

Engineering Enzymes for Non-Natural Carboligation Reactions



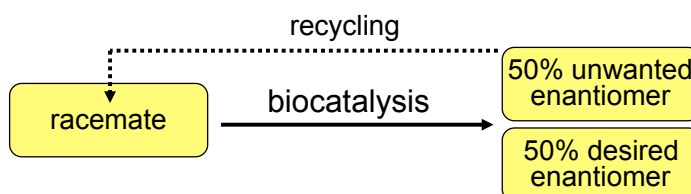
Wolf-Dieter Fessner
TU Darmstadt
Germany

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Routes to Chiral Building Blocks

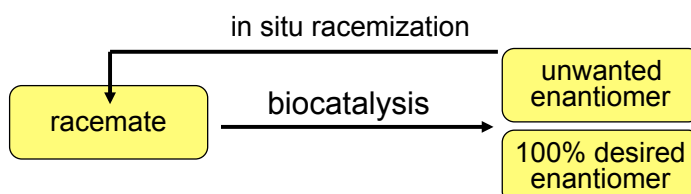
kinetic resolution

several steps
50% yield limit



dynamic resolution

several enzymes
100% yield



asymmetric synthesis

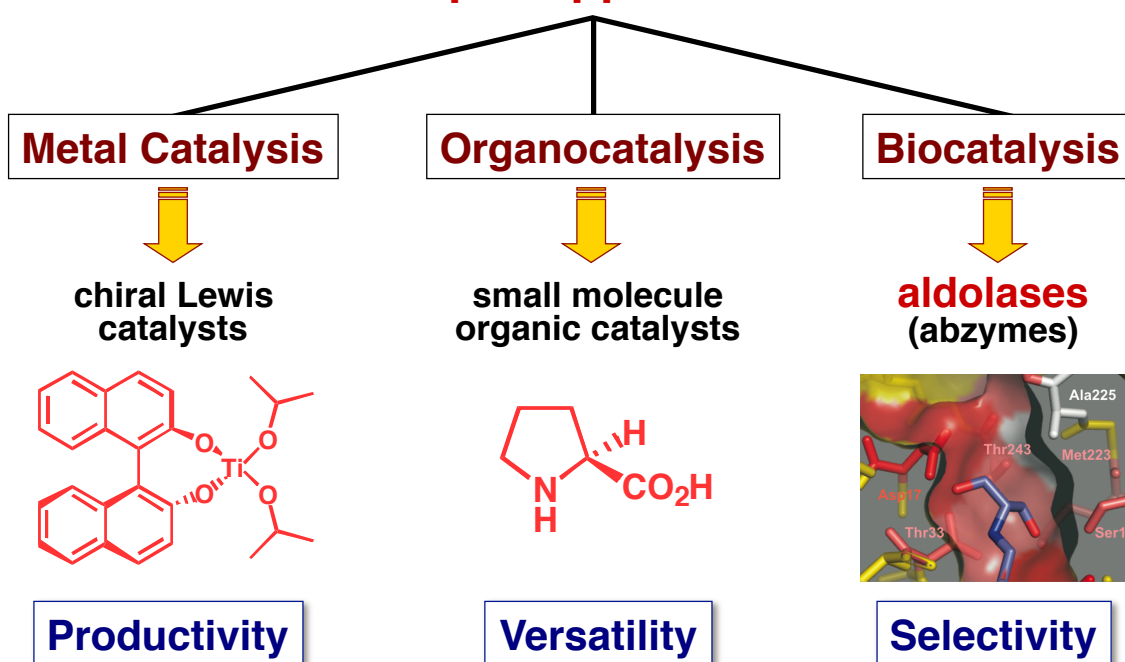
one enzyme, one step
100% yield



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Catalytic Asymmetric Carboligation

Multiple Opportunities



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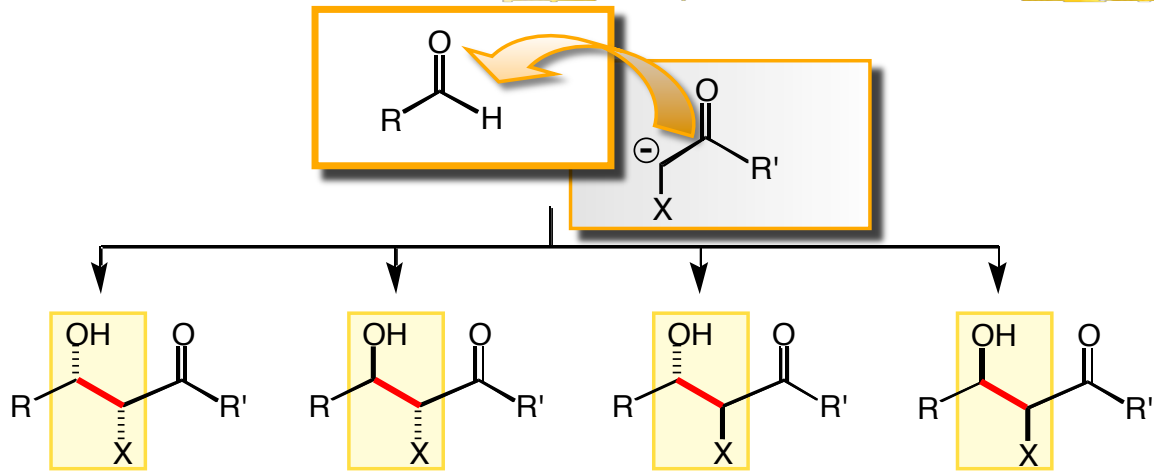
Overview

■ Asymmetric Carboligation

- Fructose-6-phosphate Aldolase from *E. coli*
- Transketolase from *G. stearothermophilus*

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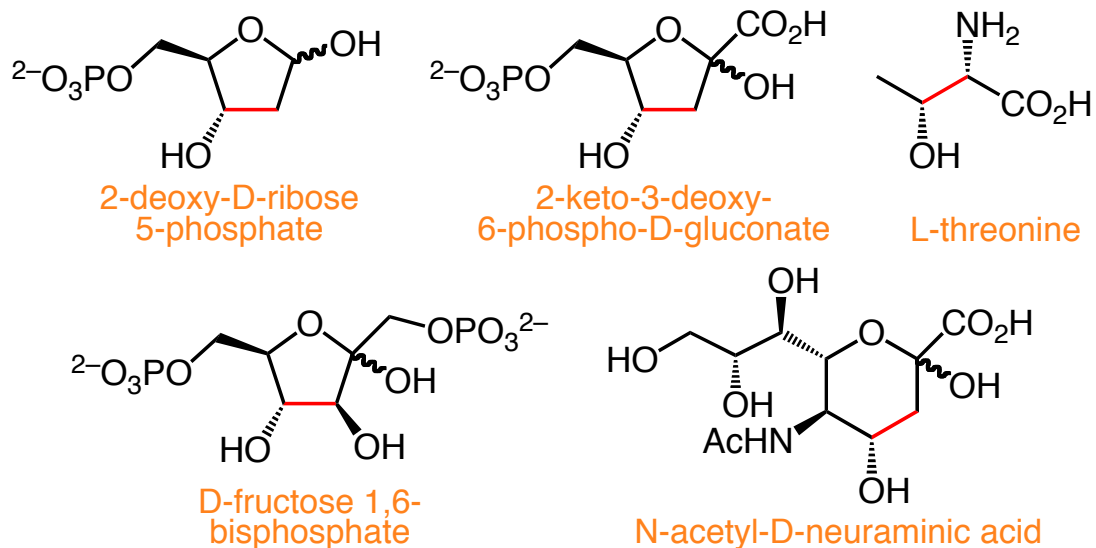
Aldol Opportunity



high stereoselectivity, fully controlled by enzyme
 broad substrate tolerance for acceptor component
 many enzymes well studied & commercially available

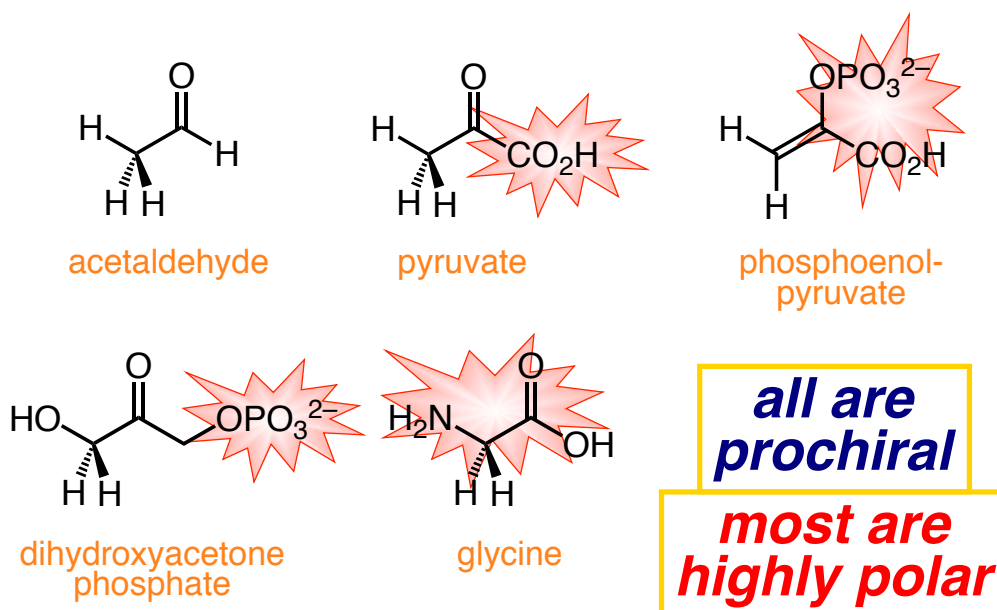
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Nature's Aldol Products



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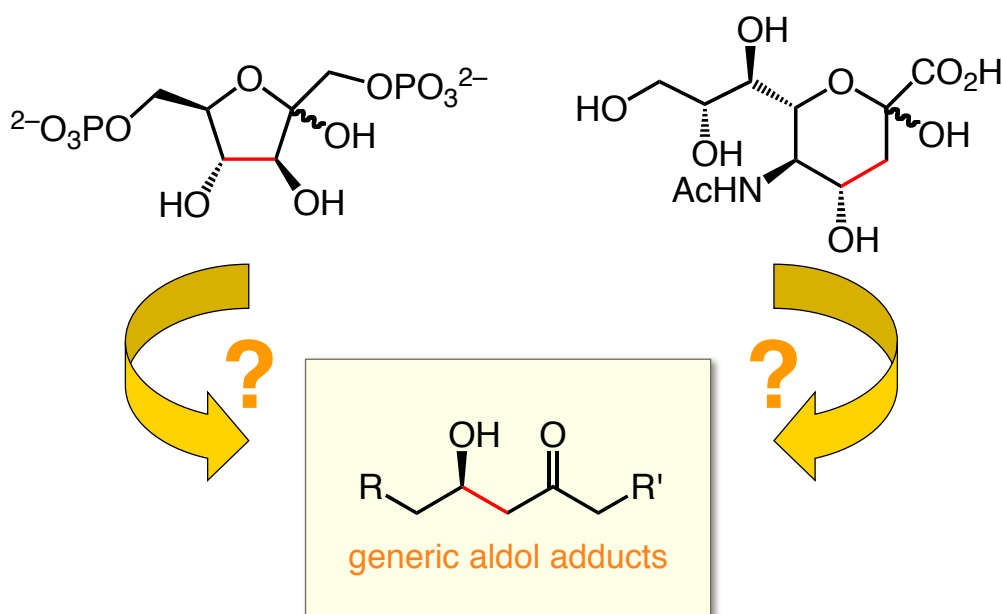
Nature's Nucleophiles



Fessner et al. *Curr. Opin. Chem. Biol.*, 2010, 14, 154-167

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The Challenge



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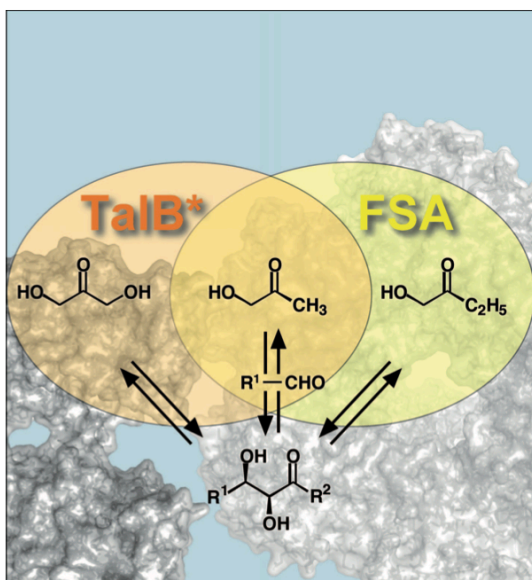
New Enzymes: FSA + TaIB^{F178Y}

CHEMBIOCHEM

DOI: 10.1002/cbic.201100072

The Transaldolase Family: New Synthetic Opportunities from an Ancient Enzyme Scaffold

Anne K. Samland,^[a] Madhura Rale,^[b] Georg A. Sprenger,^[a] and Wolf-Dieter Fessner^{*(b)}

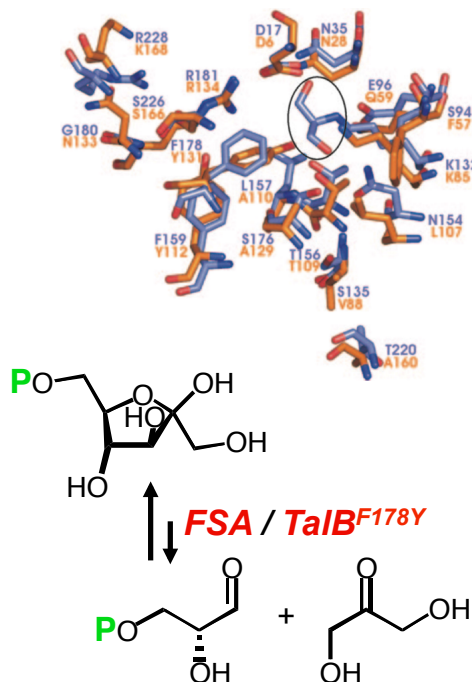


1454

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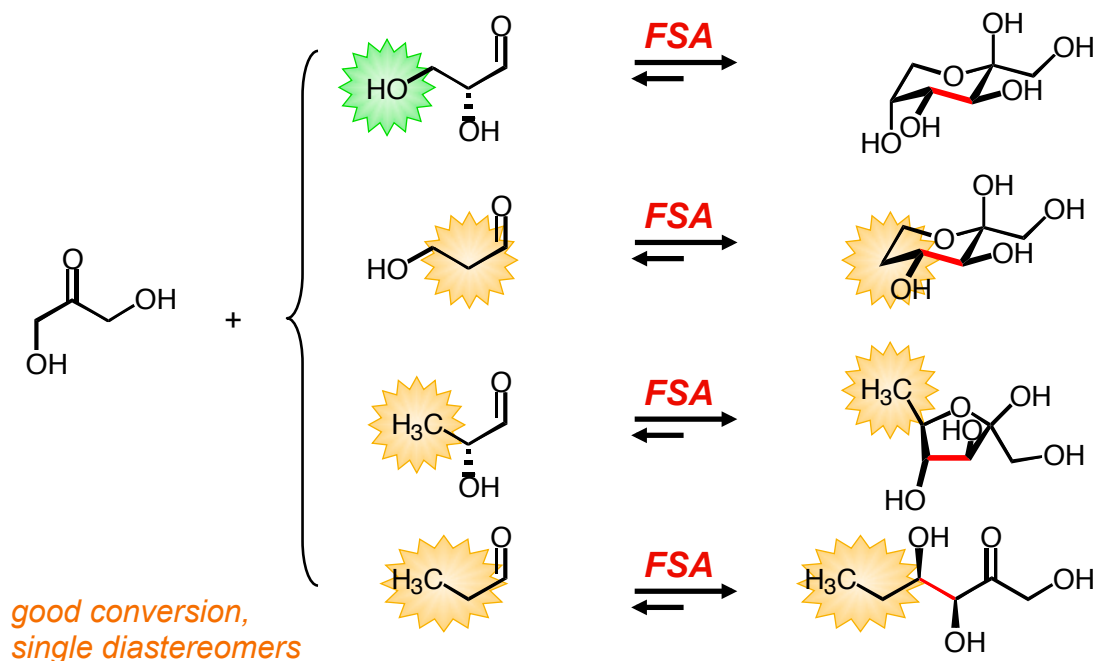
© 2011 Wiley-VCH Verlag GmbH & Co. KGaA, Weinheim

ChemBioChem 2011, 12, 1454–1474



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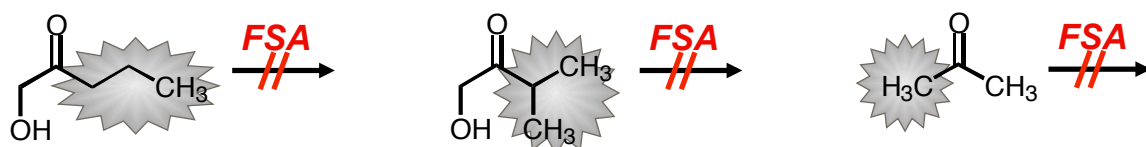
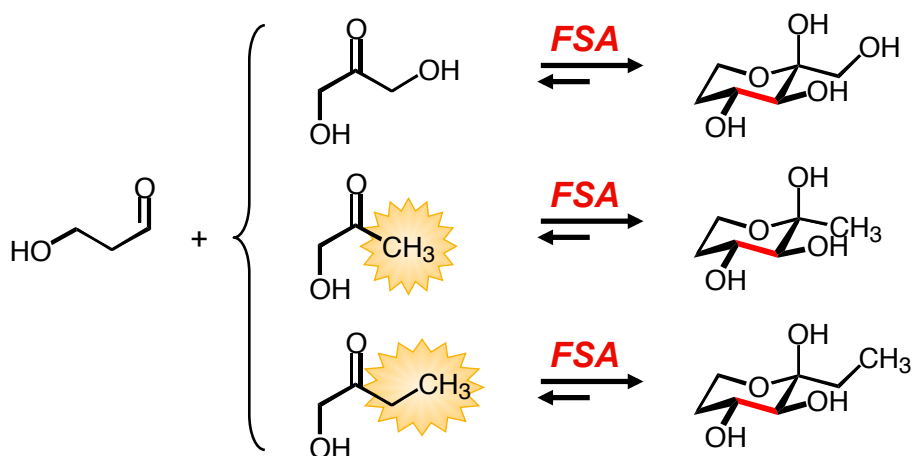
Substrate Tolerance: Electrophiles



Fessner et al. *Chem. Eur. J.*, 2011, 17, 2623-2632

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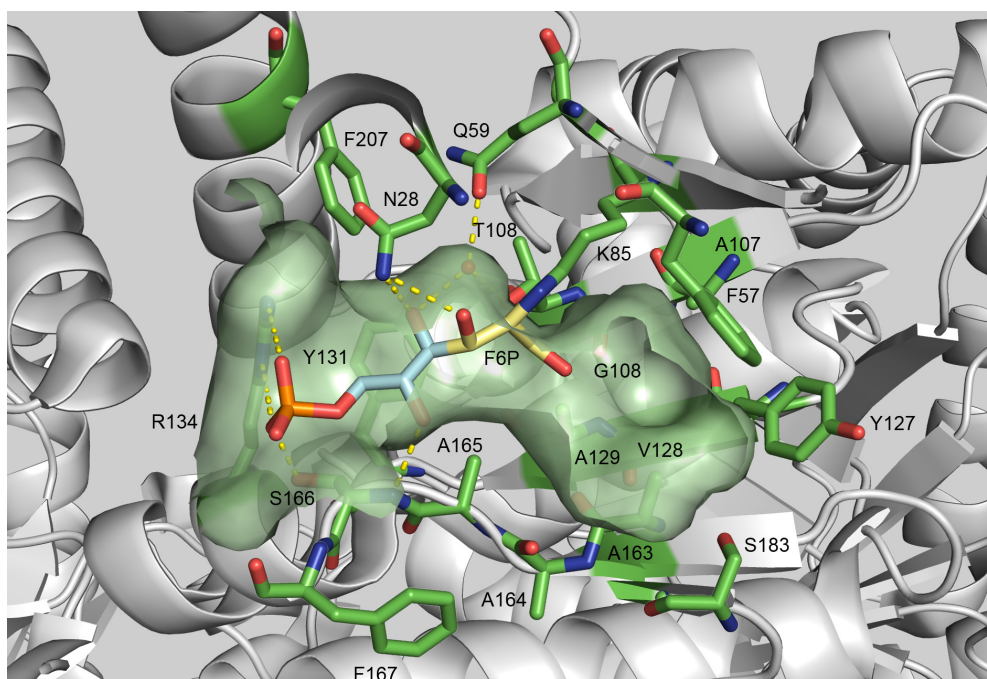
Substrate Tolerance: Nucleophiles



Fessner et al. *Chem. Eur. J.*, 2011, 17, 2623-2632

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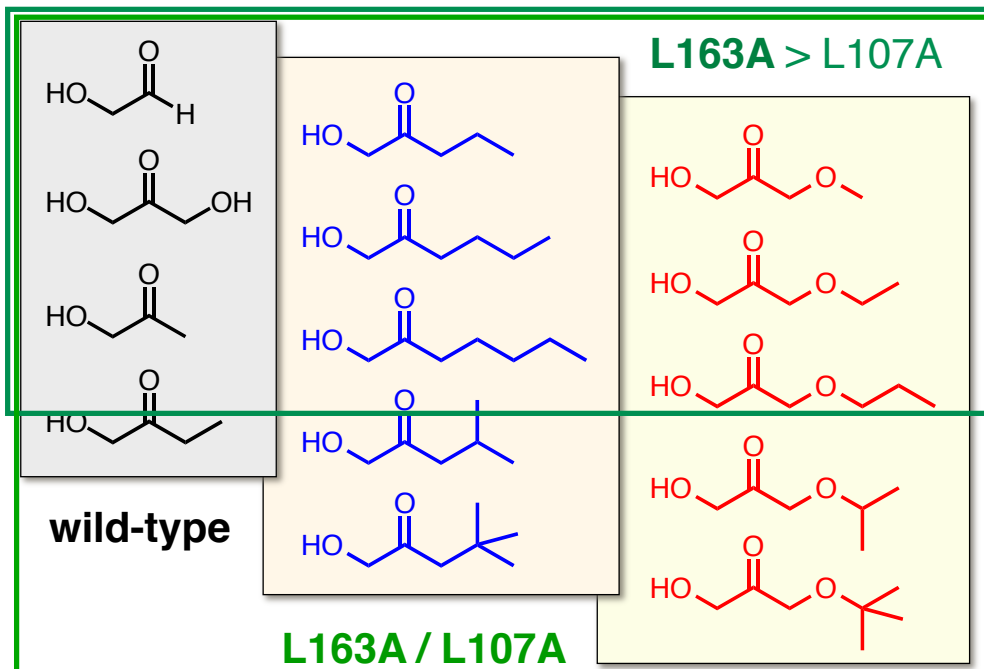
Protein Engineering



structure-based active-site mutations to expand donor tolerance

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Donor Variations



Fessner et al. *manuscript in preparation*

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Protein Stability

**Destabilization by Leu→Ala
Cavity-Creating Mutations:**

Matthews et al.
Science
1992, 255, 178-183

$$\Delta\Delta G = 2.0 \text{ kcal mol}^{-1} + n \times (0.03 \text{ kcal mol}^{-1})$$

transfer constant

cavity volume term per Å³

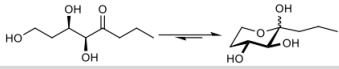
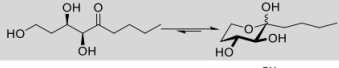
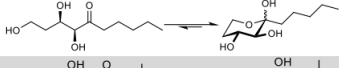
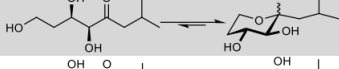
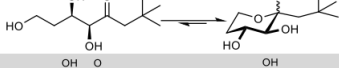
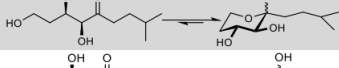
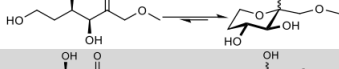

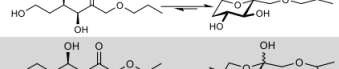
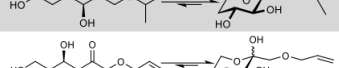
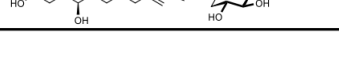
Variant	T _m [°C]
FSA _{wt}	87.0
FSA (L107A)	90.4
FSA (L163A)	78.4
FSA (L107A / L163A)	81.4

**Differential
Scanning
Fluorimetry**

Pantoliano et al.
J. Biomol. Screening
2001, 6, 429-440

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Product Range

	Isolated yield	ratio OC/ α/β
	75%	67/5/34
	76%	54/5/41
	50%	62/4/34
	28%	64/2/34
	25%	66/2/32
	25%	54/5/41
	89%	-/1/99
	82%	-/-/100
	28%	-/-/100
	30%	-/3/97
	45%	-/1/99

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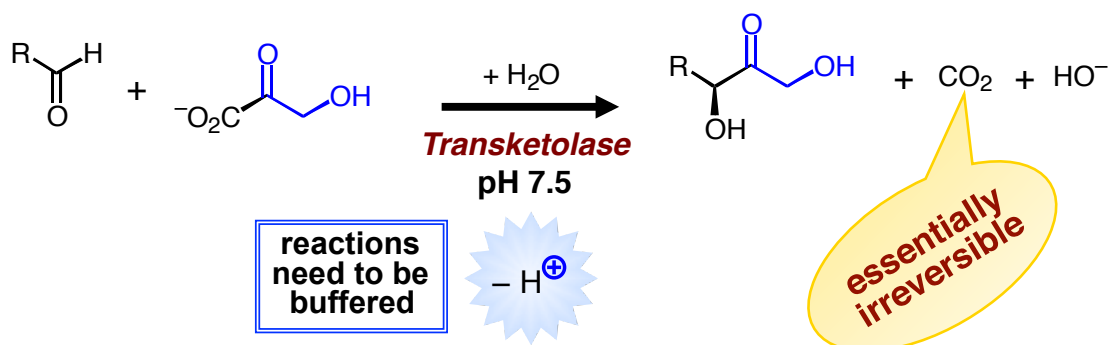
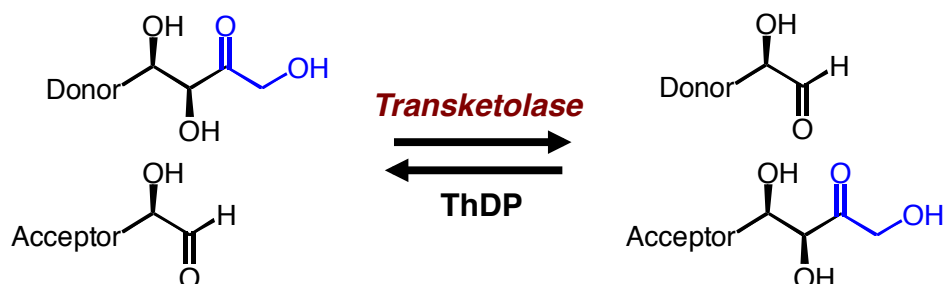
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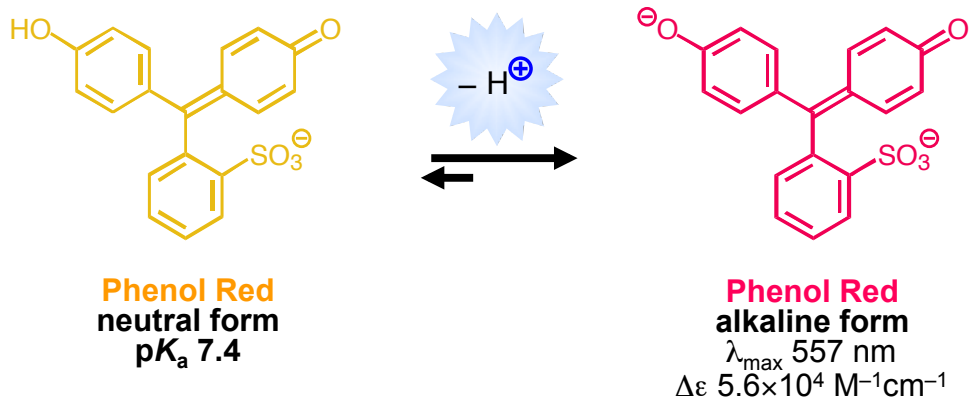
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New Enzymes: Transketolase

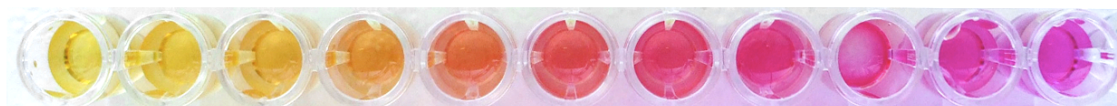


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pH Dependent Assay Technology

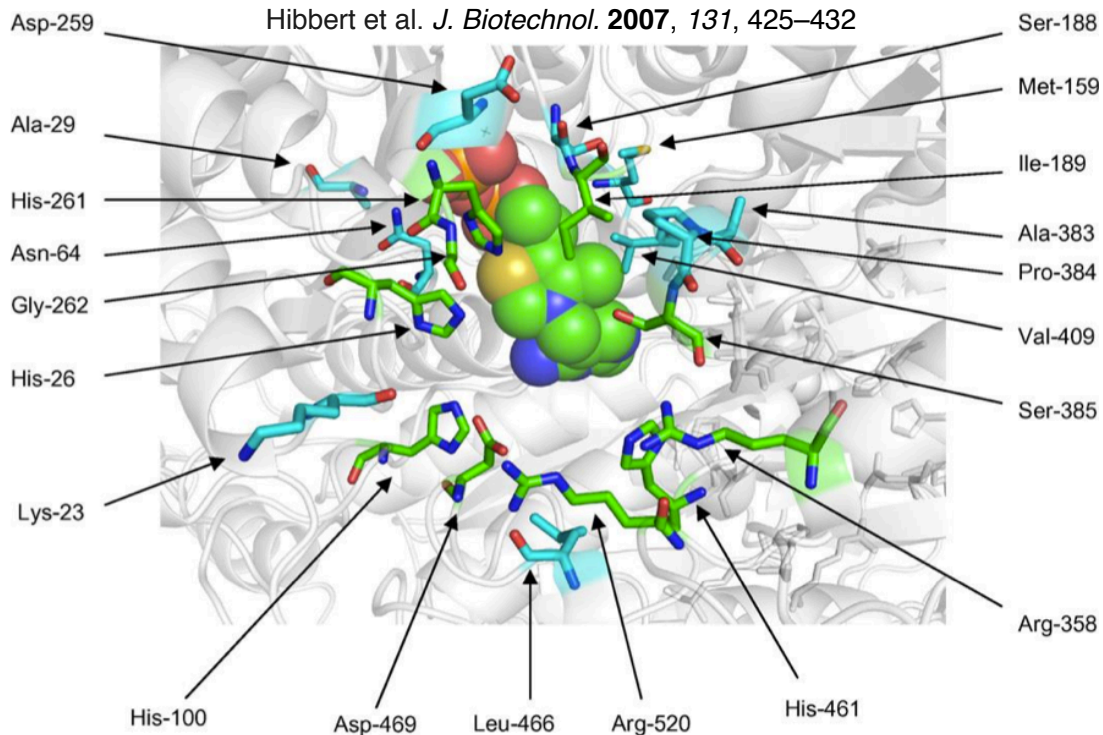


pH 6.42 6.61 6.83 7.02 7.27 7.50 7.68 7.89 8.13 8.32 8.73



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Catalytic Environment



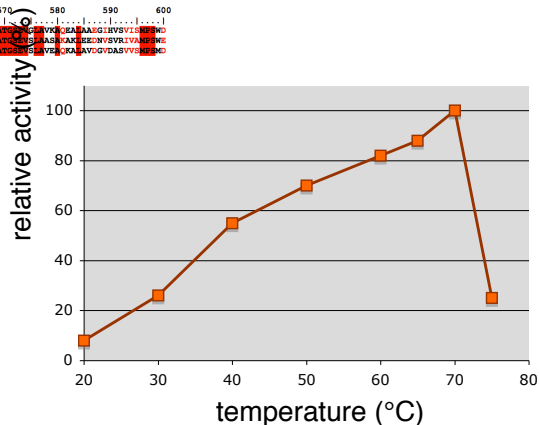
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TK from *G. stearothermophilus*

	10	20	30	40	50	60	70	80	90	100	110	120
GetTK	RSLS	EL	IF	ST	IK	ST	ST	ST	ST	ST	ST	ST
BanTK1	RSLS	EL	IF	ST	IK	ST	ST	ST	ST	ST	ST	ST
BanTK2	RSLS	EL	IF	ST	IK	ST	ST	ST	ST	ST	ST	ST

high similarity to
transketolase gene
from *B. anthracis*

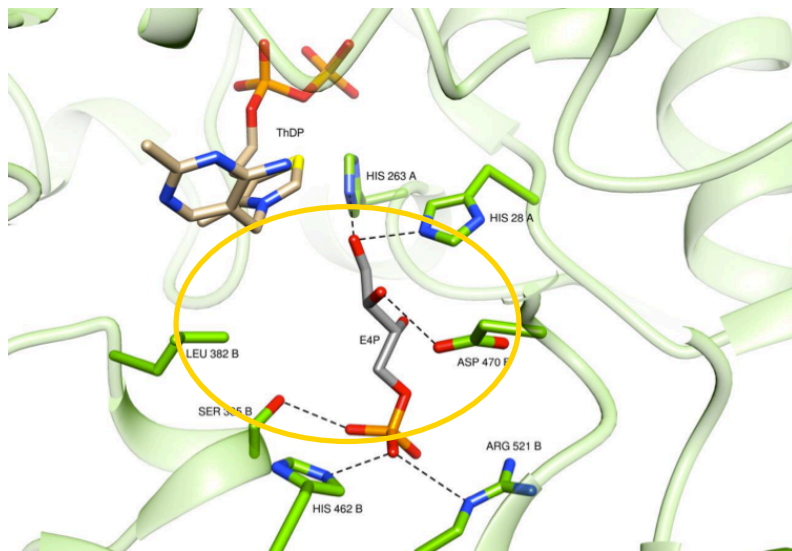
excellent protein over-
production in *E. coli*
highly thermostable
enzyme activity



Fessner et al. *Adv. Synth. Catal.*, 2013, 355, 116–128

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Acceptor Discrimination

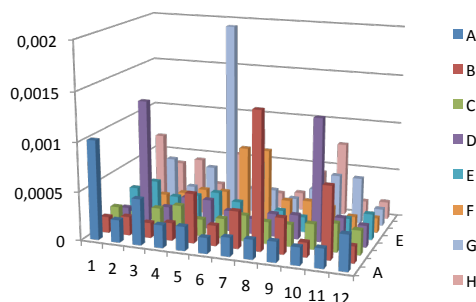
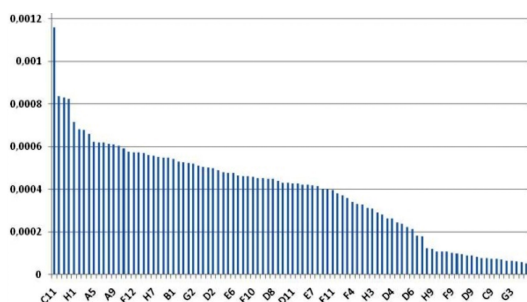


hydrogen bonding contacts to (*R*)-hydroxyaldehydes

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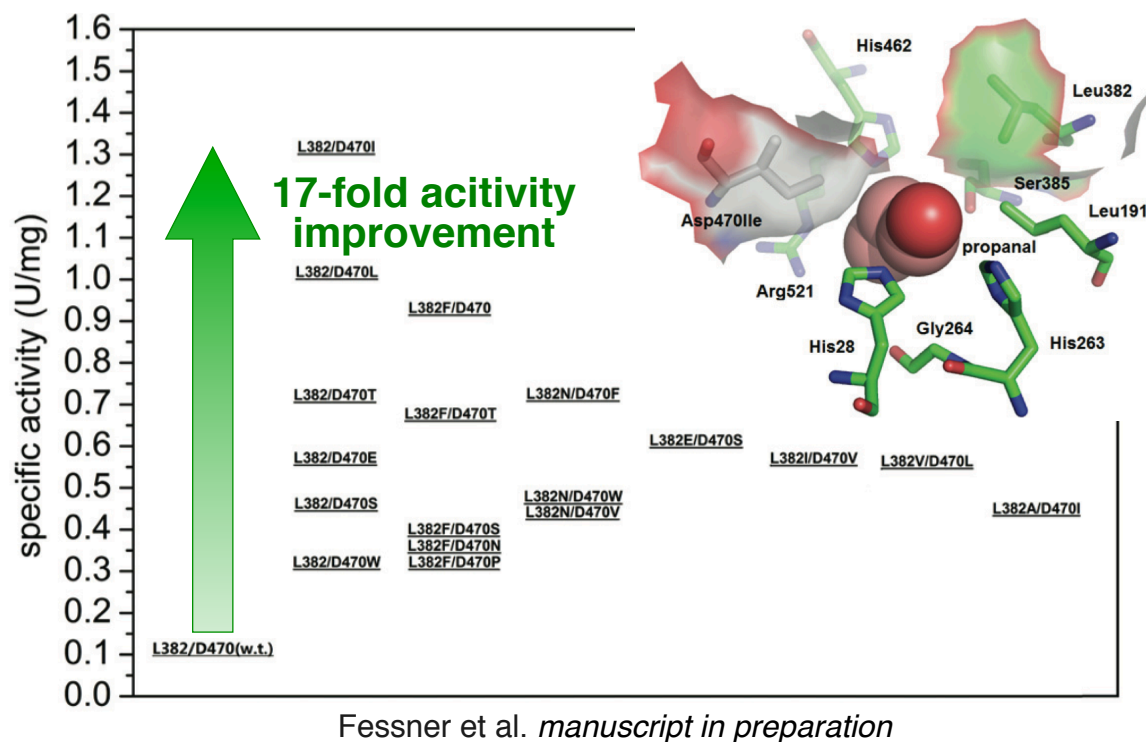
Improved TK Variants

- structure-guided molecular evolution
- focused “small, but smart” libraries
- saturation mutagenesis at Leu381/Asp469
- residues involved in binding of 2-hydroxyaldehyde
- >3,400 colonies (>95% coverage)
- screening for propanal / butanal / methoxyethanal



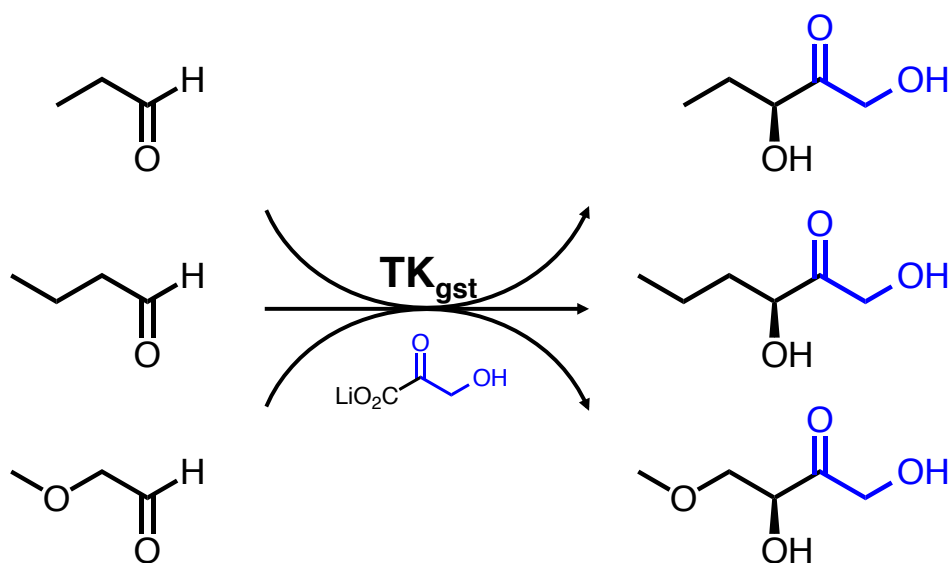
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Hydrophobic Substrates



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Less Polar Products



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Summary

- **novel carbonylation catalysts**
 - thermostable aldolase FSA
 - thermostable transketolase
- **broad substrate tolerance**
 - engineered donor tolerance for FSA
 - engineered acceptor tolerance for TK
- **high enantio- and diastereoselectivity**
- **one-step creation of structural complexity**

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Acknowledgments



Dr. Madhura Rale
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BMBF / DAAD
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Abbott



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