

1 nnnn: vector backbone
2 NNN: homology arm/intergenic region
3 NNN: homology arm/coding region
4 NNN: attP
5 NNN: Gal4
6 NNN: mini-white

7
8 GTCCGAACAACATGGTCATAGTGCAGGTATTATAGTGTAAAAGCCAAAAGTGAGAAAACCA
9 GCAGCCAAAGGACTACGCCACGAGCATACAAATGCCAGCCATGCCGTCGCAACTTGTGATA
10 AGCGGCACCGCCGGCTTACCAGCAAATAAACAGCGAACACAGGGACACCAGCCAGCAAAC
11 TCGAGACGCAGCAGCAGCAAACGCAGCAGGATACCAGCCAGGAGCAGCAGCAGCAGCA
12 GCAGCATCAGCATTCGCACAGCCAACTctagaCCCAGGTCAGAAGCGGTTTTCGGGAGTAG
13 TGCCCCAACTGGGGTAACTTTTGAGTTCTCTCAGTTGGGGGCGTAGGGTCGCCGACATGAC
14 ACAAGGGGTTagatctATGAAGCTGCTGAGTAGTATTGAACAAGCGTGTGACATATGCCGA
15 CTTAAGAAGCTGAAGTGCAGCAAGGAAAAGCCGAAGTGCGCCAAATGTCTTAAGAACAATT
16 GGGAGTGCCGGTACTCGCCAAAGACGAAGCGAAGCCCTTTGACACGCGCTCATTTGACTGA
17 AGTGGAGAGTCGCCTGGAGCGCCTTGAGCAGTTGTTCCCTCCTCATTTTTCCCGTGAAGAC
18 CTCGACATGATACTGAAGATGGATAGTCTCCAAGATATTAAGGCCCTGCTCACTGGCCTGT
19 TTGTTCAGGATAACGTCAACAAGGATGCCGTGACGGACCGCCTTGCCCTCGGTGGAGACAGA
20 TATGCCACTTACCTTGCGGCAGCATCGCATCTCGGCTACCAGCAGTAGTGAAGAAAGTTCC
21 AATAAGGGTCAACGGCAACTGACGGTATCGATCGACAGTGCGGCCACCACGACAACAGCA
22 CCATTCCACTGGACTTCATGCCTCGTGACGCACTGCATGGATTTGACTGGAGCGAGGAGGA
23 TGATATGAGCGACGGACTCCCATTCCTGAAGACCGACCCCAATAACAATGGTTTCTTTGGC
24 GACGGCAGCCTGCTGTGTATCCTGCGAAGTATTGGTTTTAAGCCGGAAAATTACACGAACA
25 GTAATGTCAATCGTCTTCCCACAATGATCACCGACCGTTACACGCTGGCCAGTCGCAGTAC
26 TACAAGCCGCCTGCTGCAAAGCTATCTTAACAACCTCCACCCCTACTGCCCCATAGTGCAT
27 AGTCCTACTCTCATGATGCTTTACAACAATCAGATTGAGATAGCATCGAAGGACCAGTGGC
28 AGATCCTCTTCAACTGCATTCTCGCCATCGGGCCTTGGTGTATTGAAGGCGAGAGCACAGA
29 TATCGACGTGTTTTACTATCAAAACGCCAAGTCGCACCTCACCTCGAAGGTCTTCGAAAGT
30 GGAAGCATTATCCTCGTGACAGCCCTCCATCTGCTGAGTCGTTATACCCAGTGGCGTCAGA
31 AAATAACATCCTACAACCTTTCATTCGTTTCAGCATTTCGGATGGCGATTAGCCTGGGACT
32 TAATCGTGACCTCCCAAGTAGCTTCAGCGATTTCGTCGATCCTGGAACAGCGCCGGAGGATT
33 TGGTGGAGTGTGTACAGCTGGGAGATCCAGCTGTCCCTCTTGTACGGACGCAGCATTACAGC
34 TGAGTCAGAACACGATTTCCCTTCCCGTCCAGCGTGGACGACGTTCAACGGACAACACTACGGG
35 CCCGACCATCTACCACGGTATCATTGAGACGGCCCGCTTGCTTCAAGTGTTTTACGAAGATT
36 TATGAGCTGGATAAGACAGTGACCGCGGAGAAGAGTCCGATCTGCGCCAAGAAATGCCTGA
37 TGATCTGCAATGAGATAGAGGAGGTGAGCCGCCAGGCACCCAAGTTCCCTGCAGATGGACAT
38 AAGCACTACGGCGCTGACTAATCTCCTCAAAGAACACCCCTGGCTGAGTTTTACCCGCTTC
39 GAGCTGAAGTGGAAACAGCTGAGTCTGATTATATACGTTTTGCGGGACTTTTTACAAATT
40 TTACGCAGAAGAAGTCGCAGCTTGAACAAGATCAGAACGACCATCAGTCTACGAGGTAAA
41 GAGGTGCAGCATAATGCTCTCCGACGCAGCCAGCGCACCGTGATGTGCGGTGTCGTCCTAT
42 ATGGACAACCATAATGTGACCCCTTATTTGCCTGGAATTGCTCCTACTATTTGTTCAACG
43 CCGTGTGGTCCCAATCAAGACTCTTCTGAGCAACTCCAAATCGAACGCCGAGAATAACGA
44 GACAGCACAGCTCTTGACGCAAATTAACACCGTGTGATGCTGTTGAAGAAACTCGCTACA
45 TTCAAGATTCAGACTTGCGAAAAGTACATCCAGGTGTTGGAAGAGGTGTGCGCCCCATTCC
46 TTCTTTTCGCAATGCGCAATTCCTCCCGCACATAAGTTACAATAATTCACACGGCTCCGC
47 TATCAAGAATATCGTTCGGCTCCGCTACAATCGCGCAGTATCCACGCTGCCTGAGGAAAAT
48 GTTAACAACATATCGGTGAAGTATGTAAGTCCAGGCAGCGTCGGCCCGAGCCCTGTGCCGC

49 TGAAAAGCGGAGCCTCCTTCAGCGATCTGGTCAAGCTGCTCAGCAACCGCCCACCATCCCG
50 CAACTCCCCTGTCACTATAACCCCGATCGACCCCGTCGCATCGTAGCGTTACCCCATTCCTG
51 GGACAACAGCAGCAACTGCAGAGTCTGGTGCCACTTACGCCCTCGGCATTGTTTGGCGGAG
52 CCAACTTCAACCAGAGTGGAACATTGCGGACAGCTCCTTGAGCTTACGTTTACCAACAG
53 CTCCAACGGTCCAAACTTGATCACGACTCAGACAAACTCCCAGGCATTGTCCCAGCCCATT
54 GCCAGCTCGAATGTGCATGATAACTTTATGAATAATGAAATTACTGCCTCCAAGATTGACG
55 ACGGCAACAACAGCAAGCCCCTTTCCCCAGGCTGGACTGACCAGACGGCCTACAATGCTTT
56 CGGTATCACCCTGGCATGTTCAACACCACAACGATGGACGACGTATATAACTATCTTTTT
57 GATGATGAGGATACTCCACCGAACCCAAAGAAGGAGTAGgatctataacttcgtataatgt
58 atgctatacgaagtattatggtaccttgcatgccccgttattcTCTATTCTCTATTTCGTTTTG
59 TGACTCTCCCTCTGTACTATTGCTCTCTCACTCTGTGCGACAGTAAACGGCAGCTATT
60 CTCGTTGCTTCGAGAGAGCGCGCCTCGAATGTTTCGCGAAAAGAGCGCCGGAGTATAAATAG
61 AGGCGCTTCGTCGACGGAGCGTCAATTCAATTCAAACAAGCAAAGTGAACACATCGCGAAG
62 CGTAAGCTGAGCAAACAAACAAGCGCAGCTGAACAAGCTAAACAATCTGCAATAAAGTGCA
63 AGTTAAAGTGAATCAATTAAGTAACCAACAACCAAGTAATTAAGTAAACTAAAACTGCAACT
64 ACTGAAATCAACCAAGAAGTCATTATTGAAGACAAGAAGAGAACTCTgaataggtcgatag
65 cgtcaattcggcacgagcgggtttcgtgacgaagctccaagcggtttacgccatcaattaa
66 caciaagtgctgtgccccaaactcctctcgcttcttatttttGTTTGTTTTTTGAGTGATTG
67 GGTGGGTGATTGGTTTTGGGTGGGTAAGCAGGGGAAAGTGTGAAAAATCCCGGCAATGGGC
68 CAAGAGGATCAGGAGCTATTAATTCGCGGAGGCAGCAAACACCCATCTGCCGAGCATCTGA
69 ACAATGGTGACAGCGGAGCGGCTTCGCAGAGCTGCATTAACCAGGGCTTCGGGCAGGCCAA
70 AACTACGGCAGCTCCTGCCACCAGTCCGCCGGAGGACTCCGGTTCAGGGAGCGGCCAA
71 CTAGCCGAGAACCTCACCTATGCCTGGCACAATATGGACATCTTTGGGGCGGTCAATCAGC
72 CGGGCTCCGGATGGCGGCAGCTGGTCAACCGGACACGCGGACTATTCTGCAACGAGCGACA
73 CATAACGGCGCCCAGGAAACATTTGCTCAAGAACGTTTGCGGCGTGGCCTATCCGGGCGAA
74 CTTTTGGCCGTGATGGGCAGTTCCGGTGCCGAAAGACGACCCTGCTGAATGCCCTTGCC
75 TTCGATCGCCGAGGGCATCCAAGTATCGCCATCCGGGATGCGACTGCTCAATGGCCAACC
76 TGTGGACGCCAAGGAGATGCAGGCCAGGTGCGCCTATGTCCAGCAGGATGACCTCTTTATC
77 GGCTCCCTAACGGCCAGGGAACACCTGATTTTCCAGGCCATGGTGCAGGATGCCACGACATC
78 TGACCTATCGGCAGCGAGTGGCCCGGTGGATCAGGTGATCCAGGAGCTTTCGCTCAGCAA
79 ATGTCAGCACACGATCATCGGTGTGCCCGGAGGGTGAAAGGTCTGTCCGGCGGAGAAAGG
80 AAGCGTCTGGCATTGCGCTCCGAGGCACTAACCAGTCCGCCGCTTCTGATCTGCGATGAGC
81 CCACCTCCGGACTGGACTCATTTACCGCCACAGCGTCTGTCAGGTGCTGAAGAAGCTGTC
82 GCAGAAGGGCAAGACCGTCATCCTGACCATTATCAGCCGTCTTCCGAGCTGTTTGGAGCTC
83 TTTGACAAGATCCTTCTGATGGCCGAGGGCAGGGTAGCTTTCTTGGGCACTCCCAGCGAAG
84 CCGTGCAGTCTTTTCTACGTGGGTGCCAGTGTCTACCAACTACAATCCGGCGGACTT
85 TTACGTACAGGTGTTGGCCGTTGTGCCCGGACGGGAGATCGAGTCCCGTGATCGGATCGCC
86 AAGATATGCGACAATTTTGCCATTAGCAAAGTAGCCCGGATATGGAGCAGTTGTTGGCCA
87 CCAAAAATTTGGAGAAGCCACTGGAGCAGCCGGAGAATGGGTACACCTACAAGGCCACCTG
88 GTTCATGCAGTTCCGGGCGGTCTGTGGCGATCCTGGCTGTGCGGTGCTCAAGGAACCACTC
89 CTCGTAAGAGTGCAGCTTATTCAGACAACGATGGTTGCCATCTTGATTGGCCTCATCTTTT
90 TGGGCAACAACCTCACGCAAGTGGGCGTGATGAATATCAACGGAGCCATCTTCTCTTCT
91 GACCAACATGACCTTTCAAACGTCTTTGCCACGATAAATGTGTTACCTCAGAGCTGCCA
92 GTTTTTATGAGGGAGGCCCGAAGTCGACTTTATCGCTGTGACACATACTTTCTGGGCAAAA
93 CGATTGCCGAATTGCCGCTTTTTCTCACAGTGCCACTGGTCTTACGGCGATTGCCTATCC
94 GATGATCGGACTGCGGGCCGGAGTGCTGCACTTCTTCAACTGCCTGGCGCTGGTCACTCTG
95 GTGGCCAATGTGTCAACGTCCTTCGGATATCTAATATCCTGCGCCAGCTCCTCGACCTCGA

96 TGGCGCTGTCTGTGGGTCCGCCGTTATCATAACCATTCCTGCTCTTTGGCGGCTTCTTCTT
97 GAACTCGGGCTCGGTGCCAGTATACCTCAAATGGTTGTCGTACCTCTCATGGTTCCGTTAC
98 GCCAACGAGGGTCTGCTGATTAACCAATGGGCGGACGTGGAGCCGGGCGAAATTAGCTGCA
99 CATCGTCGAACACCACGTGCCCCAGTTCGGGCAAGGTCATCCTGGAGACGCTTAACTTCTC
100 CGCCGCCGATCTGCCGCTGGACTACGTGGGTCTGGCCATTCTCATCGTGAGCTTCCGGGTG
101 CTCGCATATCTGGCTCTAAGACTTCGGGCCCCGACGCAAGGAGTAGCCGACATATATCCGAA
102 ATAAGTGTGTTTTTTTTTTTTTTTACCATTATTACCATCGTGTTTACTGTTTATTGCCCCCT
103 CAAAAAGCTAATGTAATTATATTTGTGCCAATAAAAAACAAGATATGACCTATAGAATACAA
104 AAAAAAAAAAAAAAAAAAAAAAAAAAactcgaagggggcccggtgcagataacttcgtataat
105 gtatgctatacgaagttatgctagcAACCCTTGTGTCATGTCGGCGACCCTACGCCCCCA
106 ACTGAGAGAACTCAAAGGTTACCCAGTTGGGGCACTACTCCCAAAACCGCTTCTGACCT
107 GGGctcgagggcgcgAGGTCCTGCAAAGAAAAGCGGCACCACCCTATCCGTCGACCGGGCC
108 GCCTTCCTTTTGTGTCGCGTCAAGAAAGTGTTCAGAAGAAGCGACAGCAGCGCAAGGAGC
109 GACA

111 **Supplementary Data 2.** The sequence of the knock-in cassette is displayed and coloured as
112 specified to indicate the different segments.