

Supporting Information

Supporting Tables

Table S1: Codon-optimized DNA sequence for expression of C-terminally His₁₀-tagged Ldi in *E. coli*

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ATGCCTTACCCGAAACCAGTCTATTGTCTCTGCTGCACTGCTGGCCGGCTCGGCCACCGCGTGCAGCG  
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GAAATTGCGTGCCTGGCTGCCACGATCTGGATATGCCGTGAGCAAGATGAAATGCAAGCGTGTGGCGACTGG  
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TATCAGCTGGTTACTGGCAGCCGCTTATGAGGCGGAGCACGCGCATCTGACCCGATCATCCATGATGAAATTGACCC  
AACCCGTTCCGGGCATCGTTGCGAGCCTGATAACTACTTGTCAATGTAATAGCGTCGACACCTGAGCCTGTGGTG  
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TCACCATCACTAATAA
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Table S2: Codon-optimized DNA sequence for expression of C-terminally His₁₀-tagged OmpA-Ldi in *E. coli*

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CAAAGCGCCGCGCAGCGAAACCGCCGGTACGGATGATGCAAGATGGTGGTGTGGCTGGCCAGCGCCTCACCTGCTGC  
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GTGCGCTGCTGCGTATGCCAACCGGCTGCCAAGCTGGCAGGCAAGCATACCATACCATCACCA  
AA
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Table S3: Codon-optimized DNA sequence for expression of N-terminally His₁₀-tagged nosig-Ldi in *E. coli*

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ATGCATCACCATCACCATCACCATCACCGGAAC TGCCGCCAGGCCGCTGGCCACCACCGAGGACTATTTGCG
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CAAGCTGGCAGGCAAGTAATAA

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Table S4: Codon-optimized DNA sequence for expression of C-terminally His₁₀-tagged nosig-Ldi in *E. coli*

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ATGGCGGAAC TGCCGCCAGGCCGCTGGCCACCACCGAGGACTATTTGCGCAGCAGGCAAACAGGCCGTTACGCCGGA
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TGCTAAGCCGTCCATTGTCAGCGCGAGCTTGCCTTATGAGCATCCGGGTTCGCTGCTGTTGATGAATTGCTGTTCTGGCG
AAGGTCCACGCCGGTTGGTGCCTGCTGCGTATGCCGCCACCGGCTGCCAGCTGGCAGGCAAGCATACCATCACCAT
CACCATCACCATCACTAATAA

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Table S5: Primers used for amplification of Ldi constructs for cloning into pET26b(+) vector. Underlined nucleotides are marking restriction sites

Primer	Sequence (5' → 3')
Fw(Ldi_NdeI)	TATA <u>CATATGCGCTTACCC</u> CTG
Rv(Ldi_His ₁₀ _HindIII)	GCC <u>AAGCTTATTAGTGATGGTGATGGTGATGGTGATGGT</u> GATGCTGCC TGCCAGCTTG
Fw(OmpA-Ldi_NdeI)	TATA <u>CATATGAAAAGACAGCTATC</u>
Fw(nosig-Ldi_NdeI)	TATA <u>CATATGGCGGA</u> ACTGCC
Fw(His ₁₀ -nosig-Ldi_NdeI)	GCC <u>CCATATGCATCACCATCACCATCACCATCACGCGGAA</u> CTGC CGCCAG
Rv(His ₁₀ -nosig-Ldi_HindIII)	GCC <u>AAGCTTATTACTTGCTGCCAGCTTGGC</u>

Table S6: Primers used for site-directed mutagenesis of Ldi cysteine residues

Primer	Sequence (5' → 3')
Fw(Ldi_C48A)	CGTTTACAGCCGTGG <u>TGCTAGCTTC</u> GAGGCATGGGAAC
Rv(Ldi_C48A)	GTTCCC <u>ATGCCTCGAAGCTAGCACCACGGCTGTAAACG</u>
Fw(Ldi_C48S)	CGTTTACAGCCGTGG <u>TAGCAGCTTC</u> GAGGCATGGGAAC
Rv(Ldi_C48S)	GTTCCC <u>ATGCCTCGAAGCTGCTACCACGGCTGTAAACG</u>
Fw(Ldi_C101A)	CGTGAGCAAGATGAA <u>AGCTAACCGTGTGGGGCG</u>
Rv(Ldi_C101A)	CGCCCCACACACG <u>CTTAGCTTCATCTTGCTCACG</u>
Fw(Ldi_C101S)	CGTGAGCAAGATGAA <u>AGCAAGCGTGTGGGGCG</u>
Rv(Ldi_C101S)	CGCCCCACACACG <u>CTTGCTTTCATCTTGCTCACG</u>
Fw(Ldi_C196A)	GTTCGCGGG <u>CATCGTTAGCGAGCCTGATAACTACT</u>
Rv(Ldi_C196A)	AGTAGTTATCAGG <u>CTCAGCAACGATGCCCGCGAAC</u>
Fw(Ldi_C196S)	GTTCGCGGG <u>CATCGTTAGCGAGCCTGATAACTACT</u>
Rv(Ldi_C196S)	AGTAGTTATCAGG <u>CTCGCTAACGATGCCCGCGAAC</u>
Fw(Ldi_C205A)	GATAACTACTTGT <u>CAAGCTAACGATAGCGTCGCATAACCTG</u>
Rv(Ldi_C205A)	CAGGTAT <u>GCAGCCTATTAGCTTAACAAAGTAGTTATC</u>

Fw(Ldi_C205S)	GATAACTACTTGTCAA <u>AGCA</u> ATAGCGTCGCATACTG
Rv(Ldi_C205S)	CAGGTATGCGACGCTATT <u>GCTT</u> GAACAAAGTAGTTATC

Table S7: *E. coli* strains tested for expression of Ldi

Strain	Notable properties
<i>E. coli</i> BL21-CodonPlus(DE3)-RIL	Extra copies of tRNAs genes (arginine, isoleucine, leucine)
<i>E. coli</i> BL21 TaKaRa 5	Co-expression of <i>E. coli</i> Trigger Factor (TF) chaperone
<i>E. coli</i> BL21 TaKaRa 1	Co-expression of chaperones dnaK-dnaJ-grpE-groES-groEL
<i>E. coli</i> Rosetta TM (DE3)pLysS	Presence of rare tRNA genes on the same plasmid that carry the T7 lysozyme gene
<i>E. coli</i> BL21-CodonPlus(DE3)-RP	Extra copies of the tRNA genes recognizing arginine (AGA, AGG) and proline (CCC).
<i>E. coli</i> K12 ER2508	Knock out of major ATP-dependent proteases in <i>E. coli</i> cytosol (Δlon , Δzjc)
<i>E. coli</i> TUNER TM	Lac permease (lacZY) deletion mutant. Allows for uniform entry of IPTG into all cells in the population.
<i>E. coli</i> ArcticExpress	Co-expression of cold-adapted chaperonins Cpn10 and Cpn60 from psychrophilic bacterium <i>Oleispira antartica</i>
<i>E. coli</i> LEMO23(DE3)	Precisely tunable expression by modulation of T7 lysozyme level by expression from well titrable rhamnose promoter

Table S8: β -myrcene hydration activity of Ldi cysteine variants in whole cell biotransformations with *E. coli*. ‘i.a.’: Inactive

Ldi variant	(S)-(+)-linalool production from β -myrcene / ng μL^{-1}
WT	1.03E+4
C48A	ia
C48S	ia
C101A	ia
C101S	ia
C170A	ia
C170S	ia
C179A	ia
C179S	ia

Supporting Figures

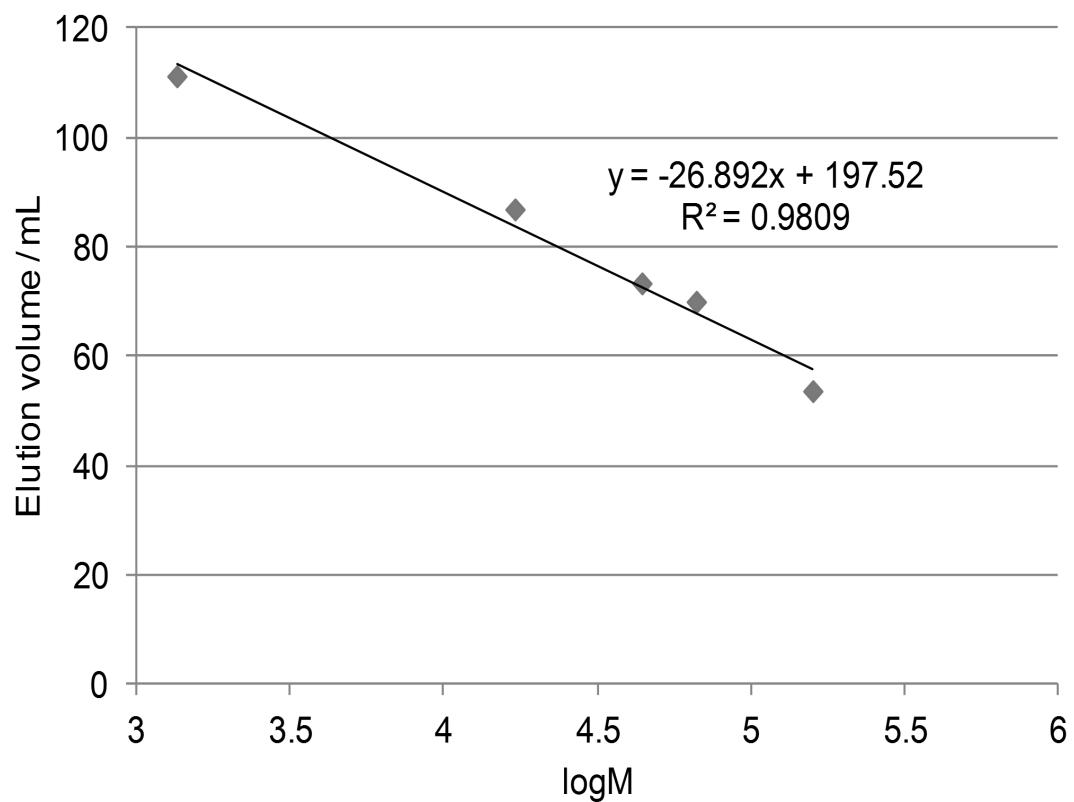
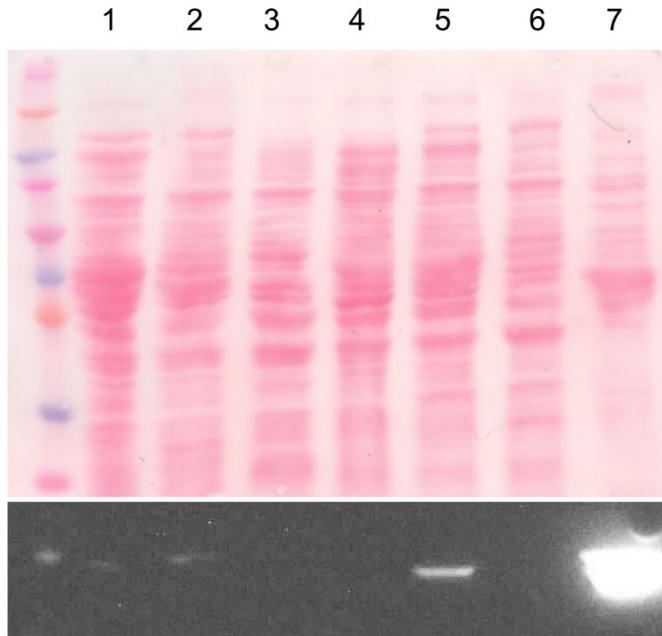


Figure S1: Gel filtration calibration curve generated with standard proteins listed in the Materials and Methods section.



Strain	Plasmid
1 <i>E. coli</i> LEMO21(DE3)	pMS470_Ldi
2 <i>E. coli</i> LEMO21(DE3)	pMS470 DsbC_Ldi
3 <i>E. coli</i> Origami	pMS470_Ldi
4 <i>E. coli</i> Origami	pMS470 DsbC_Ldi
5 <i>E. coli</i> BL21star(DE3)	pMS470_Ldi
6 <i>E. coli</i> BL21star(DE3)	pMS470 DsbC_Ldi
7 <i>E. coli</i> BL21star(DE3), insoluble fraction	pMS470_Ldi

Figure S2: Initial expression studies of Ldi in different *E. coli* strains indicated poor expression of soluble protein. Ldi was expressed from a pMS470 vector with and without co-expression of truncated DsbC. The level of Ldi in *E. coli* lysate was analyzed by immunoblotting.

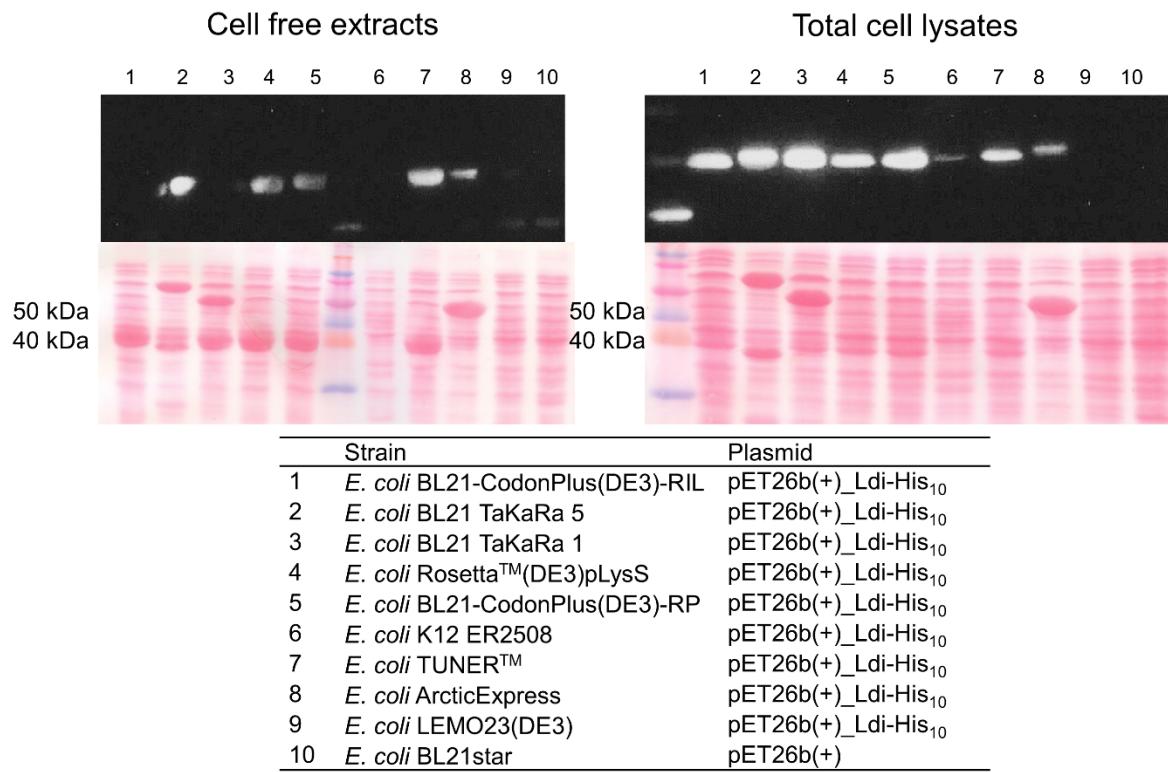


Figure S3: Expression of Ldi in different *E. coli* strains increases the yield of soluble protein in specific set ups. Nine different *E. coli* strains were tested for expression of Ldi, and after harvesting the cells, His₁₀-tagged Ldi was detected in total cell lysates and cell free extracts by immunoblotting.

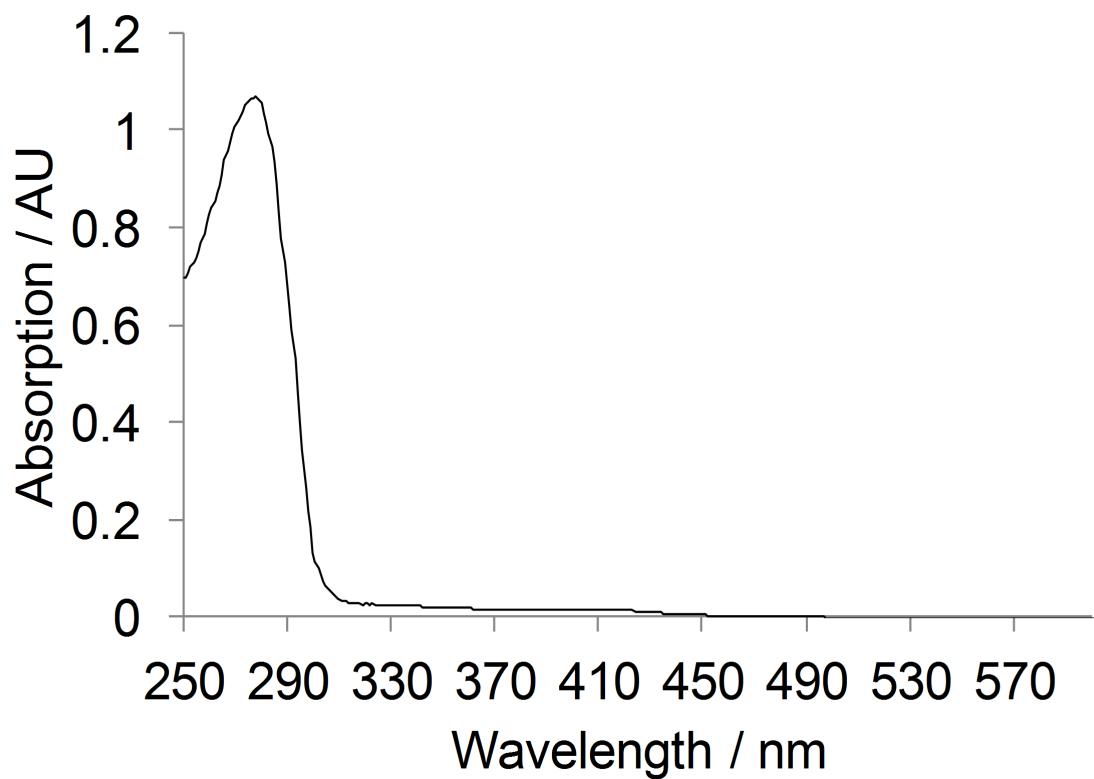


Figure S4: UV-Vis absorption spectrum of purified Ldi. The enzyme was His₁₀-tag purified from cell free extract after expression in *E. coli* BL21-CodonPlus(DE3)-RP. The lack of any noticeable absorbance in the range between 300 and 800 nm indicates that Ldi was purified as a cofactor-free enzyme.

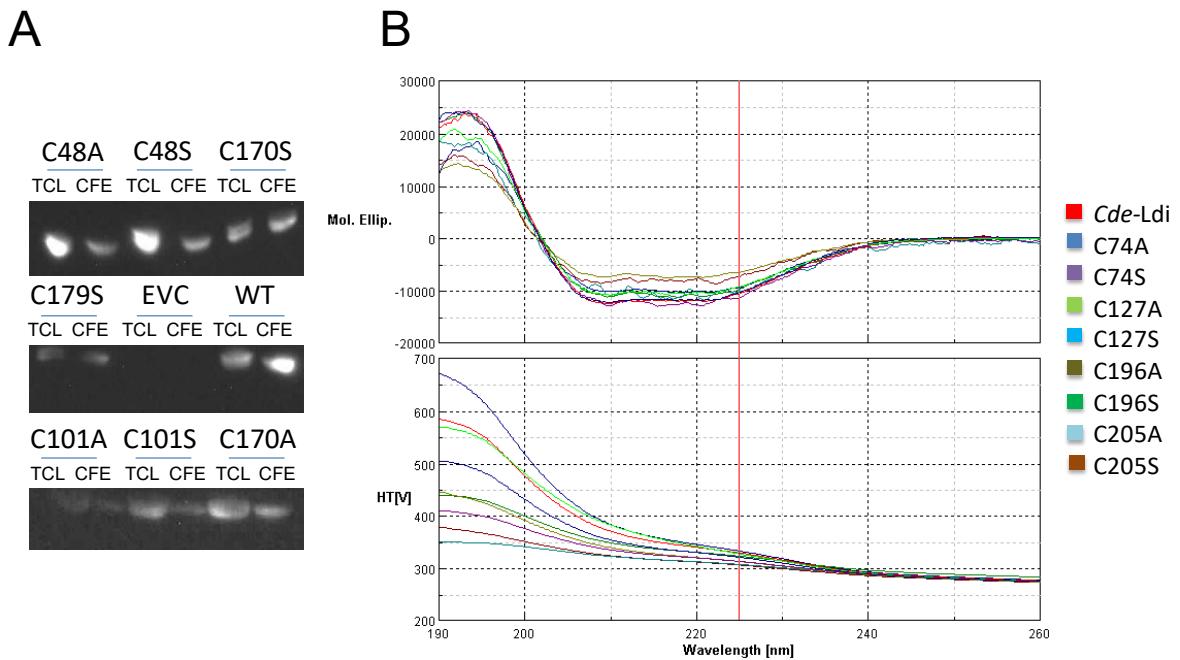


Figure S5: Expression and overall structure of Ldi cysteine variants. (A) Expression of cysteine variants and Ldi wild type (WT) in total cell lysates (TCL) and cell free extracts (CFE) was confirmed by immunoblotting. EVC: Empty vector control. (B) Comparison of circular dichroism (CD) spectra of Ldi wild type enzyme and variants confirms the overall integrity of single amino acid exchange variants. Ldi wild type enzyme and variants were purified in parallel. After dilution to an A_{280} of approx. 0.8, far-UV CD spectra were collected between 190 and 260 nm. Data are expressed in mean residue ellipticity $[\Theta]$ in $\text{deg cm}^2 \text{dmol}^{-1}$.

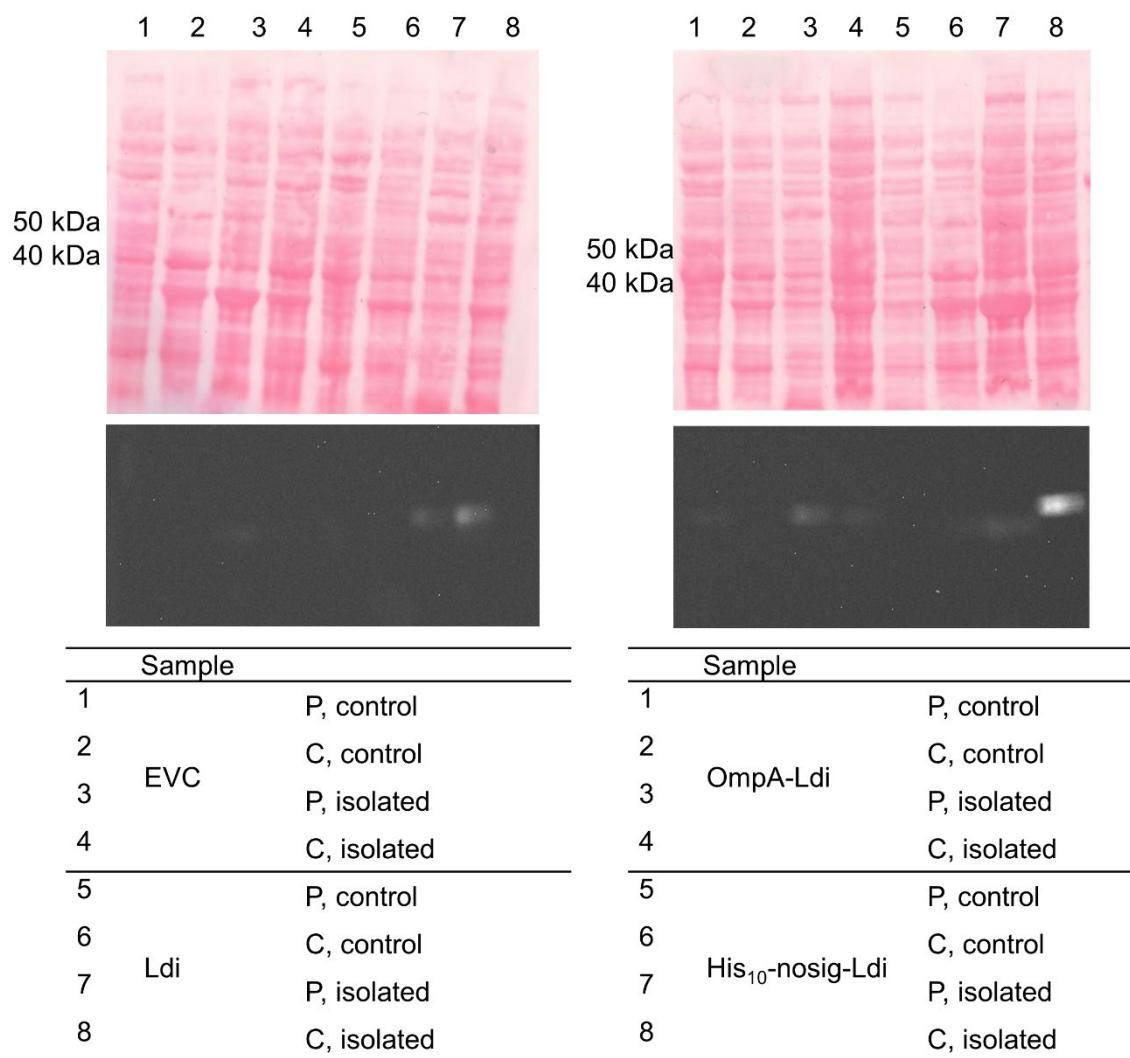


Figure S6: An N-terminal signal sequence favors localization of Ldi in the *E. coli* periplasm. After expression, *E. coli* periplasm (P) and cytosol (C) were separated as described in the Materials and Methods section. Recombinant Ldi in isolated fractions was detected via immunoblotting. Controls were treated with ddH₂O instead of the solutions inducing formation of spheroplasts.