Additional File 4

	70	80	90	360	100	110		120	130	140	150		160	
Streptomyces coelicolor A3(2) ScbR	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVELS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVVI	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
Streptomyces lividans TK24 ScbR	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENG	ELLPHVVT	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
M132 scbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVVI	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
505 ScbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVVI	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
290 scbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVV1	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
11 scbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVVI	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
380 scbRrt1 Gottelt 1 HS	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVV1	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
A3(2)N3 scbRrt1 Gottelt 5 HS	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVVT	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
J1501 scbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVVT	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
N1 scbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVV1	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
13 scbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENG	ELLPHVV1	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
210 scbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENO	ELLPHVVI	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
CH999 scbRrt1	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RRWHETLL	KLLNQAKENG	ELLPHVVI	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
M600 scbR2	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RSWHETLL	KLLNQAKENO	ELLPHVV1	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
A3(2) (Eriko Norwich) scbRrt	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RSWHETLL	KLLNQAKENO	ELLPHVVI	TDSADLY	VGTFAG	IQVVSQT	VSDYQDL
A3(2)N2 scbRrt1 Gottelt 3 HS	QAVPEQP	LRLQELII	DMGMLFCHR	RTNVVAF	AGVRLS	MDQQAHGL	DRRGPF	RSWHETLL	KLLNQAKENO	ELLPHVVT	TDSADLY	VGTFAG	IQVVSQT	VSDYQDS