**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'*