

Application of directed enzyme evolution in synthesis

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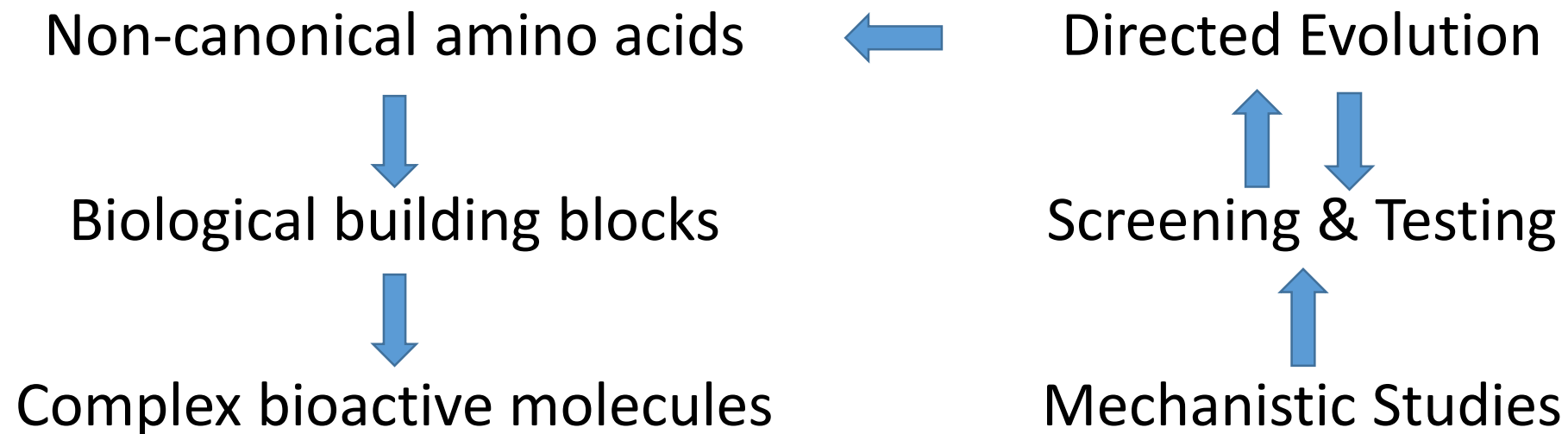
University of California, Santa Barbara

Agenda

- Background and Significance
- Directed Evolution
- **Synthesis Application-Tryptophan**
 - Tryptophan Analogs
 - Natural Products
 - Directed evolution of TrpB

“Enzymes are used in biocatalytic processes for the efficient and sustainable production of pharmaceuticals, fragrances, fine chemicals, and other products. Most bioprocesses exploit chemistry found in nature, but we are now entering a realm of biocatalysis that goes well beyond. Enzymes have been engineered to catalyze reactions previously only accessible with synthetic catalysts. Because they can be tuned by directed evolution, many of these new biocatalysts have been shown to perform abiological reactions with high activity and selectivity.”

Background



Directed Evolution

- protein engineering technique
 - mutagenesis
 - mimics natural selection
 - specific goals
-

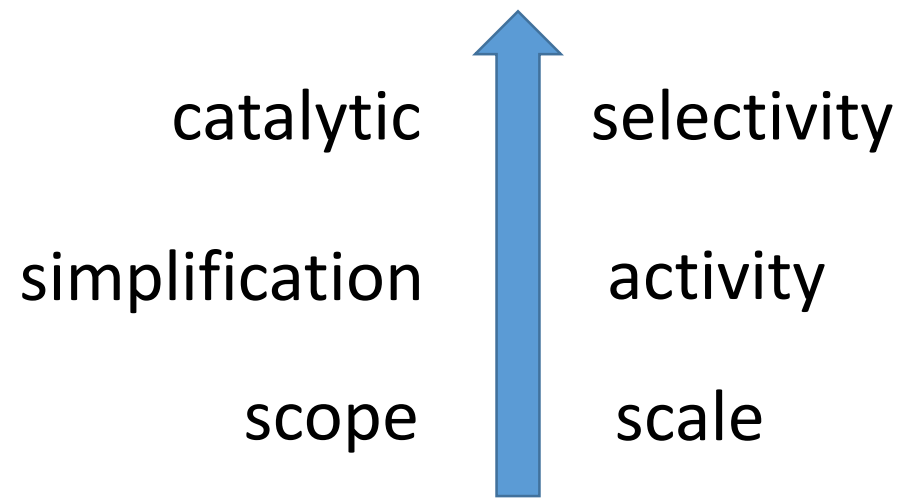
- KIE
- isotope trajectory
- R_{turnover}
- site-specific mutagenesis
- surrogate study

- computational design
- laboratory probe
- laboratory evolution



Significance

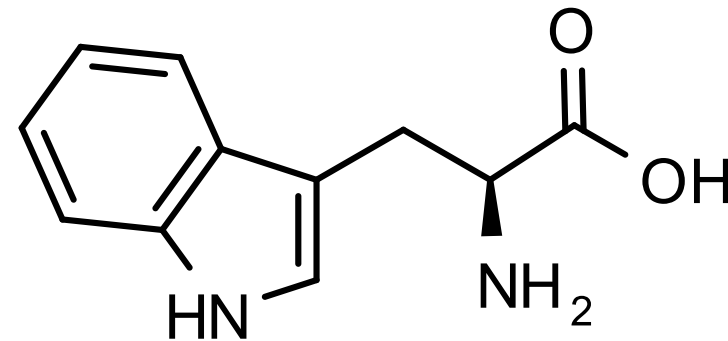
Improvement of current chemical synthesis



Abiological reactions using engineered enzymes

Target – Tryptophan Analogues

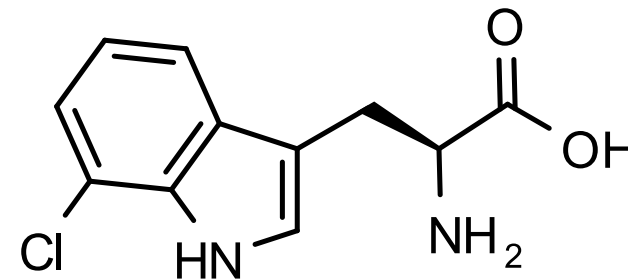
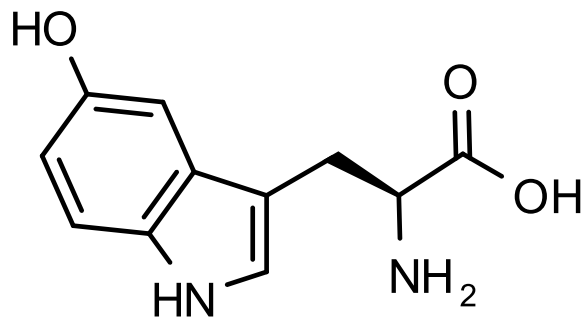
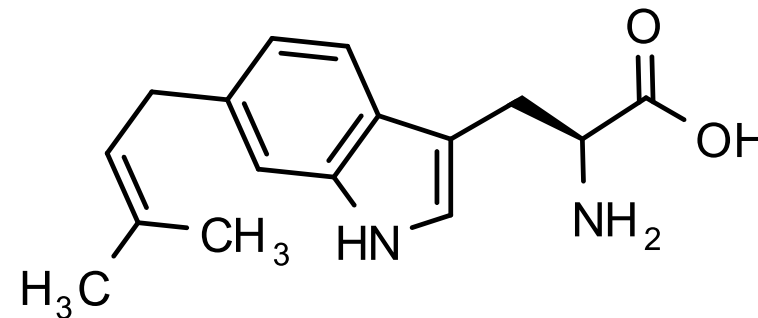
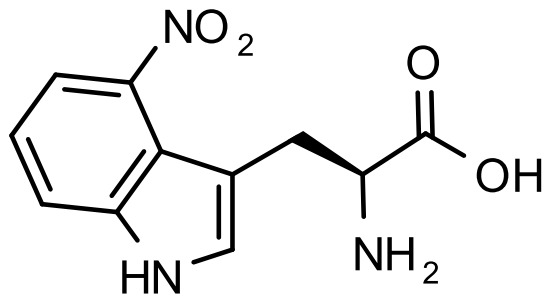
- biosynthetic precursors for a wide range of bioactivities
 - anticancer
 - antibiotic
 - antifungal
 - immunosuppressant
 - phytotoxic
 - chemical synthesis



L-Tryptophan

Current Limitations

- natural Trp derivatives: each requires a different enzyme



Current Limitations Cont.

- acylases and transaminases: only useful in setting the final stereoselectivity
- esterases: kinetic resolution (50% yield max)
- tryptophan synthase (TrpS): sensitive to the electronics and sterics of the substrates
 - complex: α -subunit (TrpA) + β -subunit (TrpB)
 - TrpA is a necessary allosteric actuator
 - TrpB is directly involved in synthesis of tryptophan analogues

Ley, S. V.; Priour, A.; Heusser, C. *Org. Lett.* 2002, 4, 711–714.

Blaser, G.; Sanderson, J. M.; Batsanov, A. S.; Howard, J. *Tetrahedron Lett.* 2008, 49, 2795–2798.

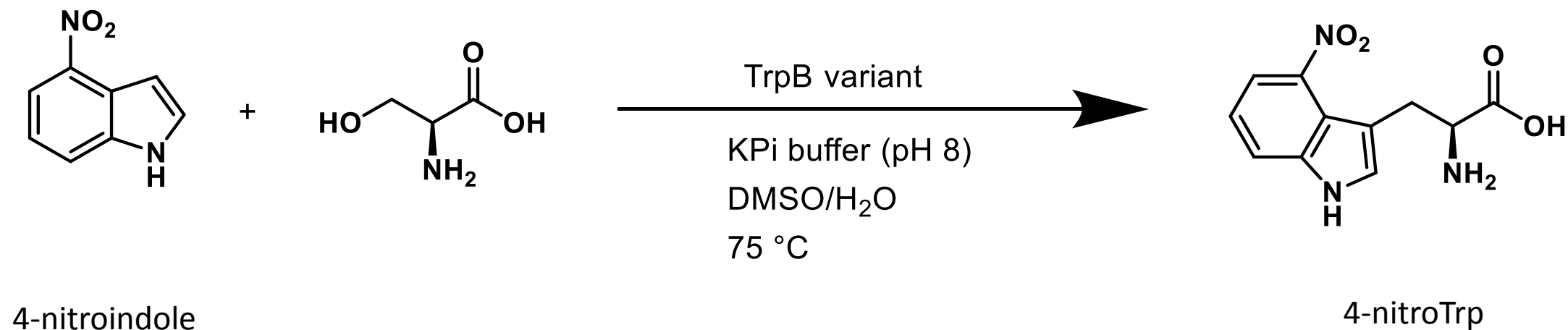
Baldwin, J. E.; Dyer, R. L.; Ng, S. C.; Pratt, A. J.; Russell, M. A. *Tetrahedron Lett.* 1987, 28, 3745–3746.

Starting Point

- recent development of TrpB variants: full activity without TrpA
- substrate: 4- substituted indoles
 - common in natural products
 - low reactivity with natural TrpB
 - 4-nitroindole

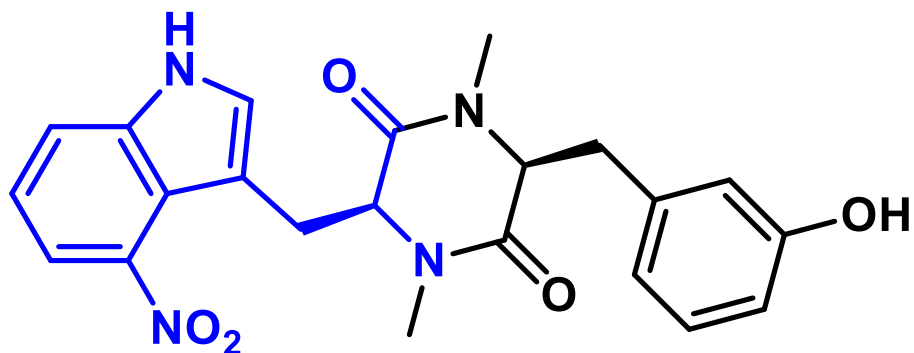
Starting Point – 4-nitroindole

- Steric hindrance
- Electron withdrawing (deactivating)

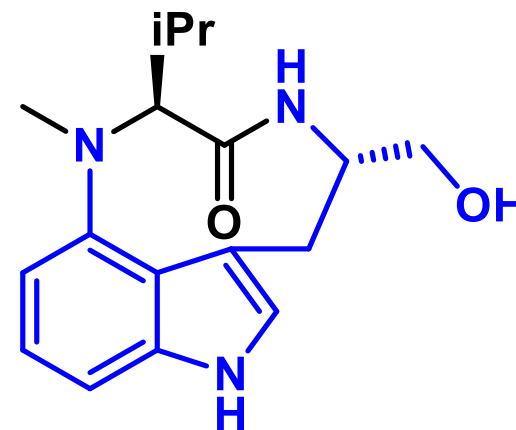


Starting Point – 4-nitroTrp

- 4-nitrotryptophan (4-nitroTrp) as a chemical precursor
 - complex enatiopure synthesis
 - limited scale using the natural enzyme



thaxtomin A



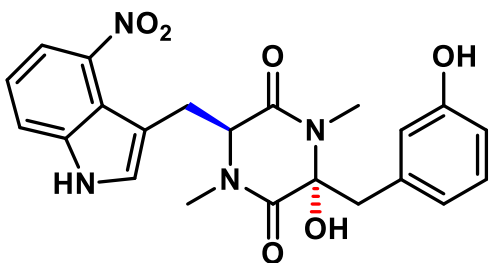
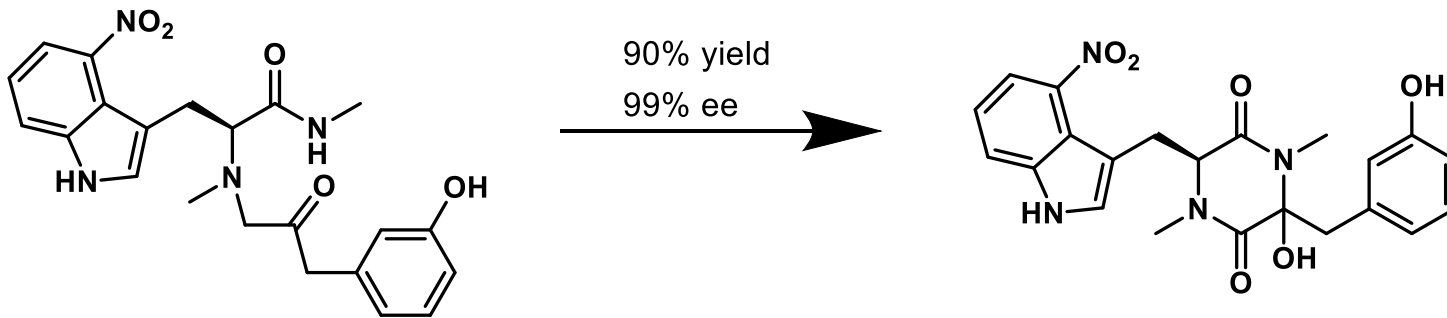
indolactam V

Zhang, H.; Ning, X.; Hang, H.; Ru, X.; Li, H.; Li, Y.; Wang, L.; Zhang, X.; Yu, S.; Qiao, Y.; Wang, X.; Wang, P. G. *Org. Lett.* 2013, 15, 5670–5673.

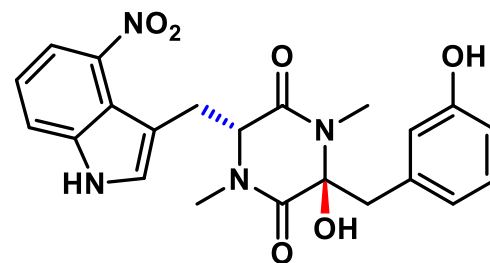
Barry, S. M.; Kers, J. A.; Johnson, E. G.; Song, L.; Aston, P. R.; Patel, B.; Krasnoff, S. B.; Crane, B. R.; Gibson, D. M.; Loria, R.; Challis, G. L. *Nat. Chem. Biol.* 2012, 8, 814–816.

Xu, Z.; Zhang, F.; Zhang, L.; Jia, Y. *Org. Biomol. Chem.* 2011, 9, 2512–2517.

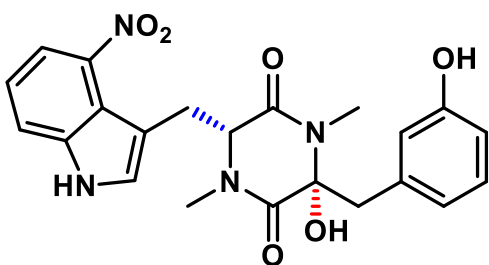
Synthesis of Thaxtomin A



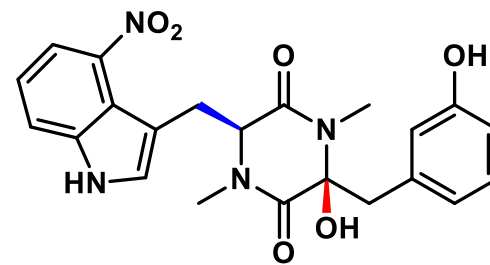
Thaxtomin A
herbicide



mirror-Thaxtomin A
fungicide



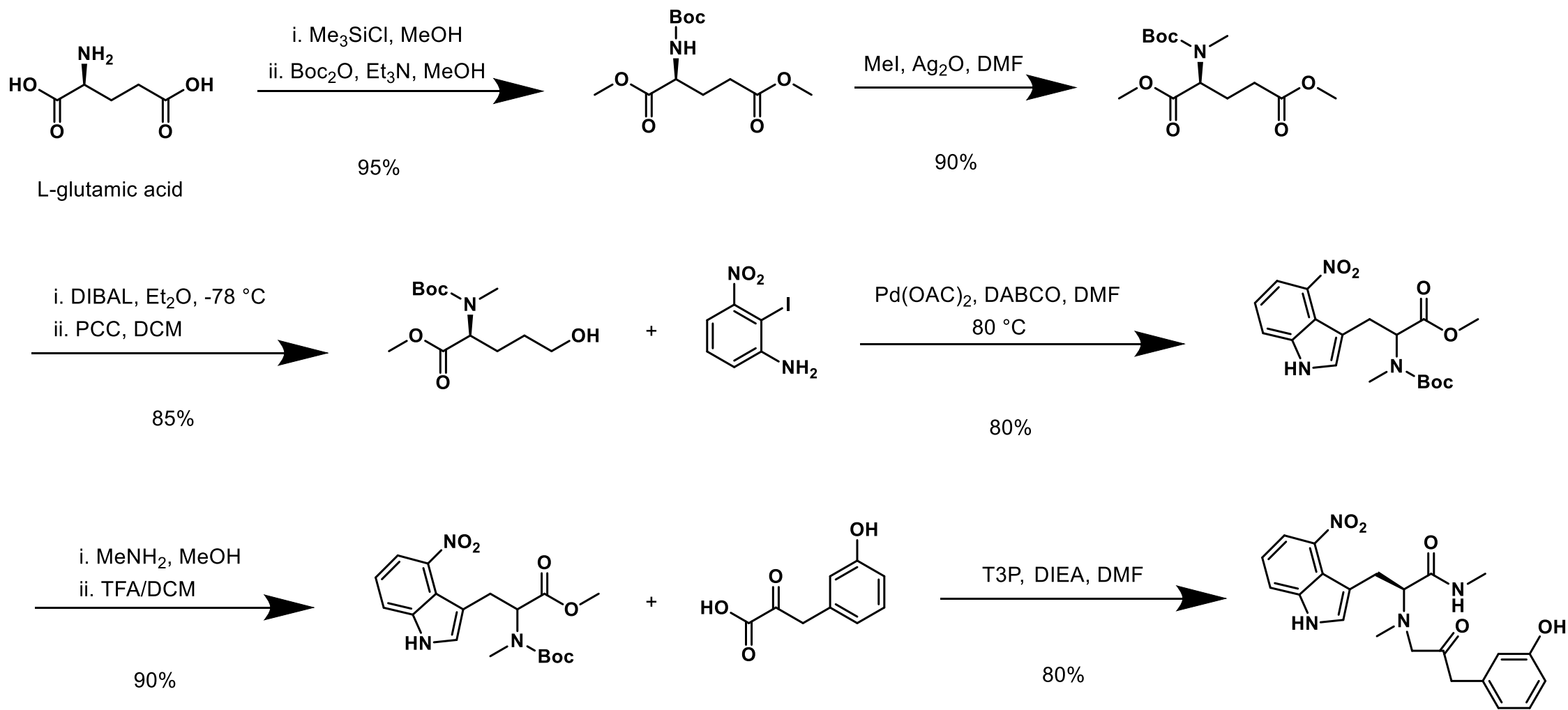
mirror-iso-Thaxtomin A
antiviral



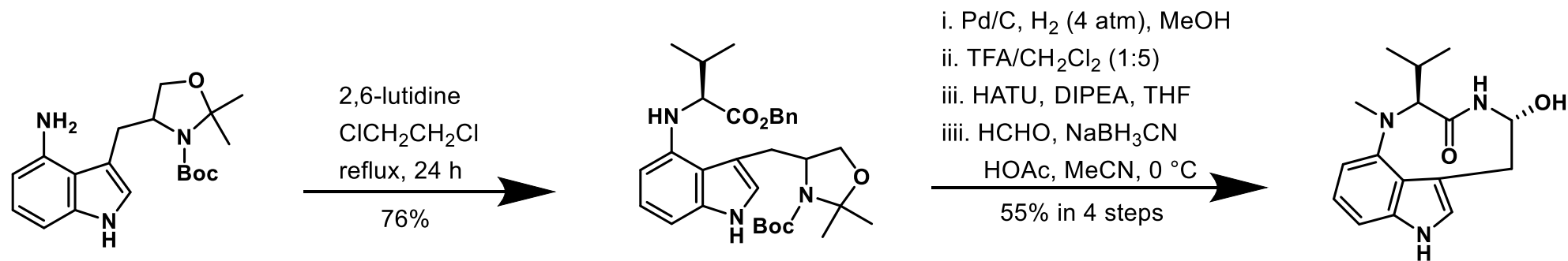
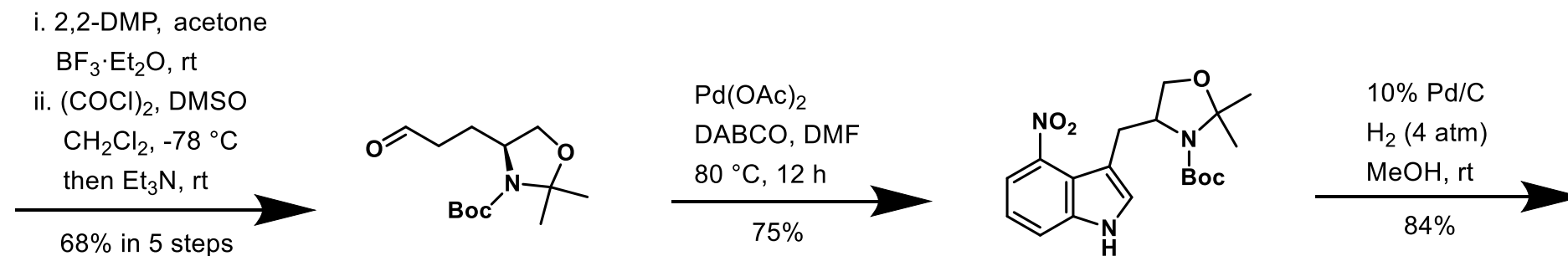
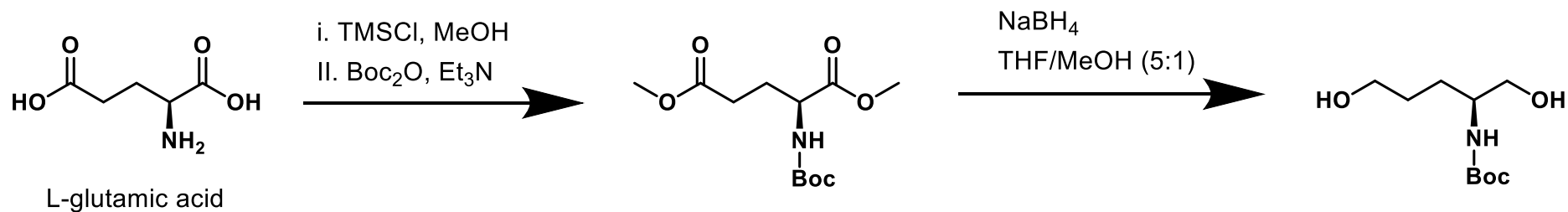
iso-Thaxtomin A
antiviral

Synthesis of Thaxtomin A

Cont.



Synthesis of Indolactam V



Screening

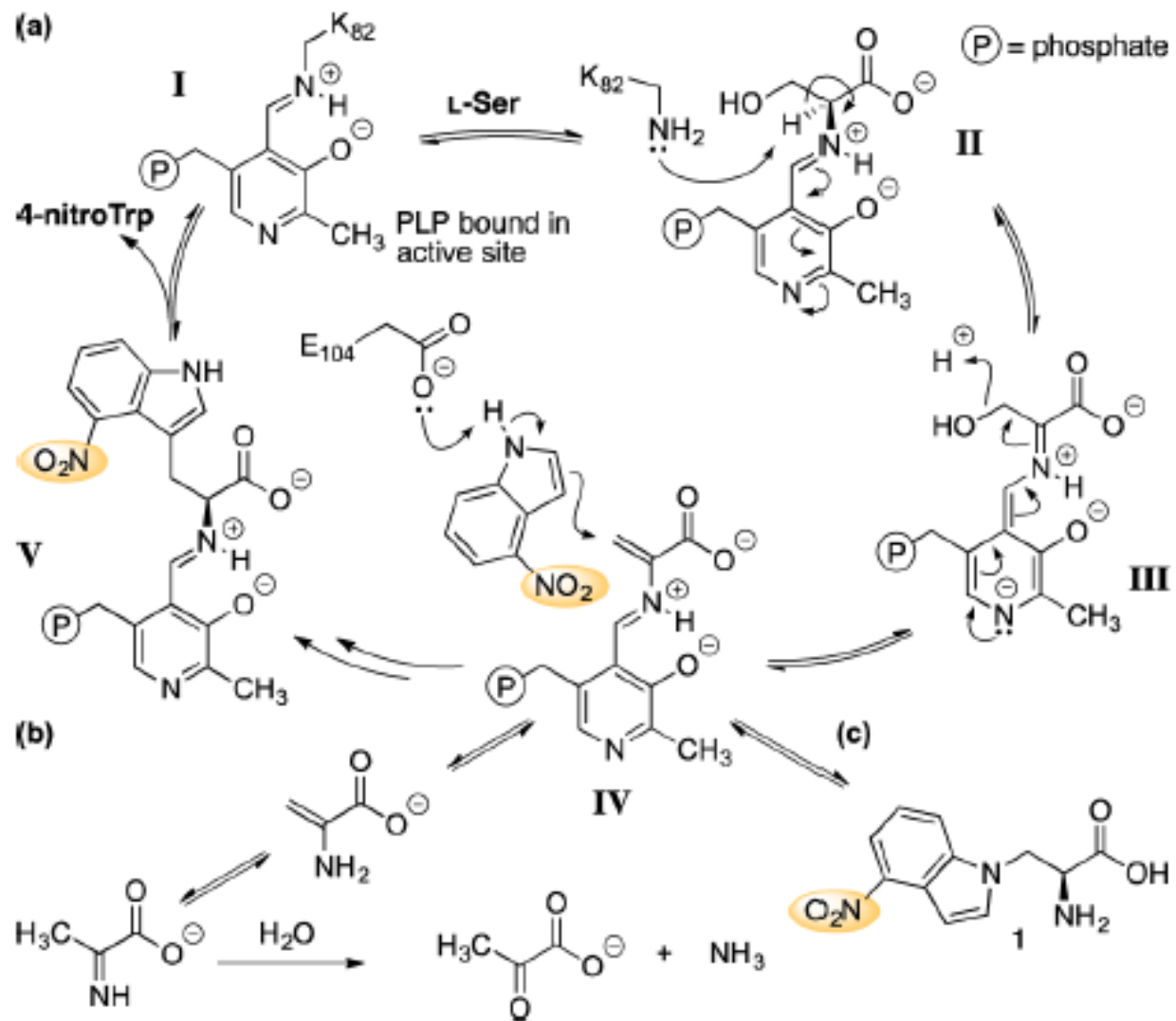


Figure 2. Putative pathways for reaction with 4-nitroindole. (a) Catalytic cycle for formation of 4-nitroTrp. (b) Enzymatic decomposition of Ser. (c) Competitive formation of isotryptophan 1.

- PfTrpB & TmTrpB
 - trace activity
- variants of 4-nitroTrp
 - Pf2B9: 18% conversion
- RDS: binding Ser
- reversible formation of side products is undesired

Active Site Mutagenesis- Sterics

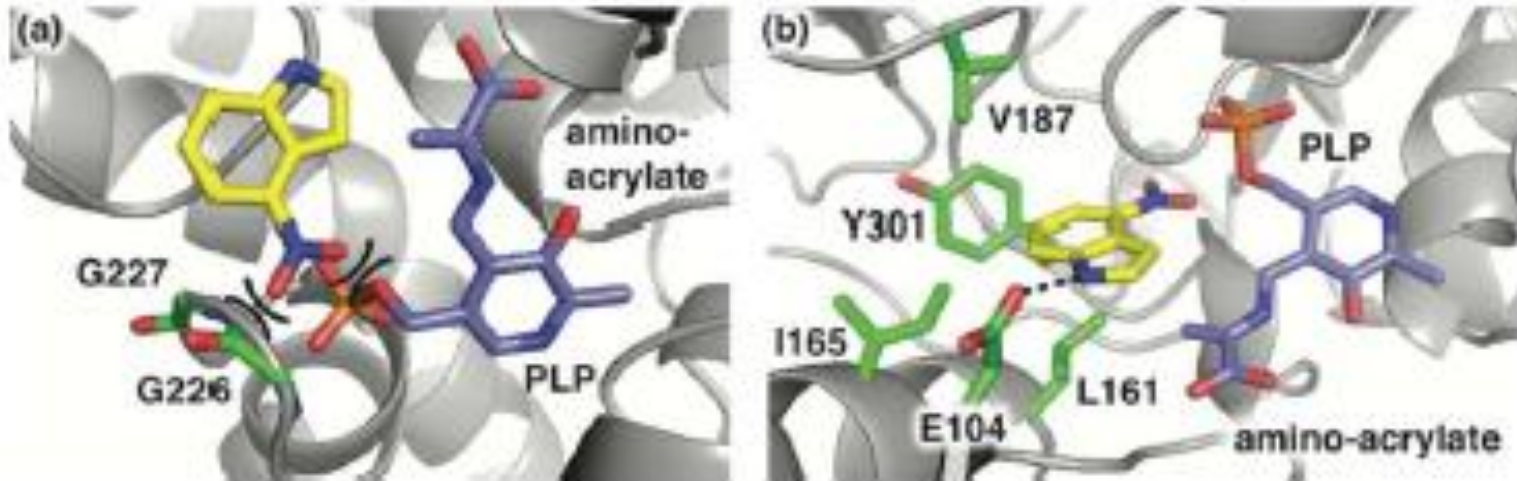


Figure 3. Model of 4-nitroindole (yellow) and the amino-acrylate in the active site of *PfTrpB* (see Section 8.13 of *Supporting Information*). (a) Nitro group clashes with the protein backbone (green) and the PLP cofactor (purple). (b) Alternative view showing side-chains extending in to the active site and hydrogen bond with E104.

- E104 – not modified
 - binds to indole through NH
 - Promotes the attack from C3
- L161V: 25%, no side product
- Not a steric problem

Random Mutagenesis

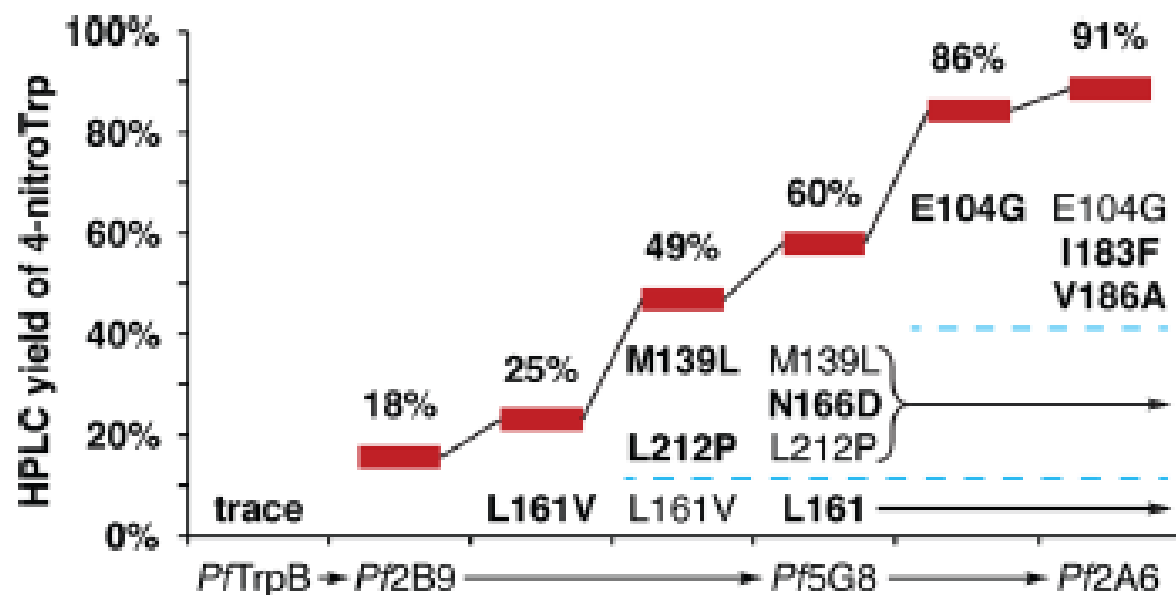


Figure 4. Evolutionary progression in production of 4-nitroTrp. Mutations in bold were added in the corresponding round of mutagenesis and screening. Dashed lines denote a new round of random mutagenesis. The horizontal axis indicates catalyst designations. See Section 8.7 of the Supporting Information for experimental details.

- Pf5G8 (leucine)
 - M139, L212, N166D: 60%, equimolar 4-nitroindole and Ser
- Pf2A6
 - E104G mutation: 86%
 - disapproves the supposition that H-bonding interaction with 4-nitroindole promotes the rxn
 - V186A, I183F & precedent: 91%
 - glycine is optimal (site-saturation screening)

Chart 2. Nitro Substitution at Other Positions

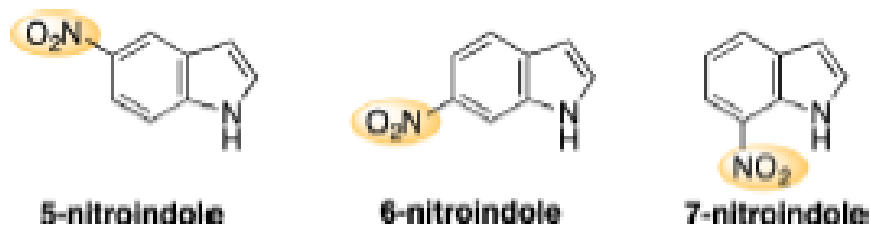


Table 1. Optimizing Catalysts for Other Nitroindole Isomers

entry	catalyst ^a	HPLC yield of nitroTrp (%) ^b			
		4-nitro	5-nitro	6-nitro	7-nitro
1	<i>Pf</i> 2B9	18	8	17	29
2	<i>Pf</i> 5G8	60	64	64	64
3	<i>Pf</i> 2A6	91	5	26	>99
4	<i>Tm</i> 2F3		76		
5	<i>Tm</i> 2F3 I184F		86		
6	<i>Pf</i> 2B9 I165F Y301H			66	
7	<i>Pf</i> 0A9			86	
8	<i>Pf</i> 0A9 E104G			91	

^aSee Table 2 for catalyst designations. ^bReactions used equimolar amounts of nitroindole and Ser. See Section 8.7 of the Supporting Information for experimental details.

- *Tm*2F3 I184F: 86% @5
 - Transferred mutations from *Pf*TrpB to the corresponding positions in *Tm*TrpB
- *Pf*0A9 E104G: 91% @6
 - 6-nitroindole has a different reactivity from 4-nitroindole

Scope of Catalyst Substrate

Table 3. Tryptophan Analogues Produced by Catalyst Panel^a

Entry	Substrate	R	Catalyst	Isolated yield (%)	Entry	Substrate	R	Catalyst	Isolated yield (%)
1		NO ₂	<i>Pf2A6</i>	95 ^b	12		NO ₂	<i>Pf0A9</i> E104G	91
2		F	<i>Tm2F3</i>	97	13		Cl	<i>Pf0A9</i>	98
3		Br	<i>Tm2F3</i>	72	14		Br	<i>Pf0A9</i>	97
4		CN	<i>Tm2F3</i> I184F	41 ^b	15		CN	<i>Pf0A9</i>	99
5		B(OH) ₂		ND	16		B(OH) ₂	<i>Pf0A9</i>	49
6		NO ₂	<i>Tm2F3</i> I184F	88 ^b	17		NO ₂	<i>Pf2A6</i>	98
7		CN	<i>Tm2F3</i>	79	18		CN	<i>Pf2A6</i>	98
8		CONH ₂	<i>Tm2F3</i>	77	19		Cl	<i>Pf0A9</i>	99
9		B(OH) ₂	<i>Pf0A9</i>	37	20		I	<i>Pf0A9</i>	91
10		I	<i>Pf0A9</i>	74 ^b	21		Br	<i>Pf5G8</i>	53
11		CF ₃	<i>Pf2A6</i>	19 ^{b,c}	22		B(OH) ₂		ND
					23		5,6-Cl ₂	<i>Pf5G8</i>	87
					24		5-Br-7-F	<i>Tm2F3</i> I184F	56
					25		5-Cl-7-I	<i>Tm2F3</i>	10

^aReactions used 0.02 mol % catalyst loading (maximum 5000 turnovers) and 1.1 equiv Ser relative to indole substrate. ^bCatalyst loading was 0.1 mol % (maximum 1000 turnovers). ^cReaction gives alkylation at nitrogen. ND, not detected.

Kinetics of the Mutations

Table 4. Initial Rates Throughout Evolution^a

entry	catalyst	initial turnover frequency (min ⁻¹)		
		to 4-nitroTrp	to pyruvate	to Trp
1	<i>Pf</i> TrpB	–	25.0 ± 0.3	19 ± 1.2
2	<i>Pf</i> 2B9	1.25 ± 0.07	12.2 ± 0.5	60.9 ± 0.16
3	<i>Pf</i> 5G8	1.8 ± 0.12	2.0 ± 0.2	9.9 ± 0.5
4	<i>Pf</i> 5G8 E104G	3.5 ± 0.2	0.9 ± 0.12	7.03 ± 0.07
5	<i>Pf</i> 2A6	7.0 ± 0.3	1.4 ± 0.10	17.6 ± 0.3

^aSee Sections 8.10, 8.11, and 8.12 of the Supporting Information.

The rate of Ser deamination: incubating the enzymes with Ser, excluding a Nu substrate and measuring the production of pyruvate.

- competing hydrolysis of amino-acrylate IM
- accelerating Nu attack of the substrate
 - Binding at the active site
 - Increasing the persistence of amino-acrylate

The New Catalyst

- M139 & N166 in open/close conformational states of TrpB
 - fully open: inactive/substrate entry and product release
 - fully closed: amino-acrylate formation, Nu addition/blocking access
 - stabilization of closed state = decreased Ser amination
 - H-bond between N166 and H275
 - rotameric switch for closure
 - basic aspartate to stabilize H-bond

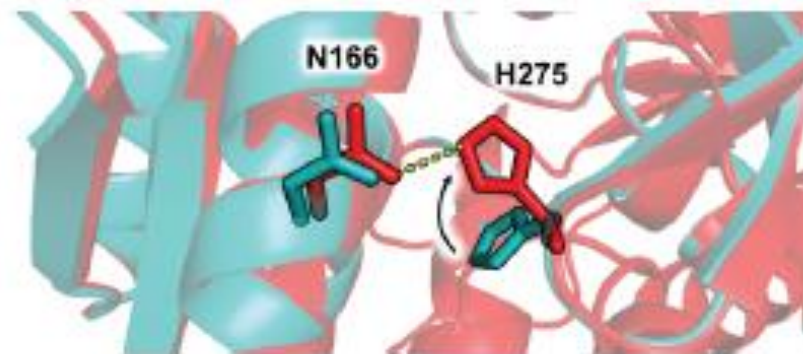


Figure 5. Overlaid crystal structures of *Pf*TrpB (PDB ID: 5DVZ) in the open state (cyan) and partially closed state (PDB ID: 5DW0, red) showing the side-chain motion of N166, and H275.

The New Catalyst Cont.

- Active site mutations
 - reshape to accommodate 4-nitroindole
 - bind in more reactive conformation
 - E104: modulates the transition to the closed state
 - ongoing study

The New Catalyst Cont.

- Directed evolution towards a challenging substrate unlocked the reactivity for many related substrates.
- The reaction was only limited by Ser hydrolysis, therefore, mutations to decrease Ser hydrolysis was the focus, and this focus was independent of substrates.

TrpB in Organic Synthesis

- challenge: Installing chiral amino acid moiety (α -C stereochemistry)
 - using existing amino acids: amine and carboxylate functional groups require protecting groups
- solution: TrpB
 - unprotected Ser
 - almost 100% enantioselective
 - straightforward synthesis
 - easy purification
 - precipitation
 - Amino-acrylate @ AS only
 - high expression level
 - thermostability

mediates 4-nitroindole

- Electronically deactivated
- Sterically hindered
- Poorly soluble

Expandable Scope

Will engineered biocatalysts open a door to
new syntheses?