

Enzymatische Modifikationen von Lebensmittelinhaltsstoffen: Verfahren zur Beeinflussung des Produktdesigns

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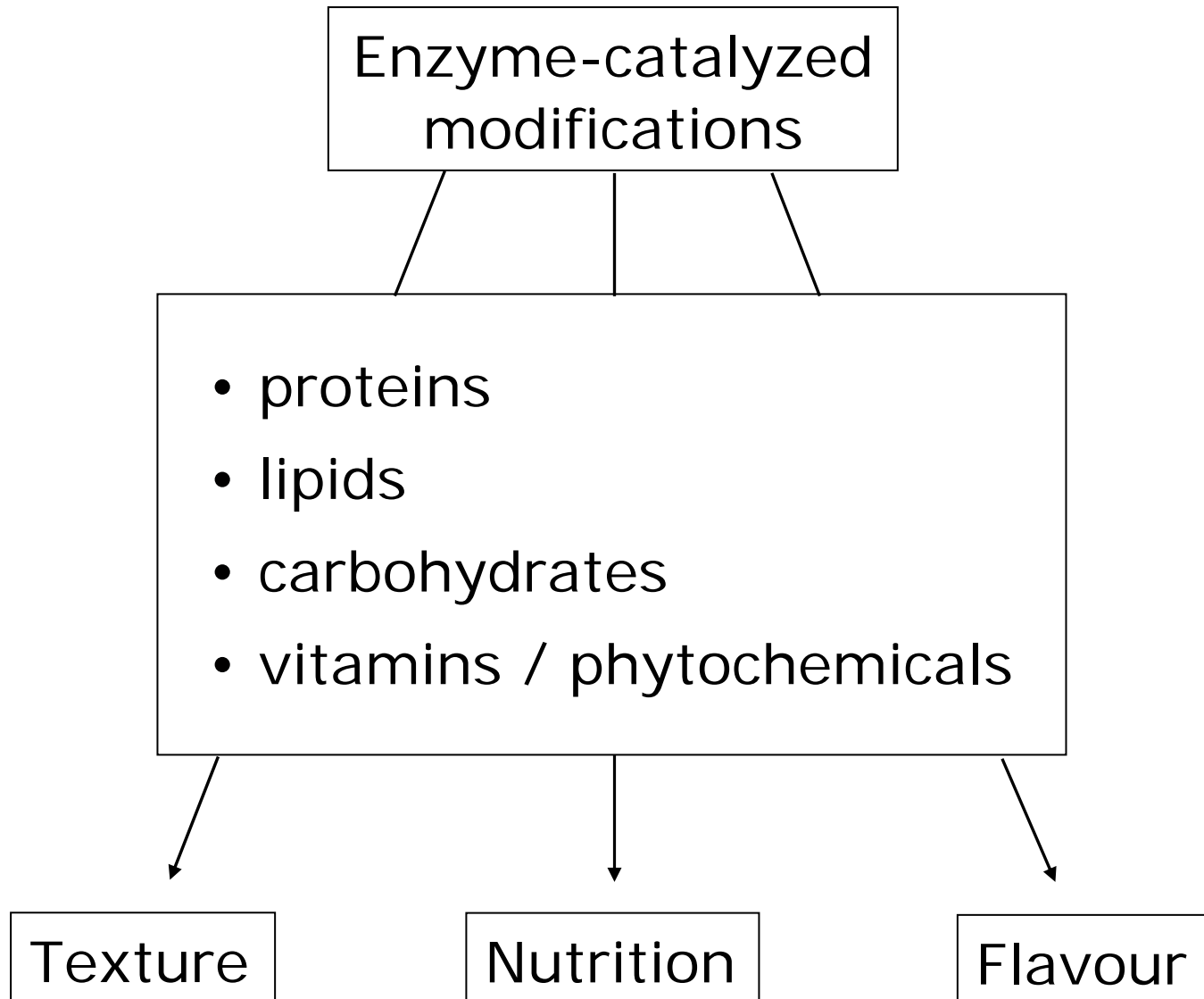
Lehrstuhl für Allgemeine Lebensmitteltechnologie

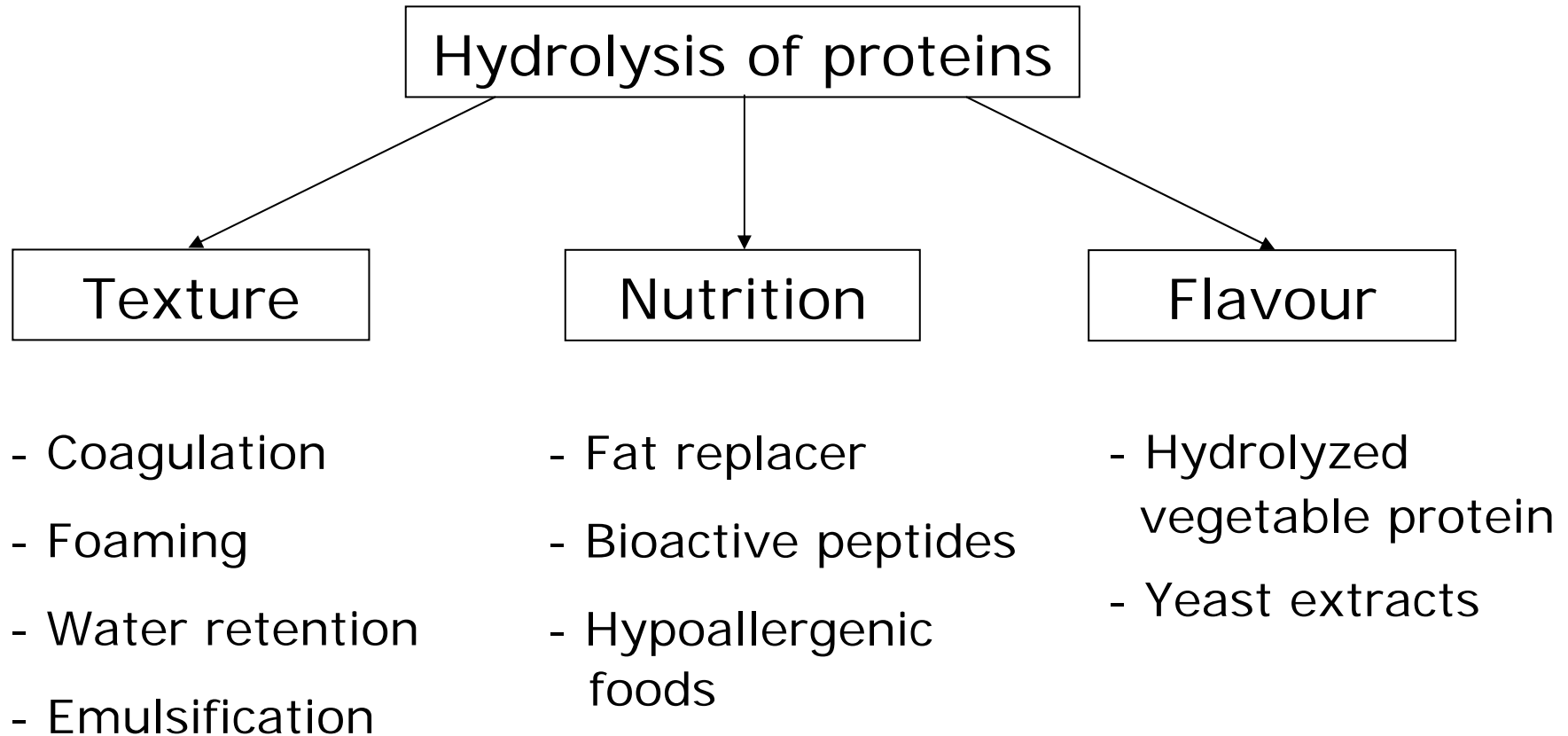
(\$ Millions)

	2005	2006	2007	2012
Technical Enzymes	1,075	1,105	1,140	1,355
Food Enzymes	775	800	830	1,010
Animal Feed Enzymes	240	260	280	375
Total	2,090	2,165	2,250	2,740

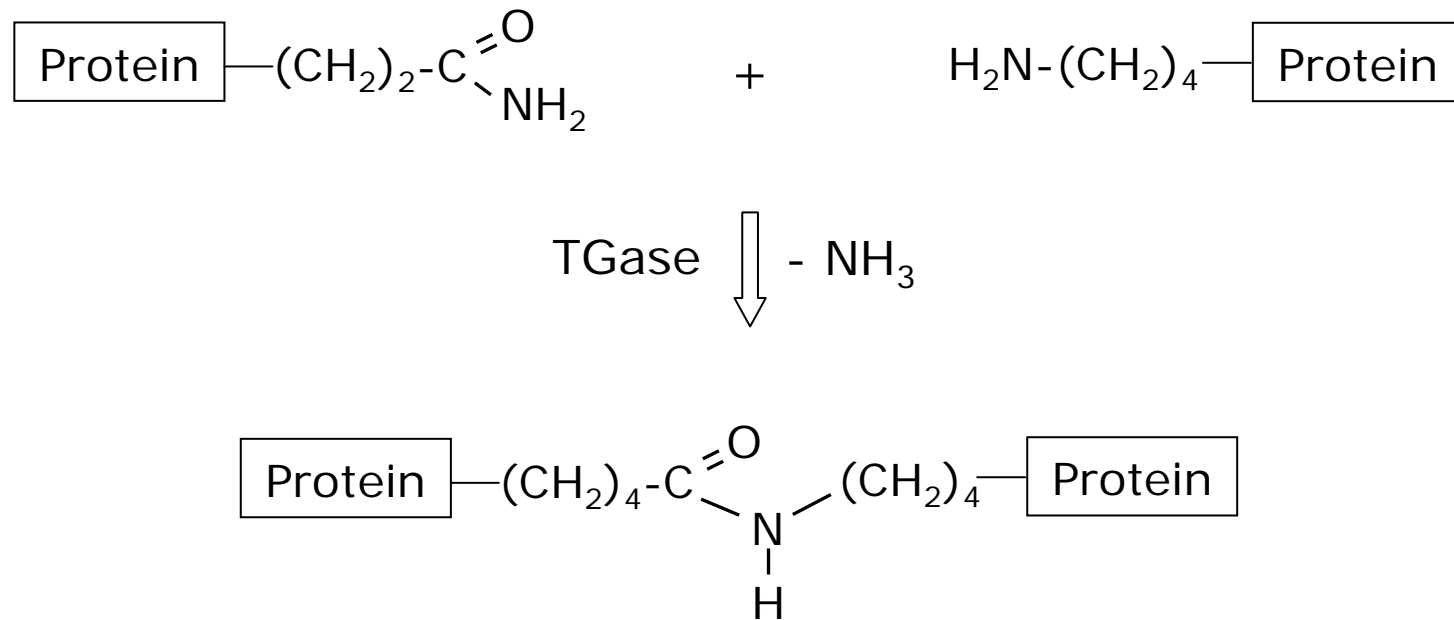
Source: BCC, Inc.

Conversion / enzyme type	Sales (%)	Main applications
Protein hydrolysis		
Alkaline proteases (e.g. subtilisin)	37.0	Detergents
Rennets (calf, microbial)	10.5	Cheesemaking, dairy
Plant proteases (papain, bromelain)	0.7	Food processing
Pancreatic proteases	1.0	Leather processing
	49.2	
Carbohydrate conversion		
Amylases	16.7	Starch processing, baking, brewing, textile, detergents
Pectinases	3.0	Beverages
Cellulases	15.3	Detergents, textile, baking
Xylanases / β -glucanases	4.5	Animal feed processing, brewing
Glucose isomerase	2.0	Food and beverages
Pullulanases	0.5	Starch
Phytases	1.2	Animal feed processing
Lactases	0.9	Dairy
	44.1	
Fat hydrolysis		
Lipases	7.0	Fat processing, dairy, detergents, organic synthesis





- Formation of isopeptide bonds



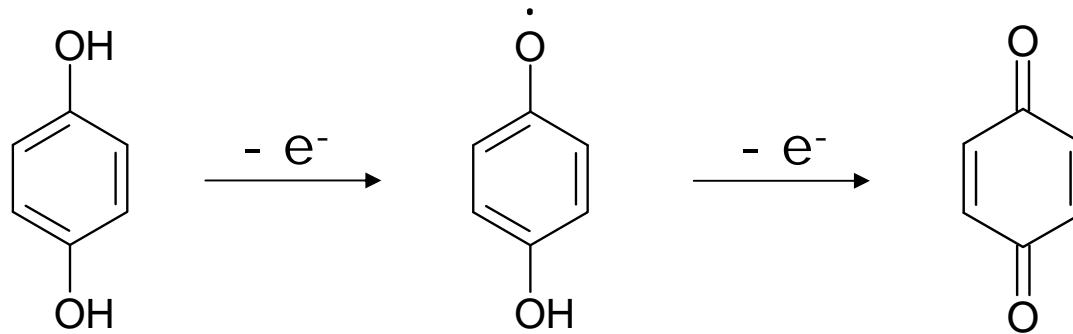
- Microbial Transglutaminase (*Streptoverticillium mobaraense*)



- viscosity
- gelation
- water retention
- syneresis

- meat products
 - sausages
 - re-structured meat
- dairy products
 - yoghurt
 - cheese
- bakery products
- pasta

Laccase

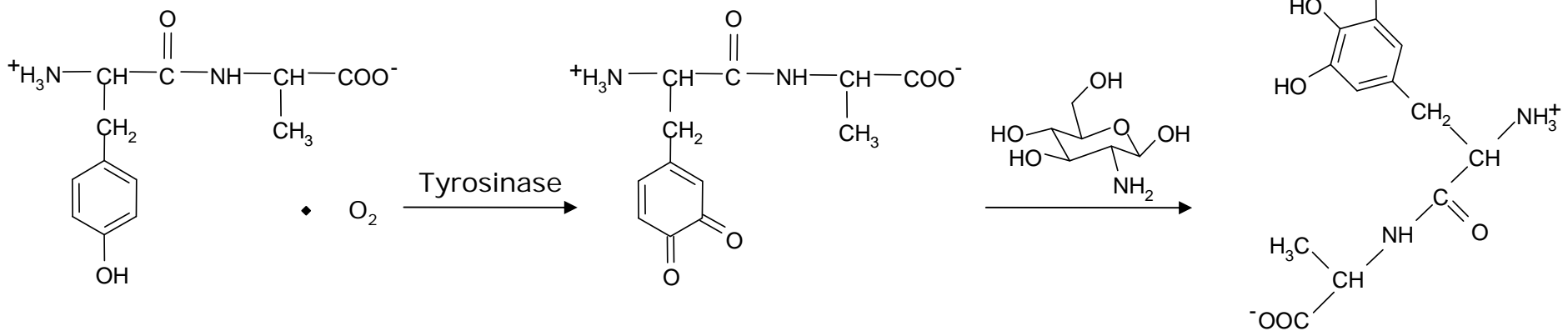
(p-diphenol oxidase)

- Wheat arabinoxylan gels (Carvajal-Millan et al., 2005)
- Sugar beet pectin gels (Kuuva et al., 2003)
- Gelatin-catechin conjugates (Chung et al., 2003)

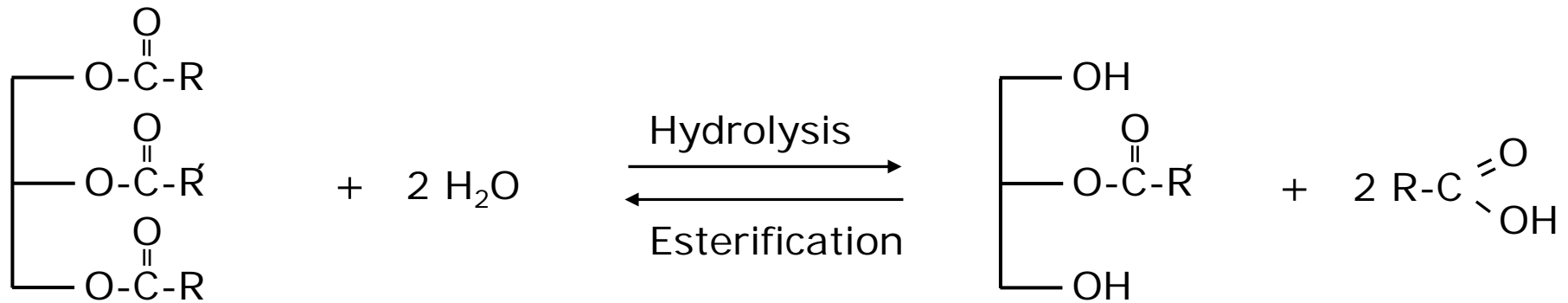
Tyrosinase

- Oxidation of peptide-bond tyrosine to quinones
- reaction with free -SH and -NH₂ groups
- tyrosine-cysteine and tyrosine-lysine crosslinks

- Enzymatic grafting of peptides from casein hydrolysates to chitosan (Aberg et al., 2004)

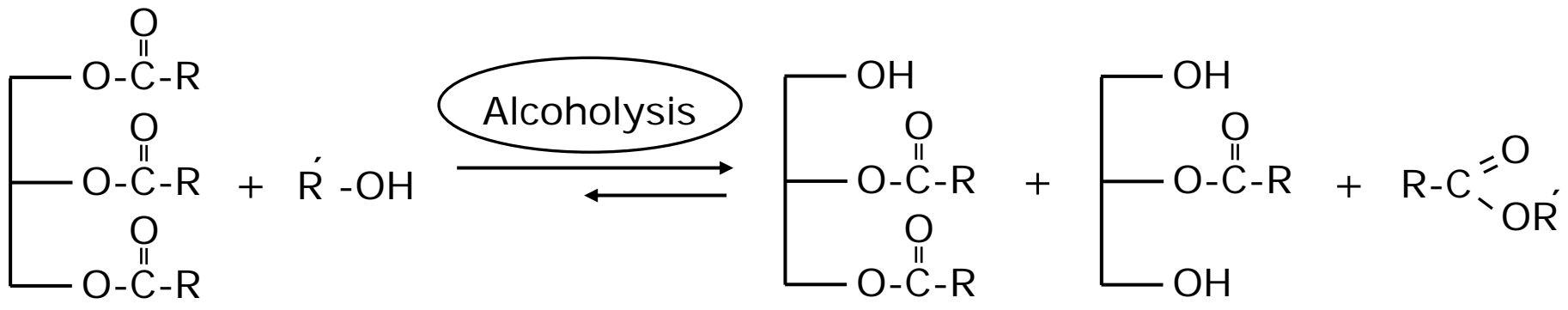
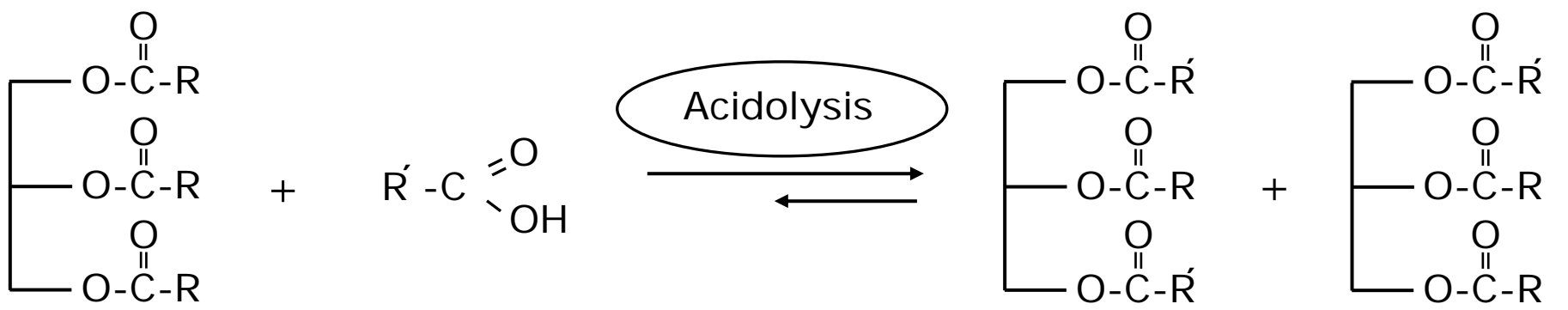
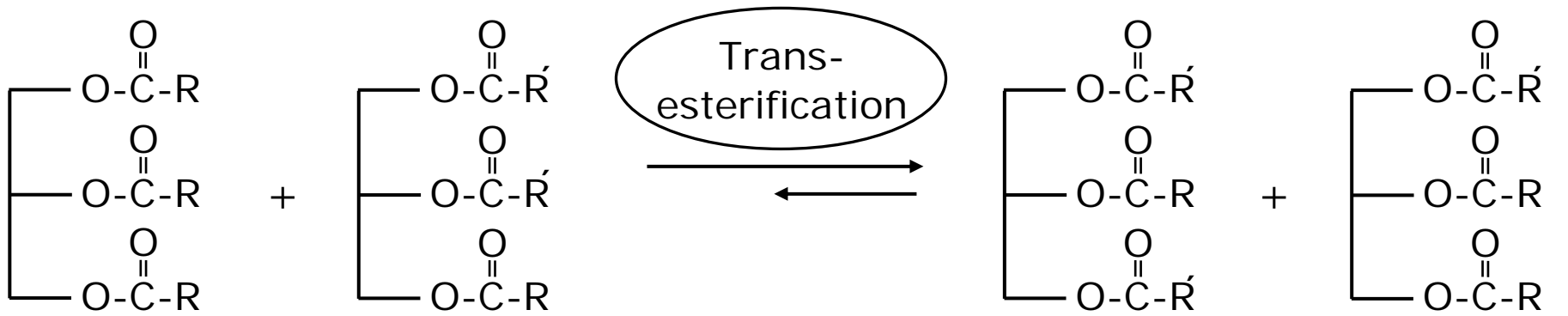


Proposed tyrosinase-initiated reaction between glucosamine and Tyr-Ala



Specificities

- position of fatty acid (e.g. 1,3-regiospecific)
- chain length of fatty acid
- number / position of double bonds in fatty acid





- Structured
 - triglycerides
 - mono-/ diglycerides
 - phospholipids
 - sphingolipids
- Incorporation of
 - short / medium chain fatty acids
 - polyunsaturated fatty acids (EPA, DHA, CLA)
- Positioning of fatty acids
 - ⇒ improved / adjusted nutritional value



┌ 8:0 (10:0 or 6:0)
 └ 18:2 $n-6$ (18:3 $n-3$)
 ┌ 8:0 (10:0 or 6:0)

┌ 18:0 / 16:0
 └ 18:1 $n-9$
 ┌ 18:0 / 16:0

┌ 8:0 (10:0 or 6:0)
 └ EPA and/or DHA
 ┌ 8:0 (10:0 or 6:0)

┌ U
 └ 16:0
 ┌ U

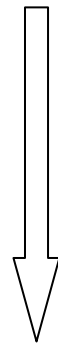
┌ 8:0 (10:0 or 6:0)
 └ CLA
 ┌ 8:0 (10:0 or 6:0)

┌ M
 └ L + ┌ M
 └ M
 ┌ L

Name	Supplier	Organism	Specificity	Matrix
Lipozyme TL IM	Novozyme A/S	<i>Thermomyces lanuginose</i> , TLL-1	sn-1,3 specific	Silica granules
Lipozyme RM IM	Novozyme	<i>Rhizomucor miehei</i> , RML	sn-1,3 specific	Macroporous ion exchange resin
Novozyme 435	Novozyme	<i>Candida antartica</i> lipase B	non-specific	Macroporous acrylic resin
Lipase PS-C	Amano	<i>Pseudomonas cepacia</i> lipase, PCL	non-specific	Ceramic particles
Lipase AK-C	Amano	<i>Burkholderia cepacia</i> lipase (formerly <i>Pseudomonas fluorescens</i> lipase PFL)	sn-1,3 specific	Ceramic particles

α -tocopherol (25 – 100 mM)

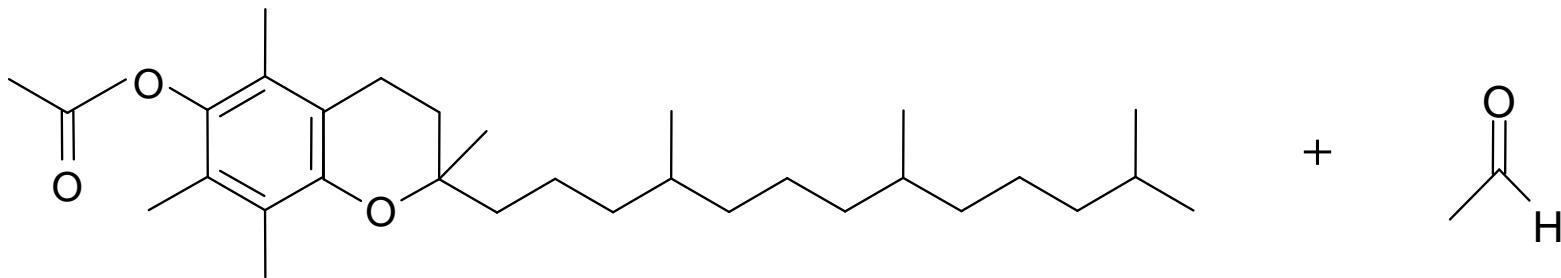
+ vinyl acetate (100 – 400 mM)



C. antarctica

lipase B (Novozym 435)

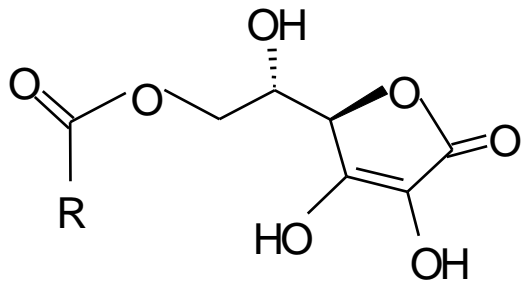
60°C; solvent: 2-methyl-2-butanol



(30%, 8 days)

L-ascorbic acid
+ vinyl palmitate

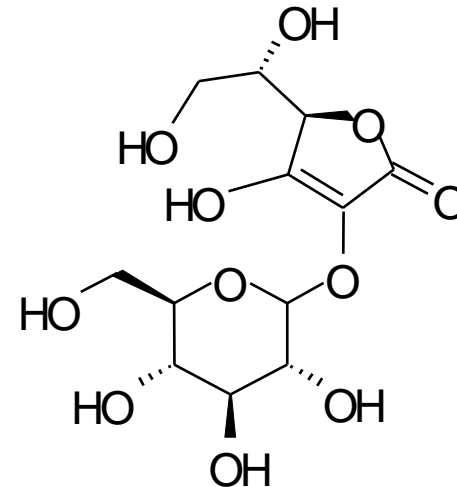
↓
Candida antarctica
lipase B
(Novozym 435)



6-*O*-palmitoyl ascorbate
(Torres et al., 2007)

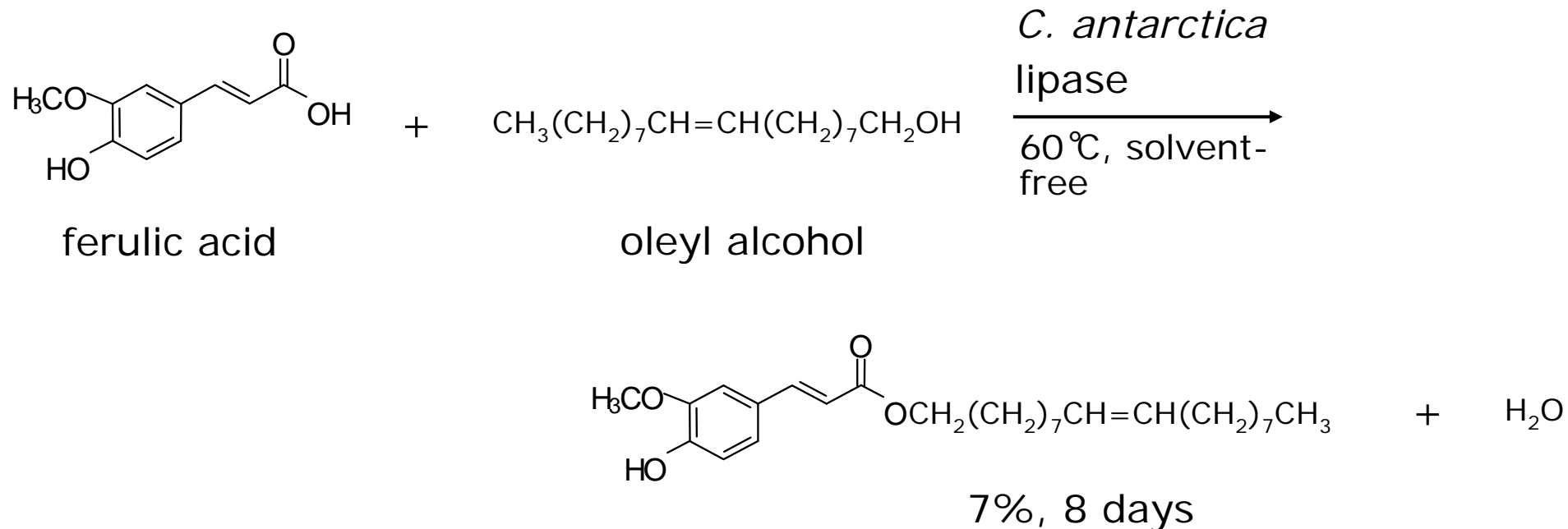
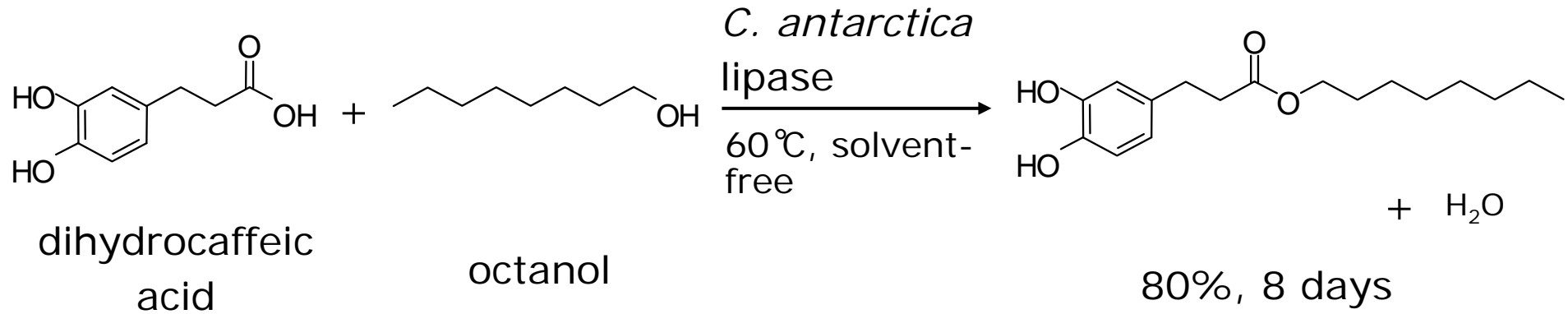
L-ascorbic acid
+ maltotriose

↓
Bacillus stearo-
thermophilus
maltogenic amylase

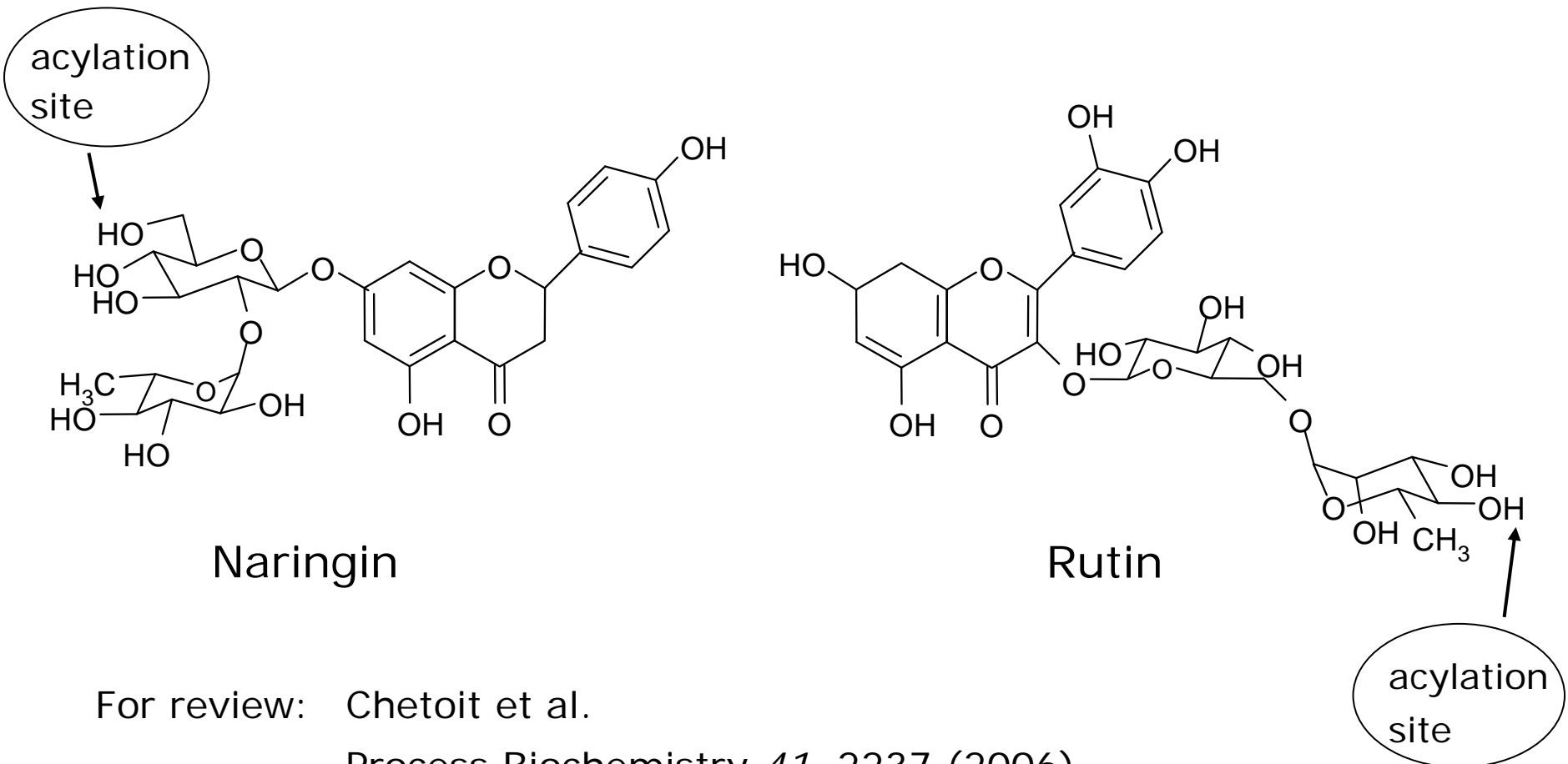


ascorbic acid 2-glucoside
(Bae et al., 2002)

Acyl donor	Solvent	Biocatalyst	Yield [%]
Methyl palmitate	2-methyl-2-butanol	Novozym 435	68
Phenylbutyric acid	<i>t</i> -butanol	Novozym 435	22
Methyl palmitate, EPA and DHA ethyl esters	2-methyl-2-butanol	Novozym 435	≤ 40
Saturated fatty acids	2-methyl-2-pentanol	Novozym 435	≤ 65
Vinyl esters (C8-C16)	<i>t</i> -butanol	Chirazyme L-2	≤ 91
Palmitic acid	hexane	Lipase from <i>Bacillus stearothermophilus</i>	≤ 97
EPA	acetone	Chirazyme L-2	47
L-Methyl lactate	2-methyl-2-butanol	Novozym 435	80
Oleic acid, Palmitic acid	2-methyl-2-butanol	Novozym 435	71-87
Saturated fatty acids (C10-C14)	acetone	Chirazyme L-2	≤ 60
Oleic acid, Linoleic acid, Linolenic acid	acetone	Chirazyme L-2	60
Oleic acid	ionic liquid	Novozym 435	61
Oleic acid, Linoleic acid	2-methyl-2-butanol	Novozym 435	≤ 45



- *C. antarctica* lipase B (Novozym 435)
 - oleic, linoleic, linolenic acid
 - solvents: acetone, *tert*-butanol



For review: Chetoit et al.

Process Biochemistry 41, 2237 (2006)



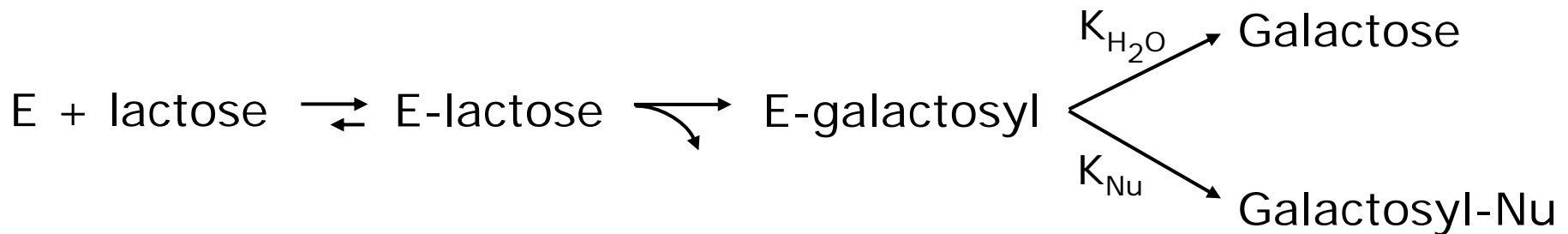
Plant cell wall materials

- Arabinoxylans
(soy, wheat, sorghum)
- Rhamnogalacturans
(apple)
- Arabinan
(sugar beet)

- endo-xylanase
(*A. tubigenis*)
→
- endo-galactanase
(*A. aculeatus*)
- rhamnogalacturonase
(*A. aculeatus*; Ultra-sp.)
→
- endo-arabinase
(*A. aculeatus*)
→

new types of
non-digestible
oligosaccharides

- hydrolysis of lactose
- formation of galacto-oligosaccharides (GOS)
 - transgalactosylation activity in lactose-concentrated solutions



GOS: α -D-Glu (1 \rightarrow 4)-[β -D-Gal (1-6)]_n

n = 2 - 10

Trivial names	Systematic name	Raw material	Functional oligosaccharide
Levansucrase (beta-fructosyltransferase)	Sucrose: 2,6-beta-D-fructan 6-beta-D-fructosyltransferase EC 2.4.1.10	Sucrose	Fructo-
Inulosucrase	Sucrose: 2,1-beta-D-fructan 1-beta-D-fructosyltransferase EC 2.4.1.9	Sucrose	Fructo-
Beta-galactosidase	beta-D-Galactoside galactohydrolase EC 3.2.1.23	Lactose	Galacto-
Beta-xylanase	1,4-beta-D-Xylan xylanohydrolase EC 3.2.1.8	Corn cob	Xylo-
Beta-fructofuranosidase	beta-D-Fructofuranoside fructohydrolase EC 3.2.1.26	Sucrose Lactose	Lacto-fructo
Beta-amylase	1,4-alpha-D-Glucan maltohydrolase EC 3.2.1.2	Starch	Isomalto-
Pullulanase	Pullulan alpha-1,6-glucanohydrolase EC 3.2.1.41		
Transglucosidase (Alpha-glucosidase)	alpha-D-Glucoside glucohydrolase EC 2.4.1.20		
Isomaltulose synthase	Sucrose glucosylmutase EC 5.4.99.11	Sucrose	Isomaltulose (palatinose)
Endoinulinase (inulinase)	2,1-beta-D-Fructan fructanohydrolase EC 3.2.1.7	Inulin	Inulo
Dextransucrase	Sucrose: 1,6-alpha-D-glucan 6-alpha-D-glucosyltransferase EC 2.4.1.5	Sucrose	Isomalto-

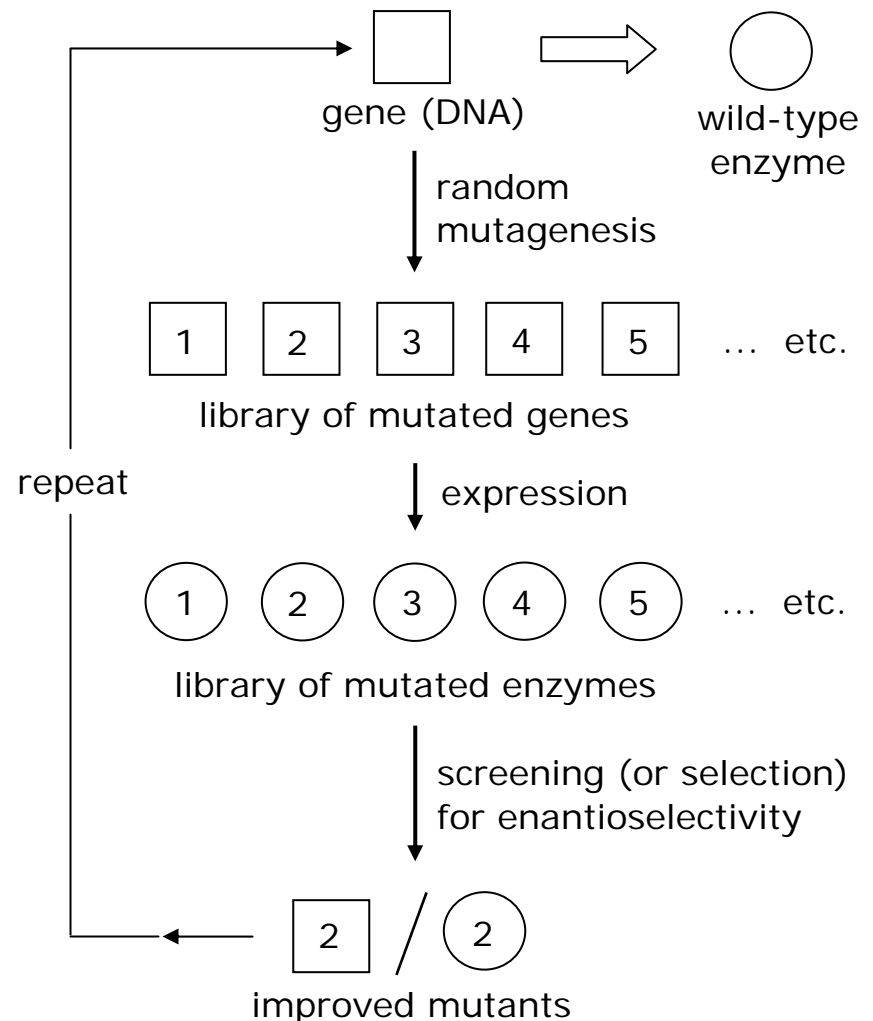


- Isolation of enzyme variants from organisms living in appropriate environments (extremophiles)
- Recombinant DNA technology
 - heterologous expression in suitable host
- Rational protein design
- Directed evolution

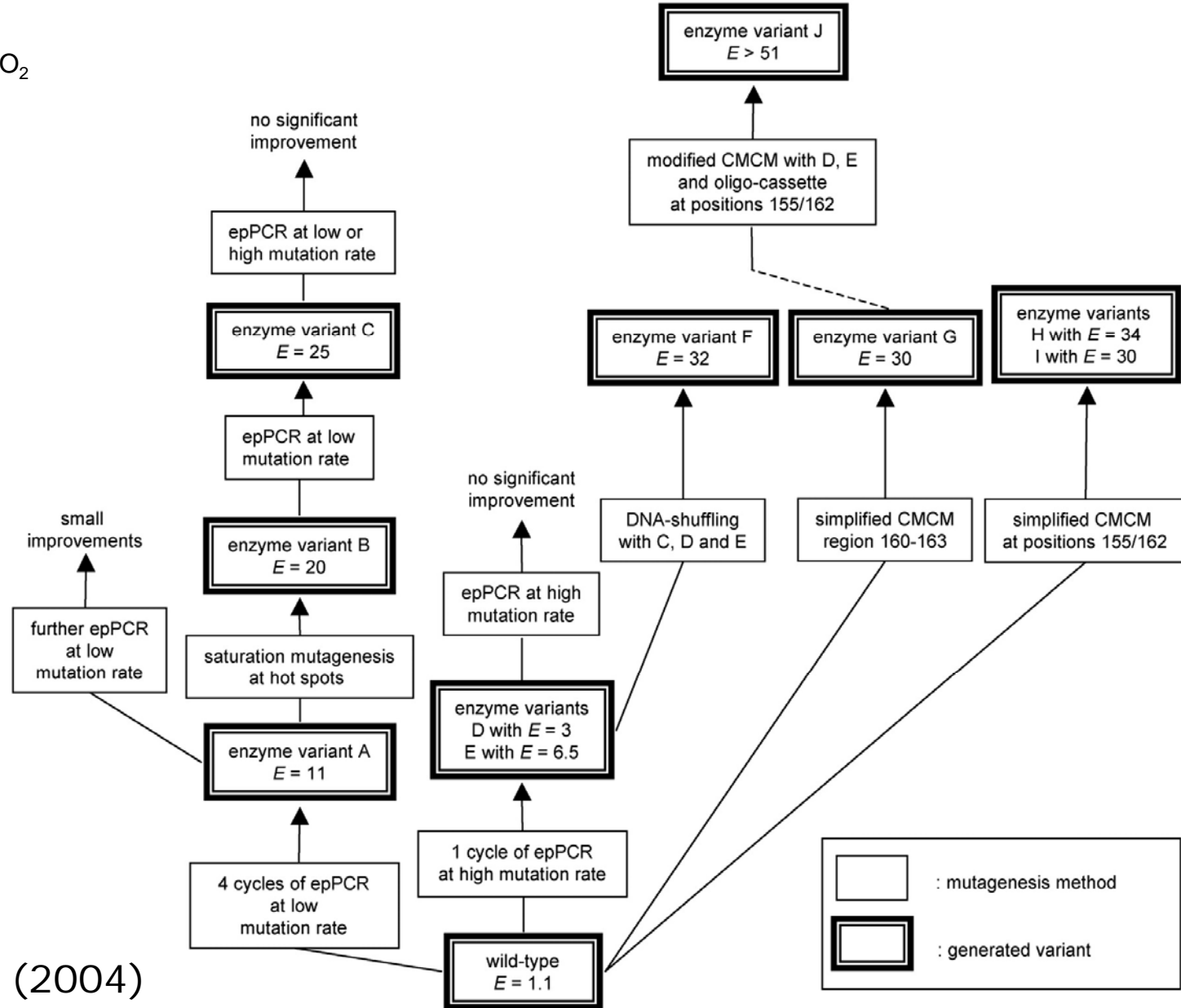
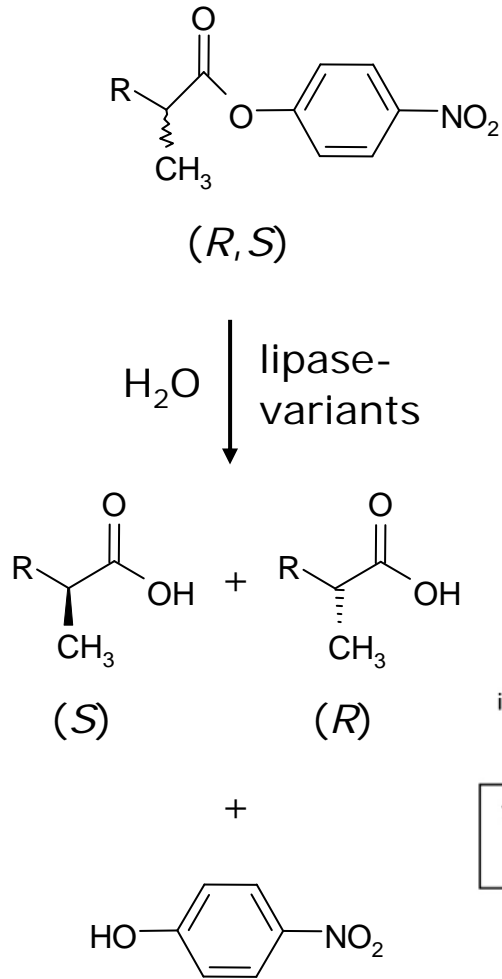


- Rational protein design
 - development of a molecular model on protein structure and function
 - generation of mutants by site-directed mutagenesis
 - transformation of vectors into host organisms
 - expression, purification and analysis of recombinant protein

- Diversification
 - random mutagenesis (error prone PCR)
 - fragmentation
 - reassembling (DNA shuffling)
 ⇒ mutant library
- Selection and screening
- Amplification



Reetz, M. T.
 PNAS 101, 5716 (2004)



Reetz, M. T.
PNAS 101, 5716 (2004)

Microorganism	Desired property	Mutagenesis method
<i>Pseudomonas aeruginosa</i> PA01	Increase in enantioselectivity toward 2-methyldecanoic acid <i>p</i> -nitrophenylester	Error-prone PCR
<i>Bacillus thermoacetenuatus</i> BTL2	Conversion of lipase to phospholipase activity	Random mutagenesis
<i>Staphylococcus hyicus</i>	Improvement in phospholipase activity	Error-prone PCR + DNA shuffling
<i>Thermomyces lanuginosus</i>	Improvement of properties in presence of detergents	Combinatorial protein engineering / phage display approach
<i>Bacillus subtilis</i>	Enantioselectivity toward <i>meso</i> -1,4-diacetoxy-2-cyclopentene	Various mutagenesis methods

Houde et al. Appl. Biochem. Biotechnol. 118, 155 (2004)

