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# Cross-linking of a polyketide synthase domain leads to a recyclable biocatalyst for chiral oxygen heterocycle synthesis†

 Lisa Wagner,  Theresa Roß, Tim Hollmann and Frank Hahn \*

The potential of polyketide synthase (PKS) domains for chemoenzymatic synthesis can often not be tapped due to their low stability and activity *in vitro*. In this proof-of-principle study, the immobilisation of the heterocycle-forming PKS domain AmbDH3 as a cross-linked enzyme aggregate (CLEA) is described. The AmbDH3-CLEA showed good activity recovery, stability and recyclability. Repetitive reactions on the semi-preparative scale were performed with high conversion and isolated yield. Similar to that observed for the free enzyme, the aggregate retained substrate tolerance and the ability for kinetic resolution. This first example of a successful enzymatic PKS domain immobilisation demonstrates that cross-linking can in principle be applied to this type of enzyme to increase its applicability for chemoenzymatic synthesis.

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Type I-polyketide synthases (type I-PKS) are multi-domain enzymes that biosynthesise the backbones of reduced polyketide natural products. Their biochemistry is related to that of fatty acid synthases, but type I-PKS have a much broader range of products due to their more flexible assembly and enzymatic mechanism.<sup>1,2</sup> Their catalytic cycle is based on the precisely tuned interplay of the individual modules and domains, which is in turn controlled by protein–protein and protein–substrate interactions.<sup>3–5</sup>

Due to their origin from this specific environment, isolated PKS domains often exhibit rather low turnover frequency and low stability in *in vitro* experiments.<sup>5–7</sup> On the other hand, PKS domains frequently show relaxed substrate specificity and, particularly in the case of domains that catalyse synthetically attractive reactions, an application in chemoenzymatic synthesis is thus conceivable.<sup>6,8–13</sup> Various recent reports have highlighted the existing potential of PKS enzymes for this purpose.<sup>14–19</sup>

Immobilisation is an effective way to increase the practical value of enzymes by providing them with higher stability and recyclability.<sup>20,21</sup> Besides the covalent and non-covalent attachment of enzymes to solid phases, the formation of cross-linked enzyme aggregates (CLEAs) or cross-linked enzyme crystals are established approaches with specific advantages.<sup>22</sup> The production of CLEAs is operationally simple and combines purification of the protein with the formation of a heterogenous catalyst. CLEAs basing on well-characterised primary metabolic

biocatalysts, such as hydroxynitrile lyases and carbohydrate-processing enzymes, have frequently been reported as well as some basing on secondary metabolic enzymes.<sup>22–25</sup>

Despite the potential of this technique, successful immobilisation or cross-linking of type I-PKS has not been reported.<sup>26</sup> This might be due to various reasons such as their problematic handling, their naturally low activity *in vitro* or their complex enzymatic cycle that involves major spatial rearrangements. Apart from improving the chemoenzymatic performance of individual domains, their immobilisation would represent a sensible first step towards immobilisation of larger PKS constituents.

It should be noted that immobilised type III-PKS have been employed for the synthesis of plant natural product libraries.<sup>27–29</sup> The analogy between type I and type III-PKS is however limited, because being single-domain synthases, the latter differ fundamentally from type I-PKS, both structurally and mechanistically. We regarded CLEA formation to be well-suited for PKS immobilisation as this method is mild, easily adaptable and minimally interfering with the pre-organised superstructure of proteins.

We have recently described the bifunctional dehydratase-cyclase domain AmbDH3 from the ambruticin type I-PKS as a new biocatalyst for the synthesis of saturated oxygen heterocycles *via* intramolecular oxa-Michael addition (Scheme 1).<sup>12,15,30</sup> Besides its broad substrate-tolerance and high stereoselectivity, this enzyme is characterised by convenient scalability of its conversions and the ability to perform kinetic resolution of chiral tetrahydropyrans (THPs).

We thus chose AmbDH3 for a proof-of-principle study on the cross-linking of PKS domains that could ideally lead to an improved heterocyclisation catalyst.

Department of Chemistry, Faculty of Biology, Chemistry and Earth Sciences, University of Bayreuth, Universitätsstraße 30, 95447 Bayreuth, Germany. E-mail: frank.hahn@uni-bayreuth.de

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**Scheme 1** AmbDH3 catalyses intramolecular oxa-Michael addition leading to chiral oxygen heterocycles. ACP: acyl carrier protein; SNAC: *N*-acetylcysteamine.

The formation of CLEAs is achieved by cross-linking of free amino groups on the protein surface by bifunctional linkers, like glutaraldehyde (GA).<sup>22</sup> A sufficient number of accessible residues is required, otherwise these need to be installed by artificial surface amination. Luckily, inspection of the AmbDH3 crystal structure (PDB-ID: 5O16) revealed the presence of various evenly distributed lysines and arginines on the enzyme surface, suggesting its suitability for CLEA formation (Fig. 1a).<sup>30</sup> The active centre of AmbDH3 contains a catalytic dyad of His-Arg within an otherwise highly hydrophobic cavity, making disturbance of this region by the cross-linker rather unlikely.<sup>30</sup> The near-surface localisation of the active site should also allow good accessibility for substrates with expectable limitations caused by the slowed diffusion through the cross-linking network.

AmbDH3-CLEAs were prepared by precipitation of the enzyme with ammonium sulphate and cross-linking with GA in the same vessel.<sup>24</sup> Two buffers and GA concentrations ranging from 0.5 to 5% were investigated (Fig. 1b and S1–S7†). The



**Fig. 1** (a) Crystal structure of AmbDH3 with highlighted lysine (light green) and arginine (blue) surface residues and the active centre (turquoise). (b) Conversion of **1** by different preparations of AmbDH3-CLEA (blue bars) as well as the lysate and the supernatant from CLEA preparation (0.5% GA, NaP buffer; grey bars). Reaction conditions: 332 nmol (2 mM) of **1**, CLEA/lysate from 4 mg *ambDH3*-expressing cells, 16 h, 30 °C.

**Table 1** Activity recovery, immobilisation yield and efficiency as well as enzyme loading in AmbDH3-CLEA compared to the reaction of the *ambDH3*-expression lysate with **1**<sup>a</sup>

Parameter	Value
Activity recovery	81%
Immobilisation yield	94%
Immobilisation efficiency	86%
AmbDH3 bound in the CLEA	<59%

<sup>a</sup> Determined from the enzymatic activity of the individual fractions from CLEA preparation. Reaction conditions as described in the caption of Fig. 2.

resulting aggregates were assessed regarding their ability to convert **1** into THP **2** in 16 h endpoint assays.<sup>15</sup> All experiments were performed using an identical quantity of *ambDH3*-expressing cells for the preparation of the cell-free lysate or the CLEA, respectively. Cross-linking with 0.5% GA in sodium phosphate (NaP) buffer resulted in a CLEA that gave similar conversion as the expression lysate and a markedly higher value than one resulting from cross-linking in HEPES buffer. A 16 h incubation with the supernatant from the precipitation step served as a control experiment and showed only very low conversion of **1** into **2**, indicating that most of the remaining enzymatic activity is bound in the CLEA. A GA concentration above 2% in NaP buffer led to a marked decrease of conversion. As preparation with 0.5, 1 or 2% GA gave no apparent difference in physical stability of the resulting CLEAs, cross-linking with 0.5% GA in NaP buffer for 2 h was used to form the AmbDH3-CLEAs in all subsequent experiments.

To more precisely describe the outcome of the immobilisation process, we determined the activity recovery in the CLEA. The enzymatic activity of the individual fractions from CLEA preparation was determined and set into relation as previously described (Table 1, Fig. S8, Tables S1 and S2†).<sup>24</sup> 81% of the lysate enzymatic activity were recovered. As obvious from the immobilisation yield and the immobilisation efficiency, both, incomplete binding of enzyme activity and a slightly lower activity of the cross-linked compared to the free enzyme, contributed to this incomplete recovery by a relevant degree. A reduced enzyme activity after cross-linking is well-described and can be attributed to disturbance of protein conformation, blockage of the active centres or diffusion restrictions.<sup>20,22</sup> Aggregates with a higher activity could be obtained in the future by optimising the preparation procedure for AmbDH3-CLEA, for example by screening other cross-linking agents or fine-tuning of the molar ratio of cross-linker to enzyme.

Attempts to determine the amount of AmbDH3 in the CLEA did not give absolutely precise values, but allowed the estimation that  $\leq 60\%$  of the AmbDH3 in the lysate were bound in AmbDH3-CLEA (Fig. S9†). Together with the observed activity recovery this suggests that relevant parts of the free AmbDH3 in the lysate were inactive and that active AmbDH3 is enriched by the precipitation step during CLEA formation.

Determination of the Michaelis-Menten parameters was complicated by practical issues like the small reaction scale and



Table 2 Stability of AmbDH3-CLEA towards long time storage and freeze-thawing

Entry	Storage <sup>a</sup> (d)	Temperature <sup>b</sup> (°C)	Conversion (%)
1	0	20	81
2	7	20	80
3	7	4	78
4	1	-20	72
5	1	-80	74

<sup>a</sup> Storage period of the CLEA before reaction under the conditions described in the caption of Fig. 2. <sup>b</sup> Temperature during storage period.

a limited solubility of the substrate **1** in the reaction buffer, leading to large errors. The  $K_m$  and  $k_{cat}/K_m$  values of the AmbDH3-CLEA ( $4.4 \pm 2.8$  mM and  $5.67 \pm 4.87$  s<sup>-1</sup> mM<sup>-1</sup>) were in the same range as those of the free enzyme ( $8.2 \pm 3.1$  mM and  $28.2 \pm 14.8$  s<sup>-1</sup> mM<sup>-1</sup>) and suggested slower catalysis as well as similar specificity of the CLEA compared to the free enzyme (Fig. S10–S17 and Table S3†). With respect to the structure of AmbDH3 this suggests that the diffusion into the active centre and its flexibility is not severely affected by the cross-linking process. A time course experiment showed a steady increase of conversion to the final value of 81% during an overnight incubation (Fig. S18†). Of these, 58% were already achieved after the initial 5 h reaction, which represents a common time frame for a biocatalytic reaction.

AmbDH3-CLEA was subjected to various storage conditions and the effect on the conversion determined (Table 2, Fig. S20 and S21†). While the CLEA stored for 7 days at 20 °C or 4 °C, respectively, transformed **1** into **2** by a similar degree as the freshly prepared one, freeze-thawing to -20 or -80 °C had a slightly negative influence on the conversion. The AmbDH3-CLEA was also applied in a recycling experiment in which recovered aggregate was repetitively inserted into further reaction-wash cycles. The CLEA showed activity in all cycles with a conversion gradually decreasing from 86% to 55% (Fig. 2 and S22–S31†). It could not be clarified under these experimental setting if the decline in conversion was caused by diminishing

### AmbDH3-CLEA recycling

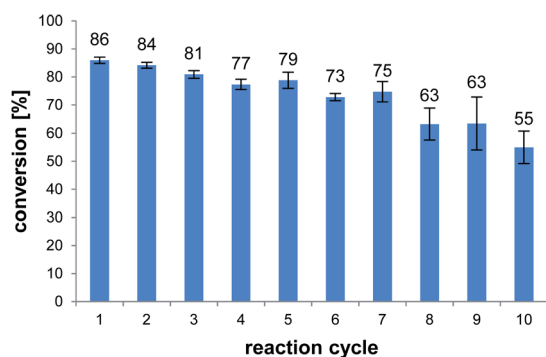


Fig. 2 Conversion of **1** into **2** in ten subsequent reaction cycles using recovered AmbDH3-CLEA. The CLEA was washed with buffer after each cycle. Reaction conditions: 332 nmol (2 mM) of **1**,  $1.9 \times 10^{-3}$  u AmbDH3-CLEA, 30 °C, 16 h.

### semipreparative-scale reactions

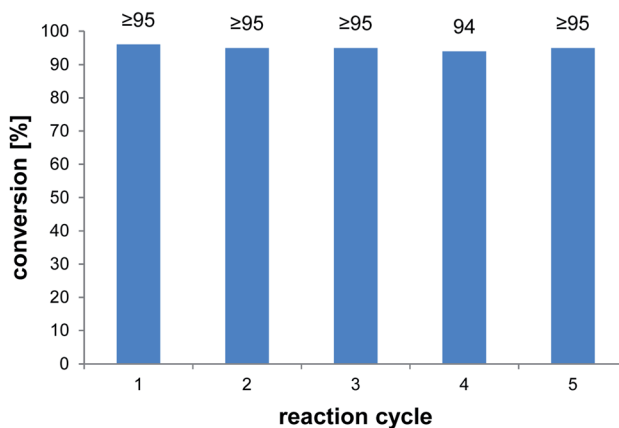


Fig. 3 Conversions of semi-preparative scale reactions of AmbDH3-CLEA with **1**. The conversion was determined by <sup>1</sup>H NMR spectroscopy. Reaction conditions: 33 μmol (4 mM) of **1**, 0.19 u AmbDH3-CLEA, 30 °C, 16 h.

activity of the CLEA itself or by loss of minor proportions of the CLEA in the workup steps of the individual cycles. These results suggest a certain stability of AmbDH3-CLEA and allow the conclusion that the aggregate retains a relevant part of its activity upon long-term storage, freeze-thawing and repetitive exposure to the conditions of a standard conversion experiment.

To clarify the performance under conditions more relevant for chemoenzymatic synthesis, we scaled up the reaction between AmbDH3-CLEA and **1** to 33 μmol (10 mg) starting material and doubled the substrate concentration to 4 mM (Fig. 3 and S32–S36†). Virtually quantitative conversion into homochiral **2** was observed in five repetitive reaction cycles, suggesting that the declining conversions over the analytical scale incubations were an artefact from the handling of small CLEA amounts. The average isolated yield of the five pooled reactions was 94%, which is significantly above the ~80%, which were observed for purified AmbDH3 or the expression lysate on the same reaction scale.<sup>15</sup> The co-precipitation of

Table 3 Substrate specificity of AmbDH3-CLEA<sup>a</sup>

Entry	Substrate	Product	Conversion/yield (%)
1			>95/94 <sup>b</sup>
2			80/41
3			35/20

<sup>a</sup> Scale: ~30 μmol (4 mM) substrate, 0.19 u AmbDH3-CLEA, 30 °C, 16 h.

<sup>b</sup> Average from five consecutive reaction cycles.



enzyme, **1** and **2** posed a major problem in the homogenous reaction with the free enzyme on reaction scales of 33  $\mu\text{mol}$  **1** or above. A 2 h incubation with proteinase prior to extractive work-up of the homogenous reaction was necessary to dissolve this co-precipitate and obtain satisfactory isolated yields. The stability of AmbDH3-CLEA and its recoverability effectively eliminate the need for this measure in the heterogenous semi-preparative scale reaction and should thus also enable simplified workup on the preparative reaction scale.

Finally, we investigated the substrate tolerance of the AmbDH3-CLEA. We incubated the aggregate with the substrate surrogates **3** and *rac*-**5** and determined conversion and isolated yield (Table 3 and Fig. S37–S41†). AmbDH3-CLEA converted **3** stereospecifically and with 80% conversion into phenyl-THP **4**. In the reaction with *rac*-**5**, AmbDH3-CLEA converted only one stereoisomer, (*R*)-**5**, into (*R*)-**6** and thus enabled one-step resolution of a chiral THP with three stereocentres. The conversions of **3** and (*R*)-**5** by AmbDH3-CLEA is lower than in the analogous reactions with the free enzyme,<sup>15</sup> which is consistent with the finding that the CLEA does not show full activity recovery.

In summary, we performed a proof-of-principle study for PKS domain cross-linking on the example of the THP-forming AmbDH3. AmbDH3-CLEA showed a good activity recovery, stability towards a couple of treatments and could be recycled at least ten times. Similar to the free enzyme, AmbDH3-CLEA exhibited substrate tolerance and kinetic resolution ability in semi-preparative scale reactions as well as higher isolated yield and simpler work-up in reactions with the standard substrate surrogate **1**. Ultimately, a heterocyclisation catalyst with improved properties was obtained.

Our work shows that type I PKS domains can be converted into an active, immobilised form with synthetically useful properties by cross-linking. It lays the foundation for applying this method to other PKS domains and larger PKS components, such as entire modules.

## Conflicts of interest

There are no conflicts to declare.

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## Notes and references

- 1 C. Hertweck, *Angew. Chem., Int. Ed.*, 2009, **48**, 4688–4716.
- 2 A. T. Keatinge-Clay, *Chem. Rev.*, 2017, **117**, 5334–5366.
- 3 T. Robbins, Y.-C. Liu, D. E. Cane and C. Khosla, *Curr. Opin. Struct. Biol.*, 2016, **41**, 10–18.
- 4 K. J. Weissman and R. Müller, *ChemBioChem*, 2008, **9**, 826–848.

- 5 M. Klaus and M. Grininger, *Nat. Prod. Rep.*, 2018, **35**, 1070–1081.
- 6 W. D. Fiers, G. J. Dodge, D. H. Sherman, J. L. Smith and C. C. Aldrich, *J. Am. Chem. Soc.*, 2016, **138**, 16024–16036.
- 7 W. D. Fiers, G. J. Dodge, Y. Li, J. L. Smith, R. A. Fecik and C. C. Aldrich, *Chem. Sci.*, 2015, **6**, 5027–5033.
- 8 G. Berkhan, C. Merten, C. Holec and F. Hahn, *Angew. Chem., Int. Ed.*, 2016, **55**, 13589–13592.
- 9 D. H. Kwan and P. F. Leadlay, *ACS Chem. Biol.*, 2010, **5**, 829–838.
- 10 S. K. Piasecki, C. A. Taylor, J. F. Detelich, J. Liu, J. Zheng, A. Komsoukianians, D. R. Siegel and A. T. Keatinge-Clay, *Chem. Biol.*, 2011, **18**, 1331–1340.
- 11 J. Wunderlich, T. Roß, M. Schröder and F. Hahn, *Org. Lett.*, 2020, **22**, 4955–4959.
- 12 G. Berkhan and F. Hahn, *Angew. Chem., Int. Ed.*, 2014, **53**, 14240–14244.
- 13 P. Pöplau, S. Frank, B. I. Morinaka and J. Piel, *Angew. Chem., Int. Ed.*, 2013, **52**, 13215–13218.
- 14 S. Sundaram, H. J. Kim, R. Bauer, T. Thongkongkaew, D. Heine and C. Hertweck, *Angew. Chem., Int. Ed.*, 2018, **57**, 11223–11227.
- 15 T. Hollmann, G. Berkhan, L. Wagner, K. H. Sung, S. Kolb, H. Geise and F. Hahn, *ACS Catal.*, 2020, **10**, 4973–4982.
- 16 C. C. Aldrich, L. Venkatraman, D. H. Sherman and R. A. Fecik, *J. Am. Chem. Soc.*, 2005, **127**, 8910–8911.
- 17 A. Pinto, M. Wang, M. Horsman and C. N. Boddy, *Org. Lett.*, 2012, **14**, 2278–2281.
- 18 E. S. Sattely, M. A. Fischbach and C. T. Walsh, *Nat. Prod. Rep.*, 2008, **25**, 757–793.
- 19 S. Friedrich and F. Hahn, *Tetrahedron*, 2015, **71**, 1473–1508.
- 20 C. Garcia-Galan, Á. Berenguer-Murcia, R. Fernandez-Lafuente and R. C. Rodrigues, *Adv. Synth. Catal.*, 2011, **353**, 2885–2904.
- 21 R. A. Sheldon, *Adv. Synth. Catal.*, 2007, **349**, 1289–1307.
- 22 R. A. Sheldon, *Catalysts*, 2019, **9**, 261.
- 23 M. R. Bennett, M. L. Thompson, S. A. Shepherd, M. S. Dunstan, A. J. Herbert, D. R. M. Smith, V. A. Cronin, B. R. K. Menon, C. Levy and J. Micklefield, *Angew. Chem., Int. Ed.*, 2018, **57**, 10600–10604.
- 24 M. Frese and N. Sewald, *Angew. Chem., Int. Ed.*, 2015, **54**, 298–301.
- 25 M. Ismail, M. Frese, T. Patschkowski, V. Ortseifen, K. Niehaus and N. Sewald, *Adv. Synth. Catal.*, 2019, **361**, 2475–2486.
- 26 L. Betancor and H. R. Luckarift, *Biotechnol. Genet. Eng. Rev.*, 2010, **27**, 95–114.
- 27 B. Ku, J. Cha, A. Srinivasan, S. J. Kwon, J.-C. Jeong, D. H. Sherman and J. S. Dordick, *Biotechnol. Prog.*, 2006, **22**, 1102–1107.
- 28 S. J. Kwon, M.-Y. Lee, B. Ku, D. H. Sherman and J. S. Dordick, *ACS Chem. Biol.*, 2007, **2**, 419–425.
- 29 M. I. Kim, S. J. Kwon and J. S. Dordick, *Org. Lett.*, 2009, **11**, 3806–3809.
- 30 K. H. Sung, G. Berkhan, T. Hollmann, L. Wagner, W. Blankenfeldt and F. Hahn, *Angew. Chem., Int. Ed.*, 2018, **57**, 343–347.

