Modified auxin improves the auxin-inducible degradation (AID) system for laid C. elegans embryos

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(A) Chemical structural formula of indole-3-acetic acid (IAA) and acetoxymethyl indole-3-acetic acid (IAA-AM). (B) Frames from time lapse videos of ieSi58 [Pefh-3::degron (AID⁎)::GFP]; ieSi57 [Pefh-3::AtTIR1::mRuby] laid embryos in IAA (1 mM) or IAA-AM (1 mM); elapsed time (hours:minutes) from the beginning of recording is indicated. GFP panels show maximum intensity projection images, and green fluorescent signal is colored with "Fire" look-up-table. Scale bar, 20 μm. (C) Plots of green fluorescent (GF) intensity in timelapse observation. Red lines: IAA-AM treatment of ieSi58; ieSi57 embryos, green lines: IAA-AM treatment of ieSi58 embryos (no TIR1 present), blue lines: IAA treatment of ieSi58; ieSi57 embryos. Mean values of GF intensity in eggshells were obtained every three minutes, and normalized by the intensity at 0 min. (D) Comparison of GF depletion between IAA and IAA-AM treatment of ieSi58; ieSi57 embryos. Mean values of GF intensity within eggshells were obtained after each treatment (two hours), and normalized with that of the beginning, *** indicates p < 0.01, p = 1.6 x 10⁻⁸ (Wilcoxon rank sum test). (E) Representative images of os169 ieSi57 adults, which are IAA or IAA-AM treated during embryos development (two hours).

Description
The targeted protein degradation systems in which a protein accompanying with specific tags can be degraded are developed as an approach of conditional loss of function analyses (Natsume and Kanemaki 2017). The insertion of tags into the gene loci by the CRISPR/Cas9 system allows us to deplete endogenous proteins in stage and cell specific manners. So far, four systems can be used to deplete the tagged proteins in *C. elegans* (Armenti et al. 2014; Zhang et al. 2015; Wang et al. 2017; Wu et al. 2017). The auxin-inducible degradation (AID) system can degrade the tagged protein by administration of the phytohormone auxin. In the AID system, a plant-derived TIR1 F-box protein can form an E3 ubiquitin ligase complex with the endogenous Skp and Cullin proteins, and in the presence of auxin, the complex interacts with a degron tag derived from the IAA17 transcriptional repressor (Nishimura et al. 2009; Yesbolatova et al. 2019). Consequently, the degron-tagged proteins are polyubiquitinated for degradation by the proteasome. With the controlled expression of TIR1 and administration of auxin, the AID system can allow us to perform spatiotemporal protein depletion. Although this system can be a powerful tool for conditional loss of function analyses, it appears to be difficult to degrade proteins in embryos, especially at late embryonic stages, since efficiencies of degradation in laid embryos surrounded by the eggshell is low compared to those in hatched larvae or adults (Zhang et al. 2015).

Here, we report that acetoxymethyl indole-3-acetic acid (IAA-AM), a cell permeable analog of natural auxin IAA (Figure 1A), is more effective in triggering the degradation of tagged proteins in embryos. To test the efficiency of protein degradation, we used the previously reported strain ubiquitously expressing both a 44-amino acid (-aa) degron derived from the *Arabidopsis* IAA17 protein (known as AID*) fused with mRuby (ieSi58 [P$_{eff}$-3:::degron (AID*)::GFP] and ieSi57 [P$_{eff}$-3:::AtTIR1::mRuby]) (Morawska and Ulrich 2013; Zhang et al. 2015). When we put laid embryos into the drop of M9 buffer containing 1 mM IAA-AM (AM form of IAA), we observed rapid decrease of green fluorescence (GF) intensity (Figure 1B, C and D). In contrast to IAA-AM, the embryos treated with IAA showed increased GF intensity due to elevated expression of degron-tagged GFP (Figure 1B, C and D). We also confirmed that IAA-AM did not cause the decreased GF intensity without the AtTIR1 transgene (Figure 1C).

Next, we tried to deplete the endogenous protein in laid embryos. WRM-1 is the *C. elegans* homolog of β-catenin and is essential for asymmetry of most cell divisions (Mizumoto and Sawa 2007). Since WRM-1 is required for asymmetric division of EMS in the 4-cell stage embryo (Rocheleau et al. 1997), conditional knockdown approaches are required for the investigation of its function in later stages. To test whether the AID system with IAA-AM allows us to perform conditional knockdown of *wmr-1* in embryos, we inserted a fusion tag composed of the mini-auxin inducible degron (mAID), a 68-aa degron derived from the *Arabidopsis* IAA17 protein that showed similar degradation efficiency with that of AID* (Li et al. 2019), and mClover at the 3’ end of *wmr-1* endogenous locus by the CRISPR/Cas9 system (Natsume et al. 2016) (Dickinson et al. 2015). We put laid embryos of the strain carrying *wmr-1*:mAID::mClover (os169) and P$_{eff}$-3::AtTIR1::mRuby (ieSi57) into drops of 1 mM IAA-AM or IAA (both with 0.2% DMSO in M9 buffer) for two hours. After washing with M9 buffer two times, we placed embryos on NGM plates without the inducing ligand. At the young adult stage, we scored elongation of gonadal arms guided by the distal tip cells (DTCs) whose production requires *wmr-1* (16/16) or *wmr-1* (15/17 showed normal gonad) (Figure 1E). In addition, all *wmr-1* animals (13/13) shows no gonadal extension. In contrast, treating with 1 mM IAA caused this phenotype only in 2/17 animals (15/17 showed normal gonad) (Figure 1E). In addition, all *wmr-1* embryos treated with 0.2% DMSO (16/16) or ieSi57 ieSi58 embryos treated with 1 mM IAA-AM (19/19) produced animals with normal gonads. These results show that the AM modification of IAA induces efficient protein degradation in laid embryos with the AID system. Since the AM modification is widely used to improve cell membrane permeability (Schultz 2003), IAA-AM is likely to penetrate into the eggshell more easily than IAA. After penetration into cells, endogenous cytoplasmic esterases are known to cleave the AM bond, making the compound again impermeant to cell membrane (Schultz 2003). We note that the chemical cleavage reaction of the AM bond produces formaldehyde (Schultz 2003), which may be toxic to cells at high concentrations. In this study, however, we found the application of IAA-AM at a concentration of 1 mM for two hours allowed normal development including gonad elongation. Our data presented here show IAA-AM will broaden the possible applications of the AID system for conditional loss of function analyses in *C. elegans*.

**Reagents**

**Plasmid construction:** For insertion of degron tag and mClover into 3’ end of *wmr-1* locus by CRISPR/Cas9 system, the fragment of mAID::mClover was amplified with the primers (F: 5’-GGCTCAGGAGCATCGGATCCGACGGCGCCAAGG-3’, R: 5’-AAAGTACAGATTCTCCTTTGTAAGCCTCGTCCATGCA-3’) from pMK290 (Addgene: #72828). The fragment of self-excising cassette (SEC) described in Dickinson et al., 2015 was amplified with the primers (F: 5’-AAAGTACAGATTCTCCTTTGTAAGCCTCGTCCATGCA-3’) from pDD287 (Addgene #70685). These fragments were assembled by InFusion HD (Takara Clontech). Then, 5’ and 3’ *wmr-1* homology arms were amplified from *C. elegans* genome with the following primers:

wmr-1RightL: 5’-AGCGAGGAAGACTTTGTGAAATCTTGTGTTCGGGTGTA-3’;
wmr-1RightR: 5’-CTATGACCAGTTATAACTGTGTTGATCGTGCTTTGG-3’
wrm-1LeftL: 5’-AACGACGGCCAGTCGTTTTGTTGAATGCAAATATGTG-3’
wrm-1LeftR: 5’-GGATCCCGATGCCTCATTAGTTGTCGATGATGCTGC-3’.

Both fragments of homology arms were assembled into the backbone and SEC fragments amplified from the above-described construct with the primers (backbone-F: 5’-CGACTGGCCGTCGTTTTACAAC-3’, backbone-R: 5’-ATAAACATGGTCATAGCTGTTTC-3’ and SEC-F: 5’-GGAGCATCGGGATCCGGTCAG-3’, SEC-R: 5’-CAAGTCTTCCTCGTGAATCAAC-3’) by InFusion HD (Takara Clontech). The vector expressing the guide RNA for wrm-1 was constructed with the primers including the target sequence (F: 5’-[ATGTGAATGAATCTTTGTGC]GTTTTAGAGCTAGAAATAGCAA-3′, R: 5’-AAGATTCACTCATAAACATTTAGTTGGAATTTCA-3‘, brackets indicate the target sequence) by inverse PCR with PU6::unc-119_sgRNA (Addgene #46469).

**Strain construction:** All *C. elegans* strains were cultured by standard methods. Plasmids (50 ng/µl mAID::mClover with wrm-1 homology arms construct, 50 ng/µl wrm-1 guide RNA construct, 50 ng/µl sur-5::GFP co-injection marker and Cas9 expression construct (pDD162, Addgene #47549)) were microinjected into N2. After picking an animal displaying a roller phenotype, self-excising cassette was excised as described (Dickinson et al. 2015). Then, we crossed this strain to the *P*eft-3::AtTIR1::mRuby strain.

**Observation:** In time lapse observation, embryos were mounted on slides with 20 μm diameter polystyrene beads (Polyscience #18329) diluted 1:30 in M9 buffer containing 1 mM IAA or IAA-AM (stocks 500 mM in DMSO)(Bao and Murray 2011). In the endogenous protein degradation experiments, laid embryos were put in drops of M9 buffer containing 0.2% DMSO, 1 mM IAA or IAA-AM for two hours, and transferred into M9 buffer two times before hatching. Washed embryos were allowed to develop on new NGM plates. All imaging was performed with Zeiss LSM700 microscope using a 63x N.A. 1.40 oil-immersion objective. Acquired images were processed by Fiji (Schindelin et al. 2012).

**Strains:**

CA1202: ieSi57 (*P*eft-3::AtTIR1::mRuby) II ; ieSi58 (*P*eft-3::degron (AID*)::GFP) IV, CGC  
HS3280: ieSi58 (*P*eft-3::degron (AID*)::GFP) IV , this study  
HS3176: ieSi57 (*P*eft-3::AtTIR1::mRuby) II; wrm-1(os169 [wrm-1::mAID::mClover]) III, this study  

**IAA and IAA-AM:**  
IAA (indole-3-acetic acid) was purchased from Nacalai tesque (#19119-61). IAA-AM was custom synthesized by Tokyo Chemical Industry Co., Ltd. Both ligands were purchased in DMSO stock solution before storing at -20°C.

**References**


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