAGCACCACCACAGTACCCGAGCAG AGCTTCAACATGGAGGAGTAGCGCGCCGCGGTACCTGGAGCACAAGACTGGCCTCATGGGCCTTCTTTCACTGC
RaCl3 (pMJ81)	CGAATTGGCGGAAGGCCGTC AAGGCCACGTGCTTTGTCCAGAGCTCGAATTCGCCACCATGGCCGCCCTGAATGGCCTGGTGTG TCTGCTGCTGACCATCAGCGCCATGTTTCATCAGCGAGTGTACAGCAGCGCGGAGAGCCAGAGCATTCAAGCGCAAGGACAGT GCGAGGAAGTGTATGTCACCCGAAGCTGAACCACTGGGAGAGCGCGTGACCAAGCGGATGCCAACCGGATGCCCTGTGCGTG ATCCGCGAGCCCGATAACGTGGACAACGCCAACGGAACCTGCTACGCGCTGATGAGCAGCACCACCACGACCACGACGACCC CAGATGGAACCAACCTCCGAAGGAGGAGTAGCGCGCCGCGGTACCTGGAGCACAAGACTGGCCTCATGGGCCTTCCGCTC ACTGC
RaCl4 (pMJ82)	CGAATTGGCGGAAGGCCGTC AAGGCCACGTGCTTTGTCCAGAGCTCGAATTCGCCACCATGAGCGCCTTCAACATCTTCGCCCT GGTGCTGGTGTGTGCGCCCTGATGATCAACGAGTGTGCTGACCAAGCCAGGAGCCACCACCCACTGAAGGCCGCCAGTCACT GACGCAACGTGAAGTGCAGCGCCGCTTCGATCAGCTGGGCAATAGTGTGACCGAGGGGTGCCAAAGCGGATGCCCTGTGCGTG TACCAGGCCACCGGTACAACCAAGGAGGCCAACGGAACCTGCTACGAGCTGATGAAGACTCGACCACCAGCAGCAGCCGAGGG AACCCAGCCCATAGCGCGCCGCGGTACCTGGAGCACAAGACTGGCCTCATGGGCCTTCCGCTCACTGC
RaCl5 (pMJ84)	CGAATTGGCGGAAGGCCGTC AAGGCCACGTGCTTTGTCCAGAGCTCGAATTCGCCACCATGAACGCCGTGATCGTGTGCGT GACCATCAGCGCCGTGCTGATCCACCAGTGTACAGTACCCGCGGAGCCACCCTGAGCATCAAGCGCGGCGATATGTGCATCG AGAAGACCTGCAACCGCTCCATCGATGCCCGGGAAGAAAGTGTGCGCGGATGCCAGGCGGCTGCCTGTGCGTGTTC AACGTGTCCGATGTGACCTACCCCGCAACGGCACGTGCTATCAGCTGGCCACCACCACGACCAATCGCCCAAGCGCCGCTGATGGA GCGCGAGCGCTAAGCGGCCGCGGTACCTGGAGCACAAGACTGGCCTCATGGGCCTTCCGCTCACTGC
RaCl7 (pMJ83)	CGAATTGGCGGAAGGCCGTC AAGGCCACGTGCTTTGTCCAGAGCTCGAATTCGCCACCATGGCCGCCCTGAATGGCCTGGTGTG TCTGCTGCTGACCATCAGCGCCATGTTTCATCAGCGAGTGTACAGCAGCGCGGAGAGCCAGAGCATTCAAGCGCAAGGACAGT GCGAGGAAGTGAAGTGCAGCGCCACCGTGAATCAGCTGGGAGTGGCCGTGACCAAGCGGATGCCAAAGTGGATGCCCTGTGCGTG ATCAGCGCCCGAGATAGCGCCGTGAACGTGAACGGCACGTGCTACCAAGCTGATGGGCTCCACCAGTACCACCACCAGCAGCAC CCCAAGCAGCGAGGATCAGGAGTAGCGCGCCGCGGTACCTGGAGCACAAGACTGGCCTCATGGGCCTTCCGCTCACTGC
RaCl1 (pMJ100)	CGAATTGAAGGAAGGCCGTC AAGGCCACGTGCTTTGTCCAGAGCTCCCATGGAAGAAGTTAAACCACCCGATTCCGAATCATC AGTGTGTTAATGCAACCTGTGAACGTAAACTGGATGCACTGGGTAATGCAAGTATTACCAATGTCGCGAGGGTGTCTGTGTGTT GTTCGTGGTGCAAGCAATATTGTTCCGGCAAAATGGCACCTGTTTTACGCTGGCAACCACCAACCGCCTATGGCACGGGTGATA ATAAAGATAACAAGAAGAAGAAAGCAATTAAGCGGCCGCGGTACCTGGAGCACAAGACTGGCCTCATGGGCCTTCTTTCACTG C
RaCl2 (pMJ99)	GGCCCCATGGAGGAAGCAAA TACCACCCGATTAGCGTTAAAGATCAGTGTGCAAATGTTACCTGTCGTCGTACCGTTGATAAT CGTGGTAAACGTCATATTGATGGTTGTCCGCTGGTTGTCTGTGTGTTCTGAAAGGTCCGGATAGCAAAGATAATCTGGATGGCA CCTGTTATCTGCTGGCAACCACCCGAAAAGCACCACCACAGTACCGAACAGAGCTTTAATATGGAAGAGTAAGCGGCCGCGC GCG
RaCl3 (pMJ97)	GGCCCTCATGAGCGGTGAAAG CCGAGAGCATTACGCGTAAAGGTGAGTGTGAAGAAGTTATCTGTCATCGTAAACTGAATCATCTG GGTGAACGTGTACCAGCGGTTGTCCGACCGGTTGTCTGTGTGTTATTCTGTAACCGGATAATGTGATAATGCAAAATGGCACCT GTTATGCACTGATGAGCAGCACCACCAACCACCACGACCCGATGTTACAACCACCTCTGAAGAAGAAGATAAAGCGGCCG CGCGCG
RaCl4 (pMJ98)	GGCCCCATGGCACAAGAACC GACCACCCGCTGAAAGCAGCAAGCCAGTGTAGCAATGTTAAATGTCGTCGTGCTTTTGTATCAT CTGGGTAATAGCGTTACCGAAGGTTGTCGAGCGGTTGTCTGTGTGTTTATCAGGCAACCGGTTATAATCAAGAAGCAAACGGCA CCTGTTATGAGCTGATGAAAACCAAGTACCACCACCACAACCAGGTTACACCGGCACAGTAAGCGGCCGCGCGCG

¹Amplified from pMETαC-OmCl (Nunn, M.A. et al., *J. Immunol* **174**, 2084-2091, 2005).
²Synthetic fragments were subcloned into constructs that are indicated between brackets.
Restriction sites are underlined.

Mutations introduced with primers are in bold.
Fw = forward primer, rv = reverse primer.