

**Table S8.** Oligonucleotides used in this study. Restriction sites in the oligonucleotides are underlined, and codon changes for site directed mutagenesis are in bold.

Oligonucleotide	Sequence (5' → 3')	Source or reference
<b><u>Cloning</u></b>		
CCL2-seq fwd	AGGCCTCAGCACTCTCACTC	This study
CCL2-seq rev	GCTCTTCTGGGACTTGAGGA	This study
CCL2-fwd	<u>GGGAATTCCATATGGACT</u> CCCCAGCTGTGAC	This study
CCL2-rev	<u>GGGGGGGGATCC</u> TTAGACCTTCTCGATGACCAG	This study
CCL2-NHis fwd	<u>GGGAATTCCATATGG</u> CCATCATCATCATCATCACA	This study
	GGCGCGACTCCCCAGCTGTGACGCTC	
CCL1 fwd	CAAACCCAACTTACTTCTTCACCC	This study
CCL1 rev	CGAGTTGTGAAAGGTTACGTCCA	This study
CCL1 fwd	<u>CCCC</u> ATGGATACTCAGGCCAAACCCC	This study
CCL1 rev	<u>GGGG</u> AAATTCTCAGACCCCTCTAAAGATCCAG	This study
CCL1-NHis fwd	<u>GGG</u> CATATGGATCATCATCATCATCATCACACTCAG	This study
	GCCAAACCCCCCGCCGG	
dTomato fwd	<u>GGGGGGGGATTAATGG</u> TGAGCAAGGGCGAGG	This study
dTomato rev	<u>GGGGGGGGATCC</u> CTACATATGCTTGACAGCTCGTCATGC	This study
fut1 A	CTAAATTGGCATCCACAAACCT	This study
fut1 B	GCCATTITATTAAACAGTTCTCAT	This study
fut1 C	CCGGAGTAATTAGACCTGC	This study
fut6 A	GAATGCCACCATGCAACAT	This study
fut6 B	GAATTACCCATGATACTAGAT	This study
fut6 C	GCCCCAAATATCAATCTGC	This study
<b><u>Site directed mutagenesis</u></b>		
CCL2 Y57A	TGGATCTTAAGGAG <u>GGCC</u> ACTCGAACCGAAC	This study
CCL2 W78A	CAGCCAGATCGGG <u>GGCG</u> CTGGTAAC	This study
CCL2 L87A	CGTCCC <u>CGTC</u> <u>CCCC</u> CTCCAAACAAAC	This study
CCL2 N90A	GTCGT <u>CC</u> CTCC <u>CC</u> AC <u>GG</u> CCTACGTCTGGACT	This study
CCL2 N91A	GTC <u>CT</u> CC <u>CC</u> CCAA <u>AC</u> AC <u>GG</u> CGTCTGGACTCTG	This study
CCL2 Y92A	CT <u>CC</u> CT <u>CC</u> CA <u>AC</u> A <u>AC</u> AC <u>GG</u> CGTCTGGACTCTGACT	This study
CCL2 V93A	CT <u>CC</u> CT <u>CC</u> CA <u>AC</u> A <u>AC</u> ACTAC <u>GG</u> CTGGACTCTGACTTTGACT	This study
CCL2 W94A	CCT <u>CC</u> CA <u>AC</u> A <u>AC</u> ACTACGT <u>CG</u> <u>CG</u> ACTCTGACTTTGACT	This study
CCL2 K109A	TACAA <u>AC</u> ATT <u>CA</u> AG <u>AT</u> GG <u>CG</u> <u>CG</u> AG <u>GG</u> AC <u>CG</u> TCT <u>GG</u>	This study
CCL2 Y57A	TGGATCTTAAGGAG <u>GGCC</u> ACTCGAACCGAAC	This study
<b><u>qRT-PCR</u></b>		
Tubulin fwd	GTCATGT <u>CCGGT</u> TAC <u>CCAC</u>	[1]
Tubulin rev	GGGAAAGGA <u>ACCAT</u> GT <u>GA</u>	[1]
CCL2 fwd	CTGGTGG <u>ATAC</u> AC <u>ATT</u> CA <u>AG</u> AT <u>GG</u>	This study
CCL2 rev	AGAC <u>CTT</u> CT <u>CG</u> <u>AT</u> GC <u>CC</u> <u>AG</u> C	This study
CCL1 fwd	TGG <u>CGG</u> <u>CT</u> <u>AT</u> AT <u>CA</u> CC <u>AG</u>	This study
CCL1 rev	CAG <u>ACC</u> <u>CT</u> <u>CA</u> <u>AG</u> AT <u>CC</u> <u>AG</u>	This study

## References:

- Wälti MA, Villalba C, Buser RM, Grunler A, Aebi M, et al. (2006) Targeted gene silencing in the model mushroom *Coprinopsis cinerea* (*Coprinus cinereus*) by expression of homologous hairpin RNAs. *Eukaryot Cell* 5: 732-744.