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Supplemental information

**Biosensor and chemo-enzymatic one-pot cascade
applications to detect and transform PET-derived
terephthalic acid in living cells**

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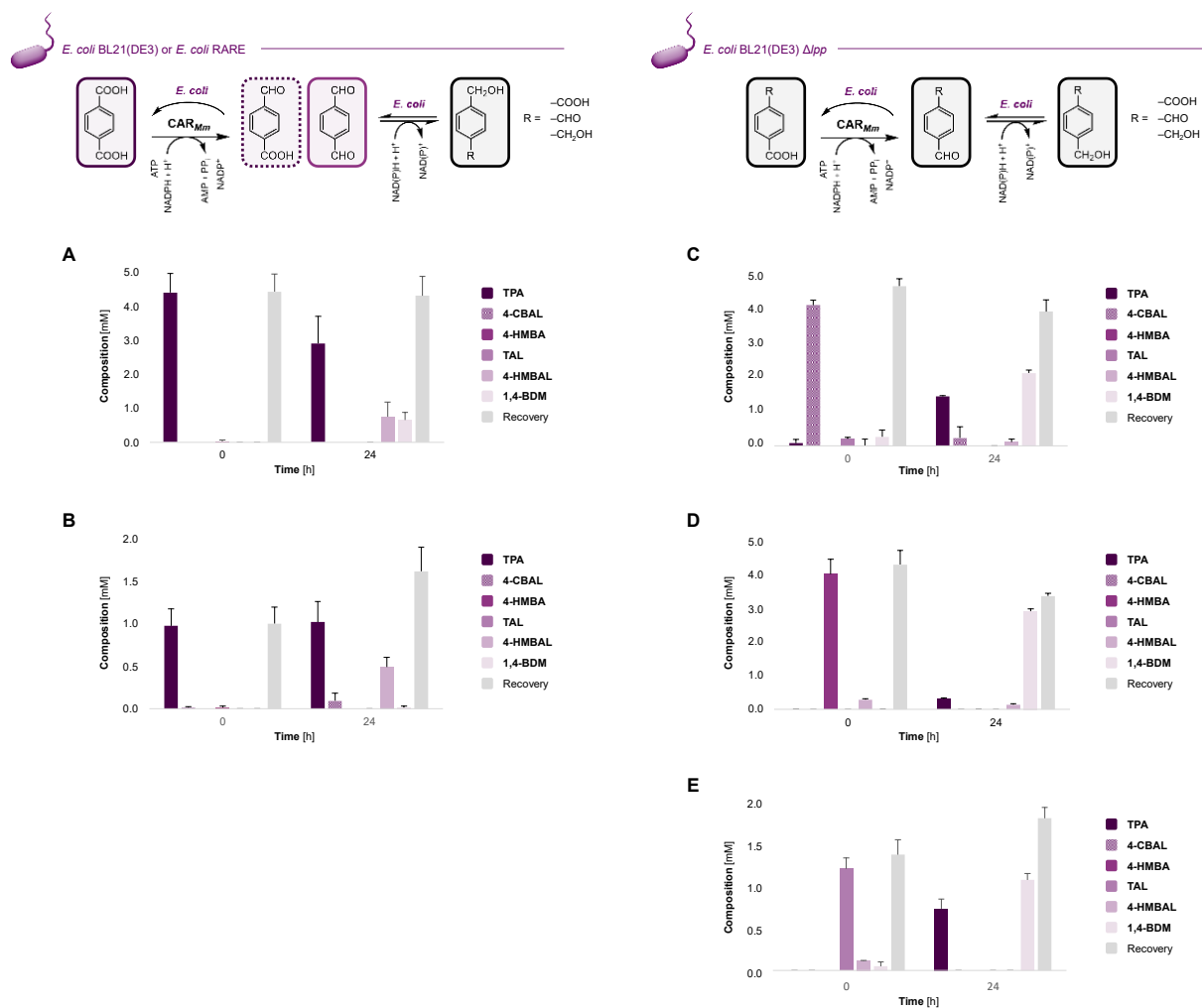


Figure S1. CAR-catalyzed reductions of TPA and derivatives *in vivo*, related to Figure 2A. (A) Bioreduction of TPA by CAR_{Mm} in *E. coli* BL21(DE3) yields a mixture of the over-reduced 4-HMBAL and 1,4-BDM, besides unreacted substrate. **(B)** Bioreduction of TPA by CAR_{Mm} in *E. coli* RARE mainly yields 4-HMBAL, indicating reduced aromatic aldehyde reducing activity compared to *E. coli* BL21(DE3), 4-CBAL, and unreacted TPA. **(C)** CAR_{Mm} reduces the carboxylate group in 4-CBAL to the corresponding TAL. Endogenous enzymes not only reduce aldehydes to the corresponding primary alcohols like 1,4-BDM but also oxidize them to the carboxylates as indicated by the detection of TPA after 24 h. This is in accordance with previous findings (Bayer et al., 2017). **(D)** CAR_{Mm} reduces the carboxylate group in 4-HMBA to the corresponding 4-HMBAL. Endogenous enzymes reduce the aldehydes to the corresponding primary alcohols like 1,4-BDM; TPA could be detected in traces after 24 h. **(E)** The highly reactive TAL is both oxidized and reduced by endogenous enzymes, yielding TPA and 1,4-BDM, respectively. On top, arrows indicate the activities of (host) enzymes; PPT_{Ni} necessary for posttranslational modification of CAR_{Mm} is omitted for clarity. Experiments were performed in RCs of *E. coli* ($OD_{600} \approx 10.0$) co-expressing enzymes from $pACYCDuet-1/car_{Mm}:ppt_{Ni}$ (Bayer et al., 2021) in the presence of 2–5 mM substrates and 5% (v/v) DMSO as organic co-solvent. Sampling: 0 h (after the addition of substrate and mixing) and 24 h. Recoveries were reduced due to low solubilities and/or the volatility of compounds. GC yields are presented as mean values + standard deviation (SD) [mM] of biological replicates ($n \geq 3$).

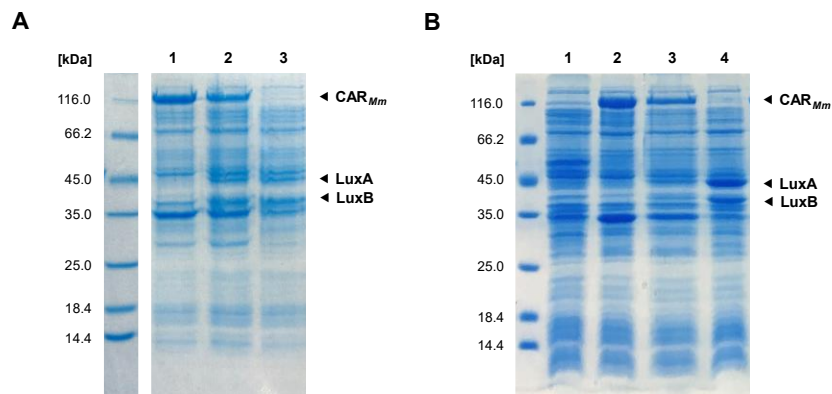


Figure S2. SDS-PAGE analysis of whole-cell samples, related to Figure 2A, Figure S1, and STAR Methods. (A) Expression of (1) CAR_{Mm}/PPT_{Ni} from pACYCDuet-1/*car_{Mm}:ppt_{Ni}* [CAR_{Mm}: 129 kDa], (2) CAR_{Mm}/PPT_{Ni} and LuxAB from pLuxAB [LuxA: 43 kDa, LuxB: 37 kDa], and (3) LuxAB in *E. coli* BL21(DE3). (B) Whole-cell samples of (1) untransformed *E. coli* BL21(DE3) Δ lpp or expressing (2) CAR_{Mm}/PPT_{Ni}, (3) CAR_{Mm}/PPT_{Ni} and LuxAB, and (4) LuxAB; PPT_{Ni} [23 kDa] was not detectable due to low expression levels in corresponding samples. Proteins were produced from pACYCDuet-1/*car_{Mm}:ppt_{Ni}* and pLuxAB (Bayer et al., 2021); the detailed protocol is given in the main text. Sample loading normalized to OD₆₀₀ = 7.0; SDS-PAGE and gel staining performed as described in the main text. Irrelevant lanes were cropped in (A), the brightness of both pictures was increased by 20%; (◄) indicate protein bands of interest.

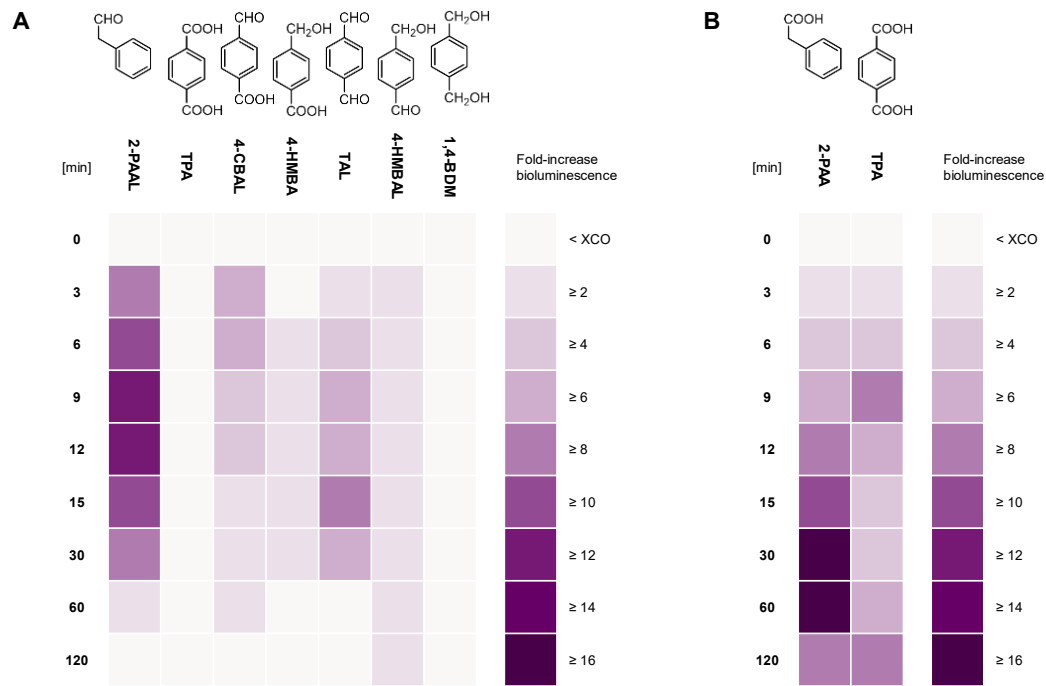


Figure S3. LuxAB-based HT detection of aldehydes in *E. coli* RARE, related to Figure 2B–C. (A) Direct detection of TPA-derived aldehydes (0.1 mM) by increasing bioluminescence over time in RCs of *E. coli* RARE expressing LuxAB from pLuxAB; 2-PAAL was used as the positive control. Whereas TPA and 1,4-BDM did not yield bioluminescence, the addition of 4-HMBA yielded background luminescence at 1 mM final concentration. **(B)** *In situ* production of aldehydes from 2-PAA and TPA (1 mM) in RCs of *E. coli* RARE co-expressing LuxAB and CAR_{Mm}/PPT_{Ni}. Experiments were performed in the presence of 1% (*v/v*) DMSO under HT assay conditions as described previously (Bayer et al., 2021) and in the main text; data presented as mean fold-increase bioluminescence obtained from biological replicates (*n* = 3).

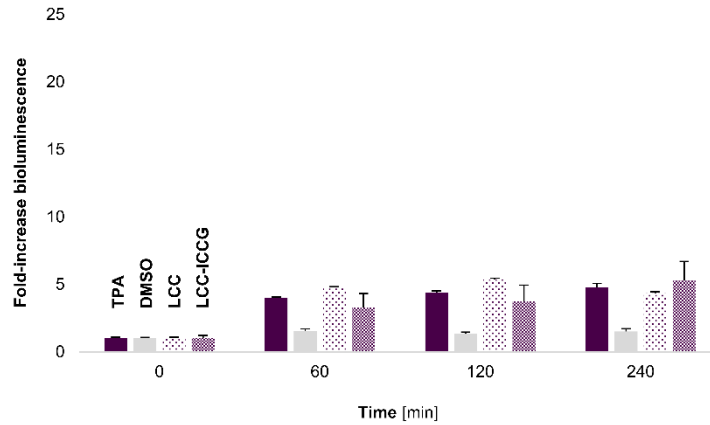


Figure S4. PET hydrolysis samples analyzed under HT conditions in *E. coli* BL21(DE3) Δlpp , related to Figure 3. The enzyme-coupled biosensor system yielded bioluminescence in the presence of 1 mM TPA (positive control) and hydrolysates obtained by the enzymatic degradation of Gf-PET films by wildtype LCC and the LCC-ICCG variant. The bioluminescence did not increase in the presence of 1% (v/v) DMSO over monitoring time (negative control). While the bioluminescence plateaued around 4-fold above background in *E. coli* BL21(DE3) Δlpp after 1 h incubation time, it increased in RCs of *E. coli* RARE proportionally to the amounts of TPA present in PET hydrolysis samples (see **Figure 3** in the main text). Experiments were performed in RCs of *E. coli* BL21(DE3) Δlpp under HT assay conditions as described previously (Bayer et al., 2021) and in the main text; data presented as mean values of the fold-increase in bioluminescence + SD of biological replicates (n = 3).

Table S1. List of DNA oligonucleotides, related to **STAR Methods**

Primer	Sequence (5' – 3')
<i>lpp</i> -up_F	gagtcgacctgcagaagcttGTAAAGAACTGGCTCTGCAGAG
<i>lpp</i> -up_R	acaggtactaCCCTCTAGATTGAGTTAATCTCC
<i>lpp</i> -down_F	atctagagggTAGTACCTGTGAAGTGAAAAATG
<i>lpp</i> -down_R	gagctgcacatgaactcgagATGAATGCACCGGATATTAAGC
pTarget_F	ctcgagttcatgtgcagctc
pTarget_R	aagcttctgcaggtcgactc
Δ <i>lpp</i> -gRNA_F	AGTAGAACCCgtttagagctagaaatagcaagtt
Δ <i>lpp</i> -gRNA_R	CTGCTGGCAGactagtattatacctaggactgagc

Table S2. HPLC yields of TPA in PET hydrolysates, related to **Figure 3** and **STAR Methods**

PET hydrolase	TPA yields [mM]
PES-H1	56.0 \pm 0.1
LCC	47.8 \pm 3.1
LCC-ICCG	111.1 \pm 15.3

Yields are given as mean values \pm SDs of independent PET hydrolysis experiments and subsequent HPLC measurement for LCC (n = 3) and LCC-ICCG (n = 2) and as mean value \pm SD of a technical replicate (n = 2) for PES-H1.