

Bacterial	-----mnitnlcpfgevfnatrfasvyawnrkrisncvadysvlynsasfstfk	49
Eukaryotic	rvqptesivrfrpmitnlcpfgevfnatrfasvyawnrkrisncvadysvlynsasfstfk	60

Bacterial	cygvsptklndlcftnvysdfvirgdevrqiapgqtgkiadynyklpddftgcviawns	109
Eukaryotic	cygvsptklndlcftnvysdfvirgdevrqiapgqtgkiadynyklpddftgcviawns	120

Bacterial	nnldskvggnynylyrlfrksnlkpferdisteiyaqgstpcngvegfcyflqsygfq	169
Eukaryotic	nnldskvggnynylyrlfrksnlkpferdisteiyaqgstpcngvegfcyflqsygfq	180

Bacterial	ptngvgyqpyrvvvlsfellhapatv-----	195
Eukaryotic	ptngvgyqpyrvvvlsfellhapatvcgpkkstnlvknkcvnf	223

Figure S1. Alignment of the RBD amino acid sequences used for bacterial expression in BL21(DE3) cells or eukaryotic expression in mammalian HEK293T cells. Asterisks show identical amino acids.