**Supplementary File 2. List of oligonucleotides used for plasmid construction**

Gene NameOligonucleotides used for plasmid construction

*MS2-GFP(x4)* 5'-GACTAGTCGGATGCAAGGGTTCGAATCCCTTAGCTCTC-3'

5'-CCGCTCGAGGGCCGCAAATTAAAGCCTTCGAGCGTCC-3'

*TIM50-Tag* 5'-CCTTATTTGAAGAGGAAAAGAAAAAGAAGAAGATTGCTGAATCCAAATAAA  
ACGCTGCAGGTCGACAACCC-3'

5'-ACATACACACATAGATACGTAGATACATGAGAAGAGGGTTTACATGAAAAG

CATAGGCCACTAGTGGATC-3'

*RPL2-Tag* 5'-ATAGAATGAAAGTAAAAGACAGACCAAGAGGCAAAGATGCAAGATTATAAA

ACGCTGCAGGTCGACAACCC-3'

5'-AACGAAGAAAAGGTCCCTATGTTGCTTCTTTTCTACTAAAATTAGCCTATGCA

TAGGCCACTAGTGGATC-3'

*ATP3-Tag* 5'-TTACTAATGAACTGGTTGATATTATTACTGGTGCTTCCTCTTTGGGATGAAAC

GCTGCAGGTCGACAACCC-3'

5'-TTCTACAAAAACAACGTCAAATAAAGAGGCAATGCAGGGTGATTTTTTTAGC

ATAGGCCACTAGTGGATC-3'

*TOM22-Tag* 5'-ATAACATATTGGCCCAAGGTGAAAAAGATGCTGCAGCAACAGCCAATTAAA

ACGCTGCAGGTCGACAACCC-3'

5'-ATGTATGGCTCCTTTTCTAAAACCCTCTCTTTTCTTTTACATCATTAAAAGCAT  
AGGCCACTAGTGGATC-3'

*ATP1-Tag 5'-GTTGGCATCTCTAAAGAGTGCTACTGAATCATTTGTTGCCACTTTTTAAAACGCTGCAGGTC*

*GACAACCC-3'*

*5'-ATTTCTTTTTGAGACGTACCTTATATTCATTTTTATTTTTTTAGTTCACAGCATAGGCCACTAG*

*TGGATC-3'*

*ATP2-Tag 5'-TGAAGATGTTGTTGCTAAAGCTGAAAAGTTAGCCGCTGAAGCCAACTAGAACGCTGCAGGT*

*CGACAACCC-3'*

*5'-TACCTTCGGTATTTCAAATTTTGCTTCCCTTGGTTTAAGCTTTATTTCTTGCATAGGCCACTA*

*GTGGATC-3'*

*ATP7-Tag 5'-GGACGTACCTGGTTACAAGGACAGATTCGGCAATTTGAATGTGATGTAGAACGCTGCAGGT*

*CGACAACCC-3'*

*5'-TGTGAAAAAAATAATAGAATATGGTGCGTAATATATAGAGGTAAAGGGTAGCATAGGCCACT*

*AGTGGATC-3'  
GPDp Sac2 F-680* 5'-TCCCCGCGGCAGTTCGAGTTTATCATTATCAATACTGCC-3'

*GPDp Hind3 R-1* 5'-CCCAAGCTTTTTGTTTGTTTATGTGTGTTTATTCGAAAC-3'

*color START Spe1 F* 5'-GACTAGTCGGTGACGGTGCTGGTTTA-3'

*color ADHt Kpn1 R* 5'-GGGGTACCTTACCCTGTTATCCCTAGCGGATCTGCCGG-3'

*C450\_2C1\_top* 5'-ATGGATCCAGTTGTCGTATTGGGTTTATGTTTGTCTTGTTTGTTACTACTTTCT  
TTGTGGAAACAATCTTACGGTGGCGGAAAGTTA-3'

*C450\_2C1\_bot* 5'-TAACTTTCCGCCACCGTAAGATTGTTTCCACAAAGAAAGTAGTAACAAACAA  
GACAAACATAAACCCAATACGACAACTGGATCCAT-3'

*TIM50 F-500 Spe1* 5'-CGGTGGCGGCCGCTCTAGAACTAGTACTGTGTTGGCTTTAACTCTTTAAATTC  
TCC-3'

*TIM50 R++500 Xho1* 5'-AATTGGGTACCGGGCCCCCCCTCGAGATGTGACGGCAGTTCCTGACCTGATA

GTG-3'

*GA\_TIM50\_F129* 5'-CAATCTTACGGTGGCGGAAAGTTACAAAAAGAAACAAAAGACGACAAGCCT

AAATC-3'

*GA\_TIM50\_R-1* 5'-ACCCAATACGACAACTGGATCCATTGCAAGCGGGTGATTTTTGGAAGTTTATT

CTAGC-3'

*TOM22 F-500 Spe1* 5'-CGGTGGCGGCCGCTCTAGAACTAGTCAAAAAGAGCTAATCAACTCCTTGAAC

TTAG-3'

*TOM22 R++500 Xho1* 5'-AATTGGGTACCGGGCCCCCCCTCGAGGTTTACGTTTTAGATTACCAAAAAGG

AAGCATAG-3'

*GA\_TOM22\_F1* 5'-CAATCTTACGGTGGCGGAAAGTTAATGGTCGAATTAACTGAAATTAAAGACG

ATGTC-3'

*GA\_TOM22\_R-1 5'-ACCCAATACGACAACTGGATCCATTTGAATGATGCTTATTTTGGGGTATATAG*

*TTCCG-3'*

*GA\_TIM50\_F562 5'-GAGCCACCTTTCCCTGATTTACTATACCAAAGGCCATTAACTCTTGTTATCACA-3'*

*TIM50\_R540 5'-TAGTAAATCAGGGAAAGGTGGCTCTTGGA-3'*

*yeGFP-CaaX\_F 5'-GGTAAGGCTAGCGGTAAAAAGAAGAAAAAGAAGTCAAAGACAAAGTGTGTAATTATGTAAC*

*TGGTCGAGTCATGTAATTAGTTATGTC-3'*

*yeGFP-CaaX\_R 5'-CTTTTTACCGCTAGCCTTACCAGCACTGCCTGCGCTATCGCTACCTTTGTACAATTCATCCA*

*TACCATGGGT-3'*