Modelling of Product Release and Identification of Export Routes in Haloalkane Dehalogenase DhaA

Martin Klvaňa Martina Pavlová Petr Kulhánek Rebecca C. Wade Jiří Damborský

Loschmidt Laboratories, Faculty of Science, Masaryk University, Brno National Centre for Biomolecular Research, Faculty of Science, Masaryk University, Brno Molecular and Cellular Modeling Group, European Media Laboratory Research, Heidelberg

1,2,3-Trichloropropane (TCP)

Dangerous environmental pollutant

Propylene → epichlorohydrin (93 %) + TCP (7 %)

Biodegradation by haloalkane dehalogenase DhaA

Low activity for commercial use

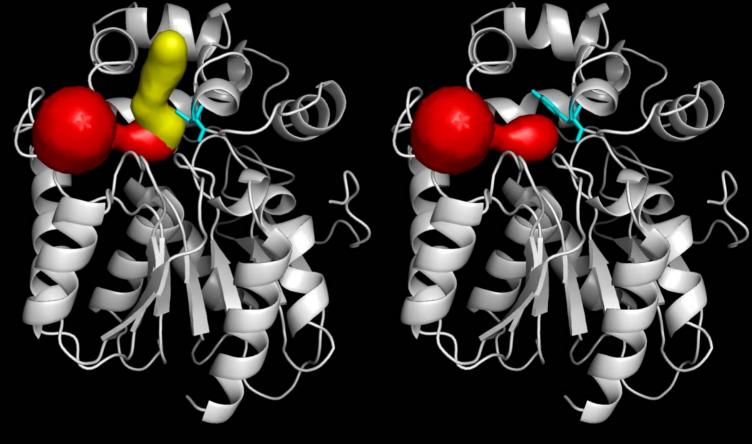
DhaA variant	Relative activity		
WT	1		
Bosma ¹	3.5		
Gray ²	4		

¹Bosma et al. (2002) *App.Environ.Microbiol.* 68, 3582-3587.

²Gray et al. (2001) Adv. Synth. Catal. 343, 607-616.

Why Export Routes?

- Main tunnel
- Slot

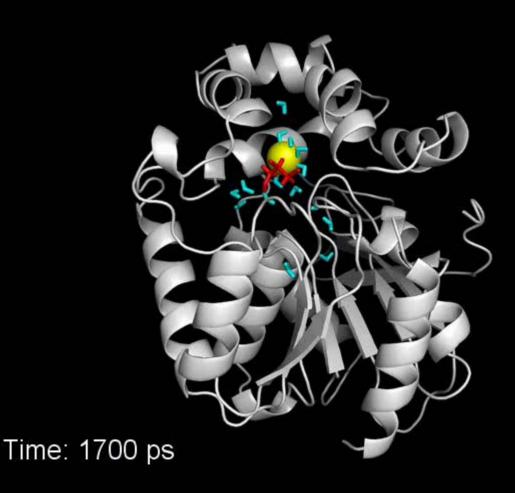


C176 (WT)

Y176 (Bosma, Gray)

Export of Cl⁻ through the Main Tunnel

Equilibration MD



Export of DCL through the Main Tunnel





Export of DCL through the Slot





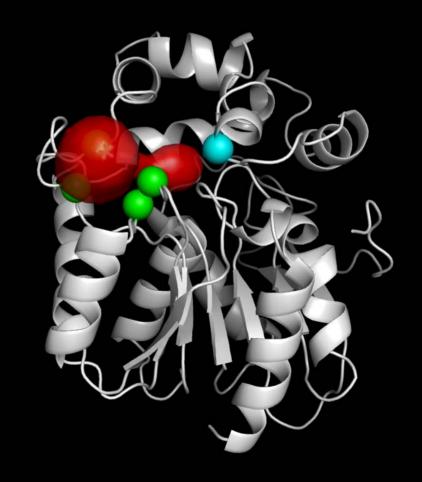
Re-design of Export Routes

Main tunnel:

C176Y

Slot:

- W141F
- I135X
- L245X
- V246X



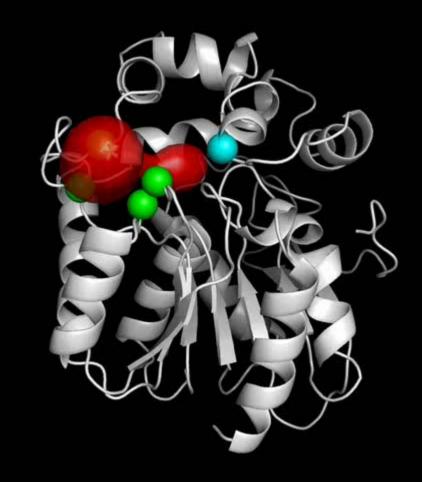
Re-design of Export Routes

Main tunnel:

C176Y

Slot:

- W141F
- I135X
- L245X
- V246X

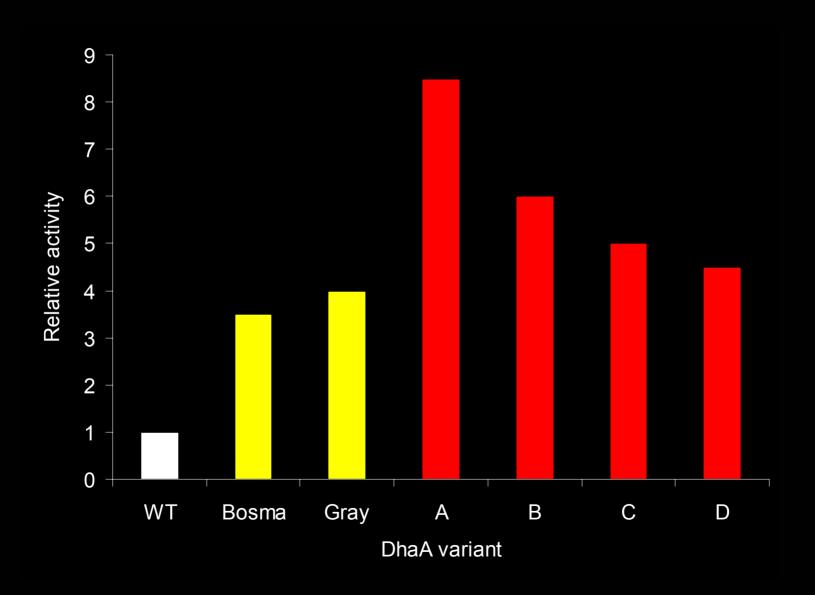


Mutagenesis

- Site-directed mutagenesis
- Directed evolution

Mutant	C176Y	W141F	I135X	V245X	L246X
А	√	✓	F	M	I
В	\checkmark	✓	Υ	M	1
С	\checkmark	✓	F	F	1
D	\checkmark	\checkmark	L	F	

Mutagenesis



Conclusions

RAMD provides useful information for protein design

Order of product release is 1. Cl⁻, 2. DCL

Export route for Cl is main tunnel

Export route for DCL is main tunnel and slot

Slot residues selected for mutagenesis are I135, W141, V245 and L246

The best mutant has 2-fold higher activity than mutant of Gray et al.