

**Table S4: List of rescue templates**

Gene	crRNA	Description	Sequence
<i>dpy-10</i>	589	Co-CRISPR misense mutation	ssODN from Arribere <i>et al.</i> 2014 (See Suppl. File S1)
<i>gtbp-1</i>	728	Ollas tag insertion at the STOP codon	GGCGGTTCGGGTGGTGCT <u>CCACGAGGAAGGAAATGCGTAGCGGTTCCAAAATGCGGGAC</u> AAAAT---Ollas---tagaagcttccgttccttttccttgtaag
<i>gtbp-1</i>	728	mTagGFP::TEV::3Xflag insertion at the STOP codon	GGCGGTTCGGGTGGTGCTCCACGAGGAGGAATGCGTAGCGGTTCCAAAATGCGGGAC AAAAT---mTagGFP::TEV::3Xflag---tagaagcttccgttccttttccttgtaag
<i>gtbp-1</i>	728	mCherry insertion at cut	GGTTCGGGTGGTGCTCCACGAGGTGGTATGCGC ---mCherry--- AGCGGTTCCAAAATGCGGGACAAAATtagaag
<i>gtbp-1</i>	772	mCherry insertion at cut	CTTCCGCCGTGGCG <u>GGTTCGGGTGGTACGCGTCCACGA</u> ---mCherry--- <u>GCTGGTATGCGCACGCGTTCCAAAATGCGGGAC</u>
<i>gtbp-1</i>	728	eGFP insertion at cut	GGTTCGGGTGGTGCTCCACGAGGTGGTATGCGC ---eGFP--- AGCGGTTCCAAAATGCGGGACAAAATtagaag
<i>fbf-2</i>	689	tagRFP insertion at cut (STOP codon)	CAAATTTCTCTCC <u>GGTAGTCACGGCGATGAT</u> ---tagRFP--- <u>taagg</u> tgaacttctaccataatctcatcc
<i>GFP/eGFP</i>	719 and 720	Replacement by tagRFP	GGAGAAGAAC <u>TTTCACTGGAGTTGTCCAAATT</u> ---tagRFP--- <u>GTCCTCTTGAGTTGTAACAGCTGCTGGGATTAC</u>
<i>glh-1</i>	723	tagRFP insertion at the STOP codon	CCAGTGTACCAACTCAAG <u>TCCCTAAGACGAGGAAGGATGG</u> ---tagRFP--- tagaaaaccgaccatttatgtatgtttcgattttatgc
<i>htp-3</i>	699	TEV::eGFP::myc::3Xflag insertion at the STOP codon	CCAGTCGCCAACATGCCAAGTAGAA <u>AGAGGAAAC</u> ---TEV::eGFP::myc::3Xflag--- <u>ttaacgatttctgg</u> acaatcgatcaattatc
<i>gtbp-1</i>	691 and 728	<i>gtbp-1</i> replacement by GFP::H2B	aaATGCAGATAACCTCGGCCTAACCCAGAATAAG---GFP::H2B--- <u>taac</u> cagggtttccaaatgcgggacaaaattagaagc

Lower case (non-coding), upper case (coding), red (silent mutations in the repair template to prevent Cas9 re-cutting), blue (insertion), bold (PAM sequence), underlined (crRNA sequence).