

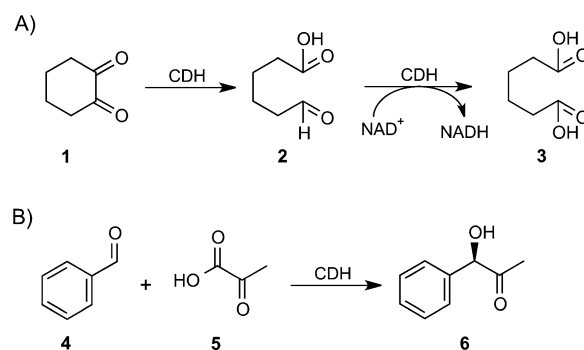
Catalytic Scope of the Thiamine-Dependent Multifunctional Enzyme Cyclohexane-1,2-dione Hydrolase

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The thiamine diphosphate (ThDP)-dependent enzyme cyclohexane-1,2-dione hydrolase (CDH) was expressed in *Escherichia coli* and purified by affinity chromatography (Ni-NTA). Recombinant CDH showed the same C–C bond-cleavage and C–C bond-formation activities as the native enzyme. Furthermore, we have shown that CDH catalyzes the asymmetric cross-benzoin reaction of aromatic aldehydes and (decarboxylated) pyruvate (up to quantitative conversion, 92–99% *ee*). CDH accepts also hydroxybenzaldehydes and nitrobenzaldehydes; these previously have not (or only in rare cases) been known as substrates of other ThDP-dependent enzymes. On a semipreparative scale, sterically demanding 4-(*tert*-butyl)benzaldehyde and 2-naphthaldehyde were transformed into the corresponding 2-hydroxy ketone products in high yields. Additionally, certain benzaldehydes with electron withdrawing substituents were identified as potential inhibitors of the ligase activity of CDH.

Thiamine diphosphate (ThDP)-dependent enzymes are involved in a wide range of metabolic pathways and catalyze a broad variety of reactions. Among these are oxidative and non-oxidative decarboxylation and asymmetric C–C, C–O, C–S, and C–N bond formation, as well as C–C bond cleavage.^[1] Prominent representatives include the enzymes transketolase, pyruvate dehydrogenase, pyruvate decarboxylase, acetolactate synthase, desoxyxylulose 5-phosphate synthase, benzaldehyde lyase, and benzoylformate decarboxylase (BFD).^[1b]

ThDP-dependent cyclohexane-1,2-dione hydrolase (CDH, EC 3.7.1.11) is the key enzyme of an anaerobic degradation pathway of alicyclic alcohols. It catalyzes C–C bond cleavage of cyclohexane-1,2-dione (**1**) to produce 6-oxohexanoic acid (**2**) as the primary product, presumably^[2] followed by oxidation of the latter to adipic acid (**3**, Scheme 1A).^[3–6] This degradation pathway of a 1,2-diketone was discovered for the denitrifying



Scheme 1. ThDP-dependent cyclohexane-1,2-dione hydrolase (CDH) catalyzes A) C–C bond-cleavage and B) asymmetric C–C bond-formation.

bacterium *Azoarcus* sp. strain 22Lin^[7] when cultivated on cyclohexane-1,2-diol as the sole carbon source and electron donor, and nitrate as electron acceptor. Additionally, CDH is able to catalyze nonphysiological asymmetric C–C bond formation. Initially, the cross-benzoin reaction of benzaldehyde (**4**) and pyruvate (**5**) (after decarboxylation) to result in the *R*-configured 1-hydroxy-1-phenylpropan-2-one (PAC, **6**, 98% *ee*) was performed on an analytical scale (Scheme 1B).^[6]

Until recently, CDH has been purified from its native source, *Azoarcus* sp. strain 22Lin, by a multistep chromatographic protocol. Together with the cofactors ThDP, FAD, and Mg^{II}, the enzyme monomer has a theoretical molecular mass of 64.5 kDa.^[3] Whereas SDS-PAGE shows a single protein band for the monomer and size exclusion chromatography indicates that CDH is a homodimer in solution, the crystal structure shows a homotetramer with one noncovalently bound FAD and one ThDP (bound to the protein with an Mg^{II} ion that is coordinated to its diphosphate moiety) per monomer.^[3–6]

The oligonucleotide and amino acid sequences of CDH do not show pronounced similarities to related enzymes, despite having typical cofactor-binding domains.^[8] Interestingly, **1** is also a substrate of ThDP-dependent YerE from *Yersinia pseudotuberculosis*;^[9] however, YerE does not display α -ketolase activity (observed with CDH for this substrate), but instead catalyzes the addition of activated acetaldehyde to result in a tertiary alcohol.^[9] We anticipated that CDH could possess further distinctive features concerning the range of catalytic possibilities. Herein, we describe the detailed characterization of CDH-catalyzed asymmetric transformations, facilitated by heterologous production of the enzyme. The results demonstrate that CDH complements other well-known ThDP-dependent enzymes with respect to substrate range and reaction scope.

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For optimal production and isolation of CDH, a synthetic gene was designed with codon optimization for expression with a C-terminal His₆ tag from pET21a in *Escherichia coli* (see the Supporting Information). *E. coli* BL21(DE3) cells were transformed with the plasmid, grown in lysogeny broth (LB), and gene expression was induced by isopropyl- β -D-thiogalactopyranoside (IPTG). Recombinant CDH was purified by affinity chromatography (Ni-NTA). The purity of the protein was confirmed by SDS-PAGE (see the Supporting Information), and the presence of the cofactor FAD was confirmed by UV/vis spectroscopy. Approximately 15 mg of pure CDH was obtained from 1 L of culture. Although riboflavin supplement in heterologous growth cultures has been reported to optimize the expression of some FAD-dependent enzymes,^[10] in the case of CDH we did not observe any effect of riboflavin (up to 50 mg L⁻¹) on the final yield of holoenzyme.

To determine the catalytic activity of recombinant CDH, both cleavage of cyclohexane-1,2-dione (**1**, Scheme 1A) and the cross-benzoin reaction of benzaldehyde (**4**) and pyruvate (**5**, Scheme 1B) were examined. When **1** was incubated with CDH in the absence of NAD⁺, the cleavage reaction stopped at 6-oxohexanoic acid (**2**). Thus, the cleavage activity of CDH could be determined unequivocally by proton NMR spectroscopy: acidification of the aqueous enzymatic assay and extraction with CDCl₃ gave **2**, which displayed a well-defined signal for its aldehyde proton ($\delta_{\text{H-6}} = 9.78$ ppm, triplet, $J_{6,5} = 1.5$ Hz); furthermore, **1** seemed to exist exclusively as its 2,3-enol form ($\delta_{\text{H-3}} = 6.15$ ppm, triplet, $J_{3,4} = 4.6$ Hz). The formation of the PAC product (**6**) was monitored by GC/MS. Preliminary experiments suggested that a 10 mM:25 mM ratio of **4** to **5** results in the highest conversion after 24 h. In the first few hours, formation of **6** showed first-order kinetics. In the linear region (0–4.7 h), the specific activity of CDH was 9.5 mU mg⁻¹ purified protein (see the Supporting Information). Conversion of **4** into (*R*)-PAC ((*R*)-**6**) was 82% after 24 h, as determined by GC/MS; *ee* was 99%, as determined by chiral-phase HPLC.

Thus, we confirmed that recombinant CDH shows the same C–C bond-cleavage and C–C bond-formation activity as reported for CDH purified from its native source, *Azoarcus* sp. strain 22Lin. Furthermore, it was shown that 20–30% (v/v) DMSO had no negative effects on the conversion.^[11] This allows the use of

hydrophobic aromatic aldehydes in screening experiments on the carboligase activity of CDH.

The scope and limitations of the C–C bond-formation capability of CDH were determined by testing 24 monosubstituted benzaldehydes **7a,b,c**–**14a,b,c** as putative acceptors and pyruvate (**5**) as donor in the formation of the corresponding PAC derivatives **15a,b,c**–**22a,b,c** (Table 1). Overall, in terms of conversion and *ee* of the PAC products, benzaldehydes with one electron-rich substituent (OH, OMe, Me) in the *ortho*, *meta*, or *para* position (substrates **11**–**13**) gave slightly better results than **7**–**10** and **14** (electron-withdrawing groups; F, Cl, Br, I, NO₂). Benzaldehydes substituted in the *para* or *meta* positions showed better conversion than the corresponding *ortho*-substituted derivatives.

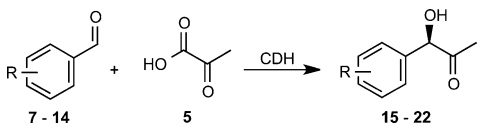
Notably, CDH accepted hydroxy- and nitrobenzaldehydes **11** and **14**, which are not (or only in rare cases)^[12] known as substrates of other ThDP-dependent enzymes. There was no conversion with *o*-nitrobenzaldehyde (**14a**), which was thus the only tested aromatic aldehyde not to be accepted by CDH. Unexpectedly, CDH showed particularly low conversion of 2-fluorobenzaldehyde (**7a**; see below).

All 18 PAC products with determined *ee* values (Table 1) were assigned as *R*-configured, as judged by the appearance of a negative band centered at 270–290 nm in circular dichroism spectra.^[13,14] All 18 products were highly enantioenriched, with *ee* values in the range of 92–99%.

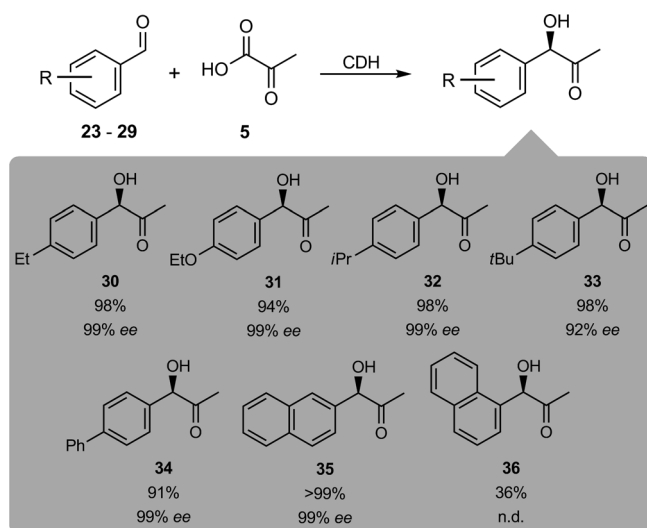
As *para*-substituted and electron-rich benzaldehydes were accepted best in terms of conversion, we applied sterically more demanding aromatic aldehydes **23**–**29** in the cross-benzoin reaction with pyruvate (Scheme 2).

In general, these substrates were almost completely converted, and products **30**–**35** were obtained virtually enantiopure. The reaction of 4-(*tert*-butyl)benzaldehyde (**26**→**33**, 98% conversion) is noteworthy. Although the 2-hydroxy ketone **33** was obtained with a relatively lower *ee* (92%), it should be emphasized that compounds bearing *tert*-butyl groups are notoriously poor substrates in enzymatic transformations.^[15] This is because of the high hydrophobicity of *tert*-butylated substrates (thus low solubility in aqueous media) and the steric hindrance imposed by the *tert*-butyl group. In our example, DMSO (20% v/v) as the solubility promoter undoubtedly alleviated the solubility challenge.

Table 1. CDH-catalyzed conversion (%) of **7**–**14** and enantiomeric excess (% *ee* in brackets) for the formation of PAC derivatives **15**–**22**.^[a]

								
R	F	Cl	Br	I	OH	OMe	Me	NO ₂
substrate	7	8	9	10	11	12	13	14
product	15	16	17	18	19	20	21	22
<i>ortho</i> (a)	5 (n.d.)	6 (n.d.)	24 (99)	45 (96)	19 (n.d.)	37 (99)	62 (99)	0 (–)
<i>meta</i> (b)	47 (98)	34 (98)	49 (96)	51 (94)	82 (99)	96 (92)	91 (98)	20 (n.d.)
<i>para</i> (c)	68 (96)	82 (96)	69 (95)	81 (96)	30 (98)	85 (98)	97 (96)	5 (n.d.)

[a] Conditions: pyruvate (25 mM), aromatic aldehyde (10 mM), CDH (1 mg mL⁻¹), buffer A (50 mM MES, 1 mM MgSO₄, 0.5 mM ThDP, pH 6.5), DMSO (20–30%, v/v), 30 °C, 48 h; % conversion was determined by GC/MS. Enantiomeric excess was determined by chiral-phase HPLC; n.d. = not determined.



Scheme 2. Screening sterically demanding aromatic aldehydes as acceptors with pyruvate (5, donor) in the CDH-catalyzed cross-benzoin reaction. Conditions as in Table 1.

Moreover, CDH used naphthaldehydes efficiently as acceptor substrates, although the sterically hindered 1-naphthaldehyde (29→36) showed only 36% conversion (>99% conversion of the less hindered 2-naphthaldehyde (28→35)). Previously, 2-naphthaldehyde (28) has been applied as a donor substrate in BFD-catalyzed cross-benzoin condensation: in the presence of solubilizing agents, only 34% conversion was obtained after 72 h.^[16] In addition, 1-naphthaldehyde (29) was not accepted at all by BFD, thus further highlighting the significance of our results with CDH. The transformations of 4-(*tert*-butyl)benzaldehyde (26→33) and 2-naphthaldehyde (28→35) were also performed on a semipreparative scale. The 2-hydroxy ketone products 33 and 35 were obtained in 90% (28 mg) and 87% (26 mg) isolated yields, respectively.

In contrast to the broad acceptor substrate spectrum, the donor substrate range of the CDH-catalyzed cross-benzoin reaction with benzaldehyde (4) was, in effect, limited to pyruvate (82% conversion). Aliphatic α -oxo acids larger than 2-oxobutanoic acid (1.3% conversion) were not accepted at all, and neither were 2-oxosuccinate, 2-oxoglutarate, or hydroxypyruvate (the last substrate was tested with 4-ethylbenzaldehyde as acceptor). In the absence of an aromatic acceptor aldehyde, highly enantioenriched (*S*)-acetoin was formed by homocoupling of pyruvate.^[17]

Analysis of the data in Table 1 with respect to the conversion of *ortho*-substituted halobenzaldehydes reveals a trend that is contrary to what was expected. On the basis of the steric requirements and the electrophilicity of the halo substituents, 2-fluorobenzaldehyde (7a) should be accepted best as a substrate, and 2-iodobenzaldehyde (10a) should be accepted least. (This trend is regularly observed with other ThDP-dependent enzymes).^[9,18] However, CDH catalysis displayed the opposite outcome (Table 1). Accordingly, we hypothesized that 2-fluorobenzaldehyde (7a) might act as an inhibitor of the carbonylase activity of CDH. To test this, we set up a competition

experiment with 7a (a potential inhibitor), 4-ethylbenzaldehyde (23, one of the best acceptors), and pyruvate (5, donor).

In the absence of 7a, conversion of 23 to 1-(4-ethylphenyl)-1-hydroxypropan-2-one (30) was 95% after 24 h (Table 2,

Table 2. Competition experiments with 2-fluorobenzaldehyde (7a), 4-ethylbenzaldehyde (23), and pyruvate (5).^[a]

	7a	23	5	15a	30
	Quantity [mm]			Product [%]	
	7a	23	5	15a	30
1	–	10	25	–	95
2 ^[b]	–	10	25	–	98
3	10	10	25	2	1
4	5	10	25	6	1
5	1	10	25	61	34
6	0.1	10	25	n.d. ^[c]	82

[a] Conditions: pyruvate (25 mM), benzaldehydes 7a and 23 (see Table 2), CDH ($c = 1 \text{ mg mL}^{-1}$), buffer A (see Table 1), DMSO (20–30% v/v), 30 °C, 24 h; % conversion (each value is an average from two independent enzymatic assays) was determined by GC/MS. [b] After 48 h. [c] Not detected.

entry 1) and 98% after 48 h (entry 2). In the presence of equimolar amounts of 7a and 23 the conversion was just 1% (entry 3); interestingly, 1-(2-fluorophenyl)-1-hydroxypropan-2-one (15a) was detected in only trace amounts. Lowering the 7a:23 ratio revealed that 7a has a strong negative impact on the formation of 1-(4-ethylphenyl)-1-hydroxypropan-2-one (30): 34% conversion of 23→30 when present as low as 10% relative to 23 (entry 5).

We then turned our attention to other aromatic aldehydes possessing (highly) electron-withdrawing substituents. A potential inhibitory effect on the formation of 30 was tested in independent competition experiments under conditions analogous to those listed in Table 2, entry 4. As anticipated, 2-chlorobenzaldehyde (8a), as well as 2,6-difluoro-, 2,4,5-trifluoro-, and pentafluorobenzaldehyde, showed the same effect as 7a: each of these benzaldehydes yielded $\leq 6\%$ conversion into the corresponding PAC products when incubated with pyruvate (5), and inhibited the formation of 30 in the same manner as 7a (<2% conversion for 23→30). Detailed kinetic studies are required to obtain a mechanistic explanation of these results and to confirm the role of these electron-deficient benzaldehydes as potential inhibitors of the ligase activity of CDH.

In summary, we have described the heterologous expression in *E. coli* and purification of recombinant His-tagged cyclohexane-1,2-dione hydrolase (CDH). In addition to its physiological C–C bond-cleavage activity, CDH catalyzes the asymmetric cross-benzoin reaction of a broad variety of aromatic aldehydes and pyruvate (up to quantitative conversion, 92–99% ee). In the case of the sterically demanding 4-(*tert*-butyl)benzaldehyde (26) and 2-naphthaldehyde (28), the respective 2-hydroxy ketone products (33 and 35) were obtained in high yield. Notably, CDH accepts several benzaldehydes, such as hy-

droxy- and nitrobenzaldehydes; these are not (or only in rare cases) known as substrates of other ThDP-dependent enzymes. An inhibitory effect of benzaldehydes with electron-withdrawing substituents on the carboligase activity was observed.

Hence, CDH expands the range of well-known ThDP-dependent enzymes suitable as catalysts in asymmetric carboligation reactions. The straightforward production and stability of this enzyme, as well as its unique behavior as a powerful multifunctional catalyst, make it an interesting starting point for synthetic applications and mechanistic studies.

Experimental Section

Cloning: The synthetic, codon-optimized, *cdh* gene from *Azoarcus* sp. 22Lin (GeneArt, Life Technologies) was cloned into a pET21a vector (Novagen), between NdeI and XhoI restriction sites. Omission of the gene's stop codon allowed expression with the plasmid-encoded C-terminal His₆ tag.

Expression and purification of CDH: For the production of His-tagged CDH, *E. coli* BL21(DE3) was transformed with pET21a carrying the *cdh* gene. A preculture was grown in LB (7 mL) with ampicillin (100 µg mL⁻¹) at 37 °C for 24 h with shaking (140 rpm). From this, an aliquot was incubated (25 °C, 130 rpm) in fresh LB (500 mL) with ampicillin (100 µg mL⁻¹). Upon reaching OD₆₀₀ = 0.6–0.7 (ca. 5.5 h), IPTG (0.02 mM) was added to induce gene expression. After 17–24 h at 25 °C (130 rpm), cells were harvested by centrifugation (5200g, 40 min, 4 °C). From the 500 mL culture, 2.5–3.0 g of cells were obtained. The cell pellet was resuspended in buffer A (15 mL; MES (2-(*N*-morpholino)ethanesulfonic acid, 50 mM, pH 6.5), MgSO₄ (1 mM), ThDP (0.5 mM)) and lysed by a French press. Cell debris was removed by centrifugation (6000g, 45 min, 4 °C), and the lysate (20 mL) was incubated with Ni-NTA resin (2 mL) at 0 °C for 1 h. After washing with buffer A containing imidazole (20 then 50 mM), recombinant CDH was eluted in buffer A containing imidazole (300 mM). The combined elution fractions were desalted by gel-permeation chromatography in buffer A in a Sephadex column (GE Healthcare). The purity of the protein was confirmed by SDS-PAGE (see the Supporting Information). Approximately 15 mg purified protein was obtained from 1 L cell culture.

CDH-catalyzed asymmetric benzoin reaction of benzaldehydes and pyruvate (analytical scale): Enzymatic reactions on an analytical scale were performed in buffer A (1.5 mL) containing DMSO (20–30% v/v). The aromatic aldehyde (final concentration 10 mM) was applied as a DMSO solution; sodium pyruvate (25 mM) and the enzyme (1 mg mL⁻¹) were added. All reactions were conducted at 30 °C and 300 rpm in a Thermomixer (Eppendorf). After 48 h, the enantiomeric excess was determined by chiral-phase HPLC after extraction of an aliquot (150 µL) of the enzyme assay with ethyl acetate (1 × 200 µL). Conversion was determined by GC/MS after extraction of another aliquot (150 µL) with ethyl acetate (1 × 150 µL). For analytical details of the PAC products, see the Supporting Information.

CDH-catalyzed synthesis of 33 and 35 (semipreparative scale): A solution of 4-(*tert*-butyl)benzaldehyde (**26**, 25.1 µL, 24.3 mg, 0.150 mmol, final concentration in the reaction assay 10 mM) in DMSO (3.75 mL, final concentration 25%, v/v) or 2-naphthaldehyde (**28**, 23.4 mg, 0.150 mmol, final concentration 10 mM) in DMSO (4.50 mL, final concentration 30%, v/v) was added to a solution of sodium pyruvate (41.6 mg, 0.378 mmol, final concentration 25 mM)

in buffer A (final total volume 15 mL) in a 15 mL Falcon tube. CDH (1 mg mL⁻¹) was added, and the mixture was incubated at 30 °C with shaking (400 rpm) for 24 h (for **28**) or 48 h (for **26**) until GC/MS indicated 99% conversion. The reaction mixture was extracted with MTBE (3 × 15 mL). The combined organic layers were dried over MgSO₄, and the volatiles were removed under reduced pressure. The crude product was isolated with an Isolera preparative flash purification system (Biotage AB, Uppsala, Sweden).

(*R*)-1-(4-*tert*-Butylphenyl)-1-hydroxypropan-2-one (**33**) was obtained in 90% yield (28 mg). (*R*)-1-Hydroxy-1-(naphthalen-2-yl)propan-2-one (**35**) was obtained in 87% yield (26 mg).

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Keywords: asymmetric catalysis • C–C bond-formation • carboligation • enzyme catalysis • ThDP

- [1] a) R. Kluger, K. Tittmann, *Chem. Rev.* **2008**, *108*, 1797–1833; b) R. A. W. Frank, F. J. Leeper, B. F. Luisi, *Cell. Mol. Life Sci.* **2007**, *64*, 892–905; c) F. Jordan, *Nat. Prod. Rep.* **2003**, *20*, 184–201.
- [2] As yet, a second enzyme for the oxidative step cannot be excluded.
- [3] S. Fraas, A. K. Steinbach, A. Tabbert, J. Harder, U. Ermler, K. Tittmann, A. Meyer, P. M. H. Kroneck, *J. Mol. Catal. B: Enzym.* **2009**, *61*, 47–49.
- [4] A. K. Steinbach, S. Fraas, J. Harder, A. Tabbert, H. Brinkmann, A. Meyer, U. Ermler, P. M. H. Kroneck, *J. Bacteriol.* **2011**, *193*, 6760–6769.
- [5] A. K. Steinbach, S. Fraas, J. Harder, E. Warkentin, P. M. H. Kroneck, U. Ermler, *FEBS J.* **2012**, *279*, 1209–1219.
- [6] S. Fraas, *PhD thesis*, Universität Konstanz (Germany), **2009**.
- [7] J. Harder, *Arch. Microbiol.* **1997**, *168*, 199–204.
- [8] a) M. Widmann, R. Radloff, J. Pleiss, *BMC Biochem.* **2010**, *11*, 9; b) <http://www.teed.uni-stuttgart.de/>.
- [9] P. Lehwald, *PhD thesis*, Albert-Ludwigs-Universität Freiburg (Germany), **2009**.
- [10] J. Alonso, J. L. Barredo, P. Armisen, B. Díez, F. Salto, J. M. Guisán, J. L. García, E. Cortés, *Enzyme Microb. Technol.* **1999**, *25*, 88–95.
- [11] CDH could be lyophilized without any significant loss of activity, and could be stored without additives in buffer A at –20 °C.
- [12] a) M. Pohl, B. Lingen, M. Müller, *Chem. Eur. J.* **2002**, *8*, 5288–5295; b) M. Pohl, C. Dresen, M. Beigi, M. Müller in *Enzyme Catalysis in Organic Synthesis*, 3rd ed. (Eds.: K. Drauz, H. Gröger, O. May), Wiley-VCH, Weinheim, **2012**, 919–946; c) M. Müller, D. Gocke, M. Pohl, *FEBS J.* **2009**, *276*, 2894–2904.
- [13] P. Lehwald, M. Richter, C. Röhr, H.-w. Liu, M. Müller, *Angew. Chem.* **2010**, *122*, 2439–2442; *Angew. Chem. Int. Ed.* **2010**, *49*, 2389–2392.
- [14] V. Kren, D. H. G. Grout, H. Dalton, D. W. Hutchinson, W. König, M. M. Turner, G. Dean, N. Thomson, *J. Chem. Soc. Chem. Commun.* **1993**, 341–343.
- [15] P. Bisel, L. Al-Momani, M. Müller, *Org. Biomol. Chem.* **2008**, *6*, 2655–2665.
- [16] T. Dünwald, A. S. Demir, P. Siegert, M. Pohl, M. Müller, *Eur. J. Org. Chem.* **2000**, 2161–2170.
- [17] M. Beigi, S. Loschonsky, P. Lehwald, V. Brecht, S. L. A. Andrade, F. J. Leeper, W. Hummel, M. Müller, *Org. Biomol. Chem.* **2013**, *11*, 252–256.
- [18] C. M. Dresen, *PhD thesis*, Albert-Ludwigs-Universität Freiburg (Germany), **2008**.

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