

# Deciphering the Crucial Roles of the Quorum-Sensing Transcription Factor SdiA in NADPH Metabolism and (S)-Equol Production in *Escherichia coli* Nissle 1917

Zhe Wang <sup>1,2</sup>, Yiqiang Dai <sup>1,2</sup>, Fidelis Azi <sup>3</sup>, Mingsheng Dong <sup>1,\*</sup> and Xiudong Xia <sup>1,2,4,5,\*</sup>

<sup>1</sup> College of Food Science and Technology, Nanjing Agricultural University, Nanjing 210095, China; wangzhe@stu.njau.edu.cn (Z.W.); 2021208005@stu.njau.edu.cn (Y.D.)

<sup>2</sup> Institute of Agro-Product Processing, Jiangsu Academy of Agricultural Sciences, Nanjing 210014, China

<sup>3</sup> Department of Chemical Engineering, Guangdong Technion-Israel Institute of Technology, Shantou 515063, China; fidelis.azi@gtit.edu.cn

<sup>4</sup> Jiangsu Key Laboratory for Food Quality and Safety-State Key Laboratory Cultivation Base, Ministry of Science and Technology, Nanjing 210014, China

<sup>5</sup> School of Food and Biological Engineering, Jiangsu University, Zhenjiang 212013, China

\* Correspondence: dongms@njau.edu.cn (M.D.); 20140034@jaas.ac.cn (X.X.)

## Supplementary Tables

**Table S1.** Nucleotide sequences of primers. Primer sequence utilized for homologous recombination is underlined.

Oligonucleotides	Sequences, 5'-3'
Pf_Pnar	GAAGGAGATATACATATGGCAGATCTCAATTGG
Pr_Pnar	ATGTATATCTCCTTCTTAAAGTTAAACAAATCTTTAAGGGCATTATACCG C
Pf_malEK (up)	GCGGTCAGCATAATCATTACCC
Pr_malEK (up)	GGC <u>GTCGACCCTAGGG</u> GAGACTGCTGCCGAAAGAGTCT
Pf_malEK (down)	<u>CCTAGGGTCGAC</u> GCCCCCTGCTGTTCAAAACGTTTTG
Pr_malEK (down)	TCCGGTTACGGTAGGCAAC
Pf_exo/cea (up)	GCTCCCATATCCCAGAACTG
Pr_exo/cea (up)	<u>GTCGACCCTAGGG</u> GCGGTCAGATTGAGTTCACCG
Pf_exo/cea (down)	ACCC <u>CCTAGGGTCGAC</u> CGGCATGGTCCCGGAAAACGGTA
Pr_exo/cea (down)	CCGCAATCATTTACGTTATCC
Pf_bglF	ATGGAAACGGAGTTAGCCAGAAAAAT
Pr_bglF	TTAGCGAATGATGGATAACAGCG
Pf_bglB	ATGAAAGCATTTCCAGAAACATTCTT
Pr_bglB	TTAAGGTGCTTTAATGGTTATTTTTTTTAATGACAG
Pf_ptsG (up)	CGTTATGTCCCCCTGGATC
Pr_ptsG (up)	CCTGAGTATGGGTGCTTTTT

---

Pf_ptsG (down)	GAGTATGGGTGCTTTTTTGGCAGAAGCAGGCGGT
Pr_ptsG (down)	CTACCGGGTTCTGGTAAGC
Pf_decR (up)	ATCCACCAGCGTCAGCAC
Pr_decR (up)	CTTCCAGCTTCGCGATGGA
Pf_decR (down)	TCGCGAAGCTGGAAGTCATGTCGGCAACCTGGAC
Pr_decR (down)	ATGTTAGATAAAATTGACCGTAAGC
Pf_HW372_01960 (up)	GGCTCCCCTGTGAAATCGT
Pr_HW372_01960 (up)	GGTTGATCGCGTCTTTAATATCG
Pf_HW372_01960 (down)	AAGACGCGATCAACCGTGATTTCAGGGGCAAGAT
Pr_HW372_01960 (down)	GCGGAAGTTGCGAGTAAAGC
Pf_yhjC (up)	TGACCGATTGTTGTTTACAACG
Pr_yhjC (up)	CTCCACCTGAATACGTAAAAAGAC
Pf_yhjC (down)	ACCCAGGTGCGGTGTATAAC
Pr_yhjC (down)	ACTGGCTAACAACAAGAAGTGTTTCGCGCCAGATAC
Pf_HW372_03545 (up)	CGCAAGCTGGCAGAACTT
Pr_HW372_03545 (up)	ATTTATCTCCGGTAGAGGTCGC
Pf_HW372_03545 (down)	CTACCGGAAGATAAATGCTACATTAATGAGCATCGTGAAG
Pr_HW372_03545 (down)	GCGCCATCATGATCAGAACATC
Pf_sdiA (up)	ACGTCGTTTGTCTGGCGG
Pr_sdiA (up)	ATAGTAAACCGCAACGCCCC
Pf_sdiA (down)	GTTGCGGTTTACTATGCAGCTGGAGTACGATTACTATTCG

---

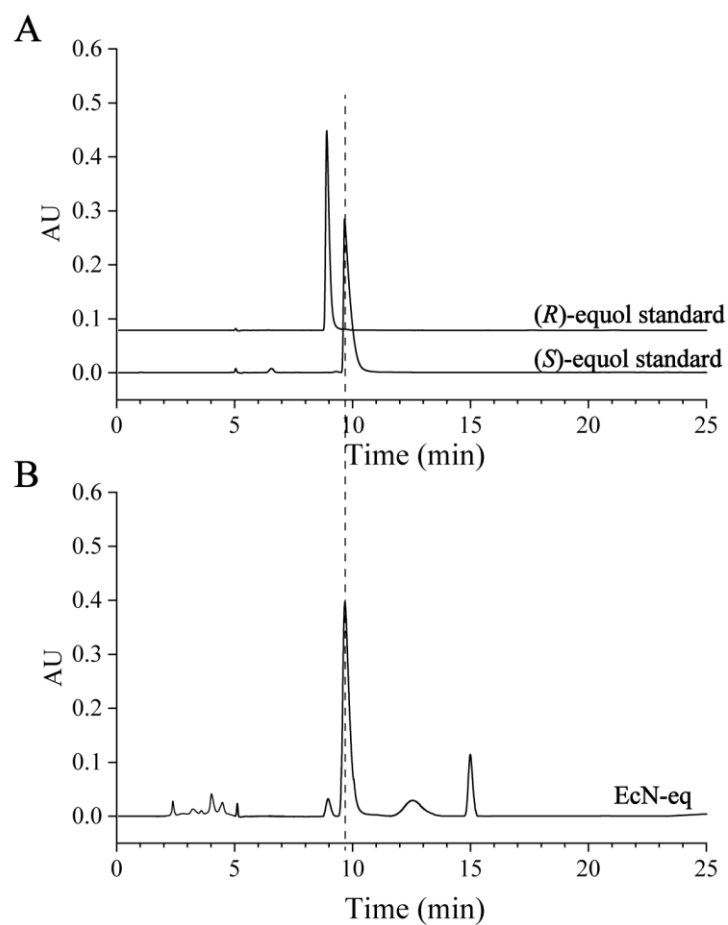
---

Pr_sdiA (down)	GCTGATGTCTTACCTTCCGCC
Pf_yhaJ (up)	ACAACATAATCAGGTCGCGTC
Pr_yhaJ (up)	CACATTCGTTTGCAAAGGAAGG
Pf_yhaJ (down)	TGAGAACGAAATGGCCTTCCGCACTTAGCTACACC
Pr_yhaJ (down)	CCACGATATTTACGCGCG
Pf_addsgRNA	GTTCCGTTTATCCGGGCAAACTAGTATTATACCTAGGACTGAGC
Pr_maleK-sgRNA	TCTGCTCGACAAACCCTTCGTTTTAGAGCTAGAAATAGCAAGTT
Pr_exo/cea-sgRNA	ACTCCATACCCTCCCCAACGTTTTAGAGCTAGAAATAGCAAGT
Pr_HW372_01960-sgRNA	TGTGTTTGGAGATGTTTCAGGTTTTAGAGCTAGAAATAGCAAGT
Pr_yhjC-sgRNA	GCAGTTGTTTCATCAAAGTCGTTTTAGAGCTAGAAATAGCAAGT
Pr_HW372_03545-sgRNA	GCGCGAGTCAGTTAACGCCTTTTTAGAGCTAGAAATAGCAAGT
Pr_sdiA-sgRNA	TTCATGGTAGACCTCTTCTGTTTTAGAGCTAGAAATAGCAAGT
Pr_yhaJ -sgRNA	ATGGATGCGATCGATCGCCGTTTTAGAGCTAGAAATAGCAAGT
Pr_ptsG-sgRNA	GTATCCGTA CTGCCTATCGAGTTTTAGAGCTAGAAATAGCAAGT
Pr_decR -sgRNA	GTCTGGTAAACAGTGTACCGTTTTAGAGCTAGAAATAGCAAGT
F_16S (qPCR)	GTTAAGTCCCGCAACGAGCGCAA
R_q16S (qPCR)	CTTTATGAGGTCCGCTTGCTCTC
F_zwf (qPCR)	CCAAGCTGGATCTGAGCTATTC
R_zwf (qPCR)	CCACTTCATCACGACGTACAA
F_gnd (qPCR)	GAACCGCTGTCGCTGATTA
R_gnd (qPCR)	GGGCCAGAGAGAACTTTAGATG

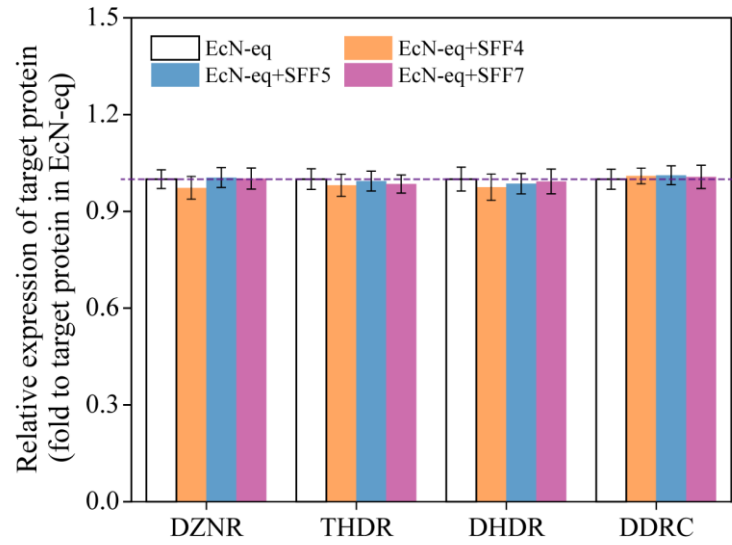
---

**Table S2.** Vectors used in this study.

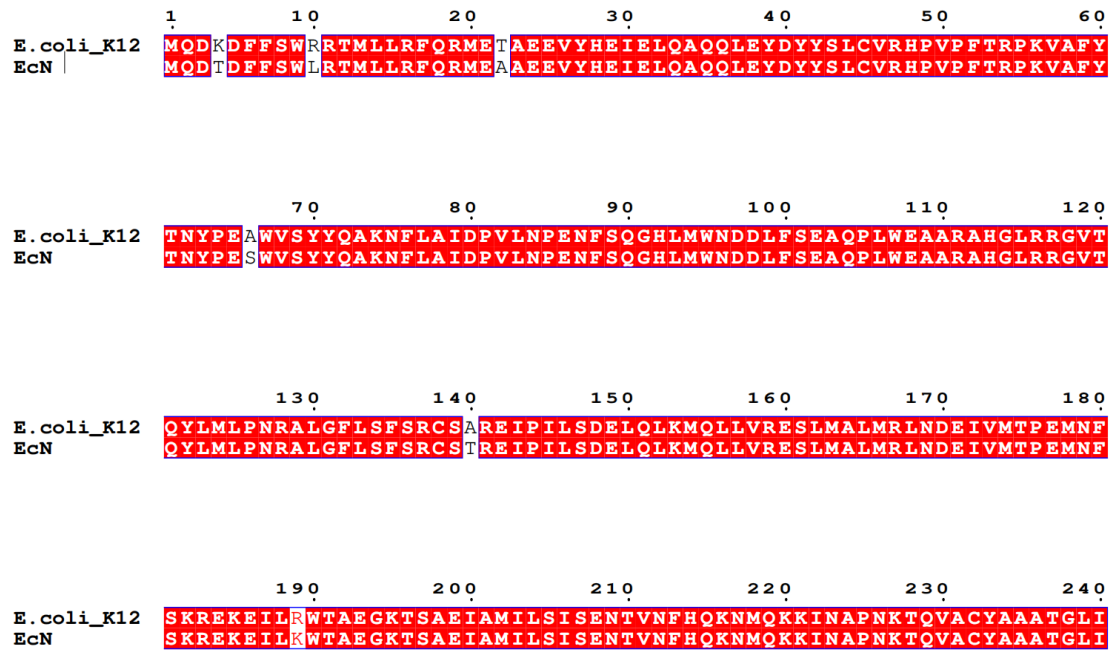
Vectors	Relevant properties	Reference
pETM6	pBR322 ori, AmpR, T7 promote	Addgene #49795
pEcCas	<i>Cas9</i> under <i>cas</i> promoter, gRNA- <i>pMB1</i> under <i>rha</i> promoter, $\lambda$ -red under <i>araB</i> promoter, pSC101 ori, <i>KanR</i>	Addgene #73227
pEcgRNA	sgRNA under J23119 promoter, pMB1 ori, <i>aadA</i>	Addgene#166581
pETM6- <i>P<sub>nar</sub></i>	pBR322 ori, AmpR, <i>nar</i> promote	This study
pUC57- <i>malEK</i>	<i>malEK</i> homology arms, pBR322 ori	This study.
pUC57- <i>exo/cea</i>	<i>exo/cea</i> homology arms, pBR322 ori	This study.
pUC57- <i>ptsG</i>	<i>ptsG</i> homology arms, pBR322 ori	This study.
pUC57- <i>yhaJ</i>	<i>yhaJ</i> homology arms, pBR322 ori	This study.
pUC57- <i>decR</i>	<i>decR</i> homology arms, pBR322 ori	This study
pUC57- <i>HW372_01960</i>	<i>HW372_01960</i> homology arms, pBR322 ori	This study.
pUC57- <i>yhjC</i>	<i>yhjC</i> homology arms, pBR322 ori	This study.
pUC57- <i>HW372_03545</i>	<i>HW372_03545</i> homology arms, pBR322 ori	This study.
pUC57- <i>sdiA</i>	<i>sdiA</i> homology arms, pBR322 ori	This study.
pUC57- <i>exo/cea-P<sub>nar</sub>-bglIF-P<sub>nar</sub>-bglB</i>	pUC57- <i>exo/cea</i> carrying - <i>P<sub>nar</sub>-bglIF-P<sub>nar</sub>-bglB</i>	This study.
pUC57- <i>malEK-P<sub>nar</sub>-dznr-P<sub>nar</sub>-ddrc-P<sub>nar</sub>-dhdr-P<sub>nar</sub>-thdr</i>	pUC57- <i>malEK</i> carrying <i>P<sub>nar</sub>-dznr-P<sub>nar</sub>-ddrc-P<sub>nar</sub>-dhdr-P<sub>nar</sub>-thdr</i>	This study.
pEcgRNA- <i>ptsG</i>	Derived from pEcgRNA, target <i>ptsG</i> in EcN	This study.
pEcgRNA- <i>malEK</i>	Derived from pEcgRNA, target <i>malEK</i> in EcN	This study.
pEcgRNA- <i>exo/cea</i>	Derived from pEcgRNA, target <i>exo/cea</i> in EcN	This study.
pEcgRNA- <i>yhaJ</i>	Derived from pEcgRNA, target <i>yhaJ</i> in EcN	This study.
pEcgRNA- <i>decR</i>	Derived from pEcgRNA, target <i>decR</i> in EcN	This study.
pEcgRNA- <i>HW372_01960</i>	Derived from pEcgRNA, target <i>HW372_01960</i> in EcN	This study.
pEcgRNA- <i>yhjC</i>	Derived from pEcgRNA, target <i>yhjC</i> in EcN	This study.
pEcgRNA- <i>HW372_03545</i>	Derived from pEcgRNA, target <i>HW372_03545</i> in EcN	This study.
pEcgRNA- <i>sdiA</i>	Derived from pEcgRNA, target <i>sdiA</i> in EcN	This study.
pETM6- <i>P<sub>nar</sub>-yhaJ</i>	pETM6- <i>P<sub>nar</sub></i> harboring <i>yhaJ</i>	This study.
pETM6- <i>P<sub>nar</sub>-decR</i>	pETM6- <i>P<sub>nar</sub></i> harboring <i>decR</i>	This study.
pETM6- <i>P<sub>nar</sub>-HW372_01960</i>	pETM6- <i>P<sub>nar</sub></i> harboring <i>HW372_01960</i>	This study.
pETM6- <i>P<sub>nar</sub>-yhjC</i>	pETM6- <i>P<sub>nar</sub></i> harboring <i>yhjC</i>	This study.
pETM6- <i>P<sub>nar</sub>-HW372_03545</i>	pETM6- <i>P<sub>nar</sub></i> harboring <i>HW372_03545</i>	This study.
pETM6- <i>P<sub>nar</sub>-sdiA</i>	pETM6- <i>P<sub>nar</sub></i> harboring <i>sdiA</i>	This study.



**Fig. S1.** Chiral HPLC analysis of daidzin metabolites converted by strain EcN-eq. (A). Reference standards of (*S*)-Equol and (*R*)-Equol. (B). HPLC spectrum displaying the reaction sample from strain EcN-eq.



**Fig. S2.** Densitometric semi-quantifications of the SDS-PAGE in Fig. 2C. Experiments in this study were conducted in triplicate, and error bars signify standard deviation (SD) with a 95% confidence interval (CI).



**Fig. S3.** Sequence alignment of *E. coli* K12\_SdiA and *EcN*\_SdiA. The red background box indicates identical sequence and red letters indicate diverse sequence.