

1 **Agrochemical control of gene expression using evolved split RNA polymerase. II**

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15 **Abstract**

16 Agrochemical inducible gene expression system provides cost-effective and orthogonal control of
17 energy and information flow in bacterial cells. However, the previous version of the Mandipropamid
18 inducible gene expression system (Mandi-T7) became constitutively active at room temperature. We
19 moved the split site of the eRNAP from position LYS179 to position ILE109. This new eRNAP showed
20 proximity dependence at 23°C, but not at 37°C. We built a Mandi-T7-v2 system based on the new
21 eRNAP, and it worked in both *Escherichia coli* and *Agrobacterium tumefaciens*. We also induced GFP
22 expression in *Agrobacterium* cells in a semi-*in vivo* system. The modified eRNAP, when combined
23 with the leucine zipper-based dimerization system, behaved as a cold-inducible gene expression
24 system. Our new system provides a means to broaden the application of agrochemicals for both
25 research and agricultural applications. Portions of this text were previously published as part of a
26 preprint (<https://www.biorxiv.org/content/10.1101/2024.04.02.587689v1>)

27 **Introduction**

28 Agrochemicals are key to health promotion and growth management in modern agriculture. The
29 development of biosensors for agrochemicals opens new avenues to control cell behavior through
30 agrochemicals applied by remotely piloted aircraft (Park, 2015, Zimran, 2022, Park, 2024). One of the
31 possible applications is to use agrochemicals to control gene expression of bacteria associated with

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34 plants or animals. Mandipropamid is an oomycete fungicide (Blum, 2010). We recently reported the
35 development of Mandipropamid inducible gene expression system (Mandi-T7) using a protein
36 proximity detection platform based on evolved split T7 RNA polymerase (eRNAP) (Yuan, 2022). Our
37 Mandi-T7 system is based on two systems developed by other groups. The first system is the
38 repurposed plant Absciscic acid receptor proteins (PYR1^{MANDI} and ABI), which will sense agrochemical
39 Mandipropamid and will be brought into proximity upon Mandipropamid binding via molecular
40 ratchet mechanism (Park, 2015; Zimran, 2022; Steiner, 2023; Park, 2024). This in turn activates the
41 eRNAP and leads to expression of genes driven by the T7 promoter (Figure 1A, Yuan, 2022). The
42 second system we borrowed is the eRNAP system. Wild type T7 RNAP split at site 179 is proximity-
43 independent and was used to construct transcriptional logic gates (Shis, 2013; Segall-Shapiro, 2014).
44 The simultaneous assembly of the split T7 RNAP was abolished through directed evolution, and the
45 eRNAP became proximity-dependent (Pu, 2017). Several small molecule inducible expression
46 systems were developed based on this eRNAP platform, such as Absciscic acid and Rapamycin (Pu,
47 2018), Mandipropamid (Yuan, 2022), and Rapalog (Martin, 2023). The molecular variety of this
48 eRNAP biosensor platform has been further expanded by fusion with the variable domains of
49 antibodies (Komatsu, 2023) or with cell-pole organizing proteins to achieve asymmetric gene
50 expression (Lin, 2021).

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51 The Mandi-T7 system was initially tested at 37°C. The failure of Mandi-T7 to work at room
52 temperature hampers engineering initiatives in other bacteria. Here, we report the alleviation of this
53 issue by adopting a new split site of T7 eRNAP. The new system Mandi-T7-v2 works at 23°C in *E.coli*
54 *in vitro* and in *Agrobacterium in vitro* and semi-*in vivo*. The modified eRNAP is also compatible with
55 a leucine zipper peptide-based dimerization system and may be used with other chemical-induced
56 dimerization systems as well.

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57 **Material and Methods**

58 **Plasmid construction**

61 The eRNAP_N (derived from 1-109 aa of T7 RNAP), the T7 RNAP_C (110-883 aa of T7 RNAP), and ZA/ZB
62 fragments were synthesized and cloned into the plasmid pJM186 (Yuan, 2022) by GenScript
63 Biotechnology (Nanjing, China). The pVS1 origin from pCambia1301 (GenBank: AF234297.1, 2488
64 bp -6266 bp), the Mandi-T7-v2 driver cassette, and the T7p::mcherry / toehold switch -sfGFP effector
65 cassette were assembled into the pGM1190 plasmid (Addgene #69994) backbone to enable single-
66 plasmid expression in *Agrobacterium*. Detailed information on these genetic parts and plasmids is
67 listed in Table S1 and S2.

68 **Mandipropamid responsive assay for evolved split T7 RNA polymerase.**

69 The response to Mandipropamid was assayed as previously reported. Briefly, the strain Top10
70 (TIANGEN biotech., Beijing, China) was transformed with the driver and reporter plasmids. Single
71 colonies were inoculated into SOC medium supplemented with Ampicillin (100 mg/mL) and
72 Spectinomycin (50 mg/mL), and the mixture was allowed to grow at 37 °C. The overnight culture
73 was transferred to a fresh medium with antibiotics at a 1:400 ratio and incubated for 3 h at 37 °C.
74 Then, induction by Mandipropamid was tested at both 23°C and 37 °C. Mandipropamid (sc-235565,
75 Santa Cruz) was added as the inducer, and DMSO as the solvent control. After incubating for 3 h at
76 37 °C or 6 h at 23°C, 100 µL of each sample was transferred to a 96-well plate. The fluorescence
77 signal (GFP, Ex: 488 nm, Em: 510 nm; mcherry, Ex: 587 nm, Em: 630 nm) and OD₆₀₀ of the culture
78 were then measured using a ThermoFisher Varioskan LUX plate Reader. For *Agrobacterium*
79 *tumefaciens*, the strain LBA4404 (Weidibio, Shanghai, China) was transformed with the plasmid pJM-
80 Mandi-T7-Ag and grown at 28°C. Single colonies were inoculated with SOC medium supplemented
81 with Apramycin (50 mg/mL) and grown at 28°C. The overnight culture was transferred to a fresh
82 medium with antibiotics at a 1:100 ratio and incubated for 3 h at 28 °C. Induction was performed
83 overnight at 23°C after adding Mandipropamid or DMSO (vehicle control). For both *E. coli* and
84 *Agrobacterium tumefaciens*, the fluorescence intensity was divided by or normalized with the OD₆₀₀
85 value. Then, normalized fluorescence intensity values from induced samples (Mandipropamid) and
86 control (DMSO) were compared to yield fold change. Raw data was provided as supplemental file

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97 “Raw_data_figure1B_1C_1D_1F_1H_1J.xlsx”. The reporter gene mcherry was used for Figure 1B,
98 Figure 1C, and Figure 1D. The reporter gene GFP was used for Figure 1F, Figure 1H, and Figure 1J.

99 **Cold responsive assay of split eRNAP fused with leucine zipper peptides.**

100 The *E. coli* strain Top10 was used. Single colonies were inoculated with SOC medium supplemented
101 with Ampicillin (100mg/mL) and Spectinomycin (50 mg/mL) and grown at 37 °C. The overnight
102 culture was transferred to a fresh medium with antibiotics at a 1:400 ratio and incubated for 3 h at
103 37 °C. Following this, half the culture was incubated at 23°C, the other half stayed at 37 °C. After 3 h
104 at 37 °C or overnight at 23°C, 100 µL of each sample was transferred to a 96-well plate. The
105 fluorescence signal (GFP, Ex: 488 nm, Em: 510 nm) and OD₆₀₀ of the culture were measured using a
106 ThermoFisher Varioskan LUX plate Reader. Raw data was provided as supplemental file
107 “Raw_data_figure1B_1C_1D_1F_1H_1J.xlsx”.

108 **Time course fluorescence measurement**

109 *E. coli* Top10 strain containing Mandi-T7_v2 and reporter plasmids was generated as mentioned
110 above. Single colonies were used to inoculate SOC medium supplemented with Ampicillin (100
111 mg/mL) and Spectinomycin (50 mg/mL) and grown at 37 °C overnight. The Overnight culture was
112 transferred to a fresh medium with antibiotics at a 1:100 ratio. Afterwards, either Mandipropamid
113 (50 mM, sc-235565, Santa Cruz) or DMSO (solvent control) was added at a 1:1000 ratio. The culture
114 was incubated at 23°C for 40 h. Samples were harvested after 1 hour, 4 h, 7 h, 15 h, 23 h, 30 h and
115 40 h. The fluorescence (mcherry: Ex: 587 nm, Em: 630 nm) and OD₆₀₀ of 100 µL of each sample were
116 measured using a ThermoFisher Varioskan LUX plate Reader. Raw data was provided as
117 supplemental file “Raw_data_figure1B_1C_1D_1F_1H_1J.xlsx”.

118 **Mandipropamid response assay in a semi-*in vivo* system**

119 *Agrobacterium tumefaciens* strain LBA4404 containing toehold switch-GFP driven by Mandi-T7-v2
120 was generated as mentioned above. Single colonies were inoculated LB medium with antibiotics at

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131 28 °C. Overnight culture was transferred to a fresh medium with antibiotics at a 1:100 ratio. When
132 OD₆₀₀ reached roughly 0.4, aliquots of 300 µL culture were added into the hollow septate stems of
133 water spinach (*Ipomoea aquatica*), which were cut into cylinder shaped and inserted into a 96-well
134 deep well plate as previously reported (Yuan, 2022). Mandipropamid (200 µM final) or DMSO
135 (solvent control) was added. The plate was shaken in an orbital shaker (1,000 rpm) at 23°C for 24 h.
136 A total of 200 µL of each sample was added to a 96-well plate. Both fluorescence (GFP: Ex: 488 nm,
137 Em: 510 nm;) and OD₆₀₀ were measured by a ThermoFisher Varioskan LUX plate Reader. The
138 fluorescence intensity was normalized with the OD₆₀₀ value. Values from induced samples
139 (Mandipropamid) and control (DMSO) were compared to yield fold change. Raw data was provided
140 as supplemental file “Raw_data_figure1B_1C_1D_1F_1H_1J.xlsx”.

141 **Statistical Analysis**

142 Datasets were analyzed within the Office Excel software. Two-tailed t-tests were used for pairwise
143 comparisons. P values were indicated in the figure legend.

144

145 **Results**

146 To apply Mandi-T7 to bacteria living at room temperature, we characterized Mandi-T7 at 23°C. The
147 results showed that the fluorescence signal of the report gene could be detected regardless of the
148 presence of the inducer (Figure 1B and Supplemental Figure 1), indicating that the eRNAP relapsed
149 into self-assembly at 23°C. Restoration eRNAP’s proximity dependency at 23°C will enable Mandi-T7
150 to work at 23°C. We speculated that a new split site in the N terminal region of T7 RNA polymerase,
151 not previously reported by the bisection mapping efforts (Segall-Shapiro, 2014), might prevent self-
152 assembly at 23°C. We scrutinized the structure of the T7 RNAP N terminal region. We selected
153 ILE109 as our candidate because ILE109 is exposed at the surface, and splitting at ILE109 / LYS110 will

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165 break the connection between the Arginine loop, which is essential for binding upstream AT-rich
166 region of the T7 promoter, with the rest of the T7 RNA polymerase (Supplemental Figure 2).

167 To assess the potential improvement in induction at 23°C, we modified the driver module of the
168 Mandi-T7 system (Figure 1A). The split site was changed to ILE109/LYS110 while retaining the point
169 mutations in the N-terminal region of the eRNAP (F21L, L32S, E35G, R57C, E63K, K98R, Q104K,
170 Q107K) as leverage for flexibility. Induction for 6 h at 23°C yielded a modest fold change over the
171 control (Figure 1B). Unexpectedly, induction by Mandipropamid at 37°C was abolished by adopting
172 the new split site (Figure 1B). Encouraged by these findings, we named this modified eRNAP as
173 eRNAP2, and our new Mandipropamid inducible expression system, Mandi-T7-v2. We proceeded to
174 characterize the induction with different concentrations of Mandipropamid. We observed maximum
175 induction by 200 μM Mandipropamid at 23°C (Figure 1C). We further evaluated the kinetic
176 characteristics of the Mandi-T7-v2 at 23°C over 30 h. We observed an induction signal after 1 hour
177 and a continuous increase over 30 h (Figure 1D and Supplemental Figure 1). However, the
178 fluorescence signal of the non-induction samples accumulated simultaneously, resulting in a
179 marginal increase in fold induction after 23 h (Figure 1D).

180 To increase the dynamic range without further engineering the Mandi-T7-v2 system, we tried to
181 incorporate the toehold switch into the reporter module (Figure 1E), which has been shown to
182 improve the performance of T7-based inducible expression systems through a coherent type 1 feed-
183 forward loop (Hwang, 2021; Greco, 2021). Indeed, after 24 h induction, incorporation of the toehold
184 switch could increase the dynamic range by two-fold (Figure 1F).

185 To test whether Mandi-T7-v2 can be applied to other bacteria, we tried Mandi-T7-v2 in
186 *Agrobacterium tumefaciens*, a plant pathogen and medium of T-DNA transformation. To generate a
187 single-plasmid-based induction system, we assembled the Mandi-T7-v2 driver module, the reporter
188 module and the pVS1 replication origin together. We tested the Mandipropamid induction in

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200 *Agrobacterium* at 23°C. The result showed a 15-fold induction (Figure 1B and Supplemental Figure
201 1), suggesting that the Mandi-T7-v2 system also works in other bacteria.

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202 To test whether induction at 23°C but not at 37°C is an inherent property of the new eRNAP2 or
203 depends on the fusion to the ABI-PYR1^{MANDI} protein pair. We replaced the ABI-PYR1^{MANDI} pair with
204 leucine zipper peptides ZA and ZB (Figure 1G), one of the model dimerization systems that will lead
205 to spontaneous dimerization and has been used for developing the evolved RNAP system before (Pu,
206 2017; Pu, 2018). We tested the ZA/ZB-eRNAP2 system at both 23°C and 37°C. The results showed
207 50-fold higher expression at 23°C over 37°C (Figure 1H and Supplemental Figure 1), reminiscent of a
208 cold inducible expression system. This result indicates the restored proximity dependency of the
209 eRNAP2 at 23°C does not require specific fusion partners.

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210 Encouraged by the results above, we evaluated the ability of Mandi-T7-v2 to function in plants at
211 23°C. We used the hollow septate stems of water spinach (*Ipomoea aquatica*) as the containers to
212 induce bacterial gene expression as previously reported (Yuan, 2022). We used toehold switch-
213 regulated GFP as the report gene. *Agrobacterium* culture containing Mandi-T7-v2 and the reporter
214 gene was incubated in the water spinach stems at 23°C for 24 h. The results showed 15-fold
215 induction (Figure 1J) over vehicle control. This result suggests Mandi-T7-v2 might be useful in other
216 complex settings such as vector-borne bacterial plant pathogens.

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217 Discussion

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218 Fortunately, We have found a new split site of eRNAP, which enables chemical inducible activity
219 control at 23°C. Induction around 23°C will be an advantage for controlling gene expression in
220 bacteria living around room temperature, especially bacteria associated with model organisms raised
221 around 23°C, such as *Arabidopsis*, *Drosophila*, and *C. elegans*. As a proof-of-principle, we have
222 shown that reported gene expression could be induced in *Agrobacterium* inside plant tissue. Our
223 new system also widens the temperature range of the existing induction system. For example,

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234 proteins produced by Mandi-T7 system in *E.coli* may not be properly folded at 37°C. Misfolding may
235 be mitigated at 23°C in our new system. This new eRNAP2 system can also be adapted to other
236 proximity-dependent systems. Our cold response assay of the ZA/ZB-eRNAP2 system exemplified
237 this possibility. For dimerization pairs that work at 23°C but not at 37°C, eRNAP2 would be preferred
238 as their interaction sensor.

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239 When we tested the performance of the Mandi-T7 system at room temperature, we found
240 constitutive activity without inducibility, indicating the disrupted spontaneous assembly of eRNAP at
241 37°C was restored at room temperature. The temperature sensitivity of the eRNAP platform may be
242 explained by its origin in the directed evolution system running at 37°C (Pu, 2017).

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243 Several split sites of T7 RNA polymerase have been identified by the bisection mapping, like around
244 67, 301, 563, and 763 (Segall-Shapiro, 2014). Except for split site 563, which has also been used for
245 engineering the thermo-repressible split T7 RNAP (Chee, 2022), and the blue light-inducible systems
246 (Han, 2017 and Dionisi, 2022), these split sites have not been developed into general or specific
247 biosensor platforms. Our work suggests it is worth exploring new split sites of T7 RNA polymerase
248 for specific applications.

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249 We incorporated a toehold switch into our simple gene circuit to improve dynamic range. This
250 provided an easy way to improve the performance of Mandi-T7-v2 system on protein-encoding
251 genes. However, the leakage issue persists for RNA devices driven by Mandi-T7-v2. Further directed-
252 evolution efforts may be required to reduce the background activity of the Mandi-T7-v2 system.

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253 The ABI-PYR1 pair has been recently expanded to be sensors of other agrochemicals (Zimran, 2022;
254 Park, 2024). Our work will readily help develop new inducible gene expression systems of these
255 agrochemicals. Both the plant growth -promoting (PGP) traits and synthetic bacteria-to-plant
256 communication pathways have been engineered in the root-associated bacteria (Haskett, 2021; Boo,
257 2024). Key PGP traits include nitrogen fixation, nutrient mobilization, antibiotics production, etc.

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(Haskett, 2021). Agrochemical control of PGP traits or bacteria-to-plant communication will add temporal modulation to these devices.

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333 **Figure 1.** Engineering the Mandi-T7-v2 system."(A) Comparison of the Mandi-T7 and Mandi-T7-v2
334 systems. The length of different parts of fusion proteins were drawn in scale. Bent arrow: T7
335 promoter; large T-shape: terminator. (B) fold change of reporter gene expression after induction at
336 23°C or 37°C. Asterisk: $p < 0.05$. (C) Dose response analysis of Mandi-T7-v2 in *E.coli* after induction
337 at 23°C for 6 h. Asterisk: the lowest concentration with significant differences ($p < 0.005$) compared
338 to the uninduced controls. (D) kinetic analysis of Mandi-T7-v2 system at 23°C. Asterisk: the earliest
339 time point with significant differences ($p < 0.05$) compared to the mock. (E) schematic of toehold
340 switch incorporated into Mandi-T7-v2 induction system. (F) fold change of report gene expression.
341 GFP: GFP expression without toehold switch. Toehold switch-GFP: A toehold switch regulator was
342 added for GFP. Asterisk: $p < 0.001$. (G) schematic of leucine zipper peptides ZA/ZB mediated
343 dimerization of eRNAP2. (H) temperature dependent expression of the reporter gene. Asterisk: $p <$
344 0.001. (J) fold change of reporter gene expression in a semi-*in vivo* system. Asterisk: $p < 0.001$.

345 **Supplemental figure 1.** cell pellet of *E.coli* and *Agrobacterium* culture post induction.
346 Samples were placed on the sill of a window with normal day light. +: Mandipropamid (50 μ M final);
347 -: DMSO.

348 **Supplemental figure 2.** the ILE109 split site of T7 RNA polymerase.
349 The N terminal region (1-109) of the T7 RNA polymerase was highlighted in dark purple. The
350 Arginine loop responsible for binding the AT-rich region was encircled by a red square. The cartoon
351 was generated from the structure of T7 RNA polymerase – T7 promoter complex (PDB ID: ICEZ).

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