

Table S1 Impact of various *nif*-gene inactivations on nitrogenase activity in different engineered strains. Relative activities with respect to that for the respective engineered strain with wild-type *nif* cluster.

Genotype	<i>E. coli</i> ^a	<i>E. coli</i> ^b	<i>Synechocystis</i> 6803 ^c
wild-type cluster	100%	100%	100%
$\Delta nifT$	-	-	4%
$\Delta nifV$	1%	25%	-
$\Delta nifB$	0%	0%	-
$\Delta nifS$	5%	-	-
$\Delta nifU$	10%	-	-
$\Delta nifX$	-	50%	<1%
$\Delta nifW$	-	-	7%
$\Delta hesA$	-	50%	-
$\Delta hesB$	-	-	9%
$\Delta nifQ$	90%	-	-
$\Delta nifF$	50%	-	-
$\Delta nifJ$	30%	-	-
$\Delta nifTZ$	-	-	<1%
$\Delta hesAB$	-	-	6%
$\Delta nifXhesA$	-	35%	-
$\Delta nifVhesA$	-	25%	-
$\Delta nifVXhesA$	-	7%	-
$\Delta nifVXNhesA$	-	2%	-
$\Delta nifENX$	110%	-	-
$\Delta nifUSV$	0%	-	-
$\Delta nifWZM$	100%	-	-

^aNitrogen fixation genes from *A. vinelandii* and *K. oxytoca*. Data from (1).

^bNitrogen fixation genes from *Paenibacillus*. Data from (2).

^cActivities shown here are based on the strain TSyNif-2 with 100% activity (this study)