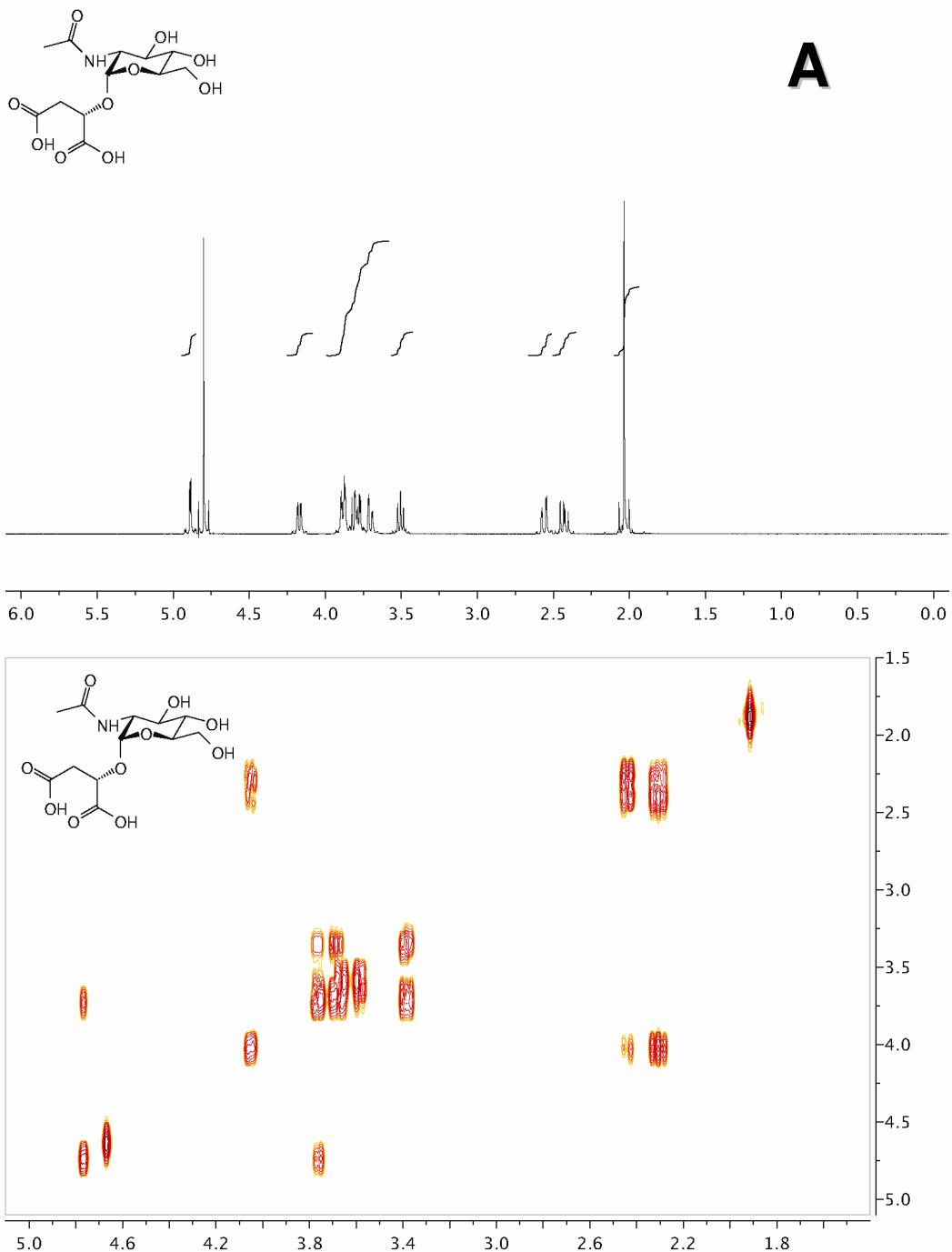
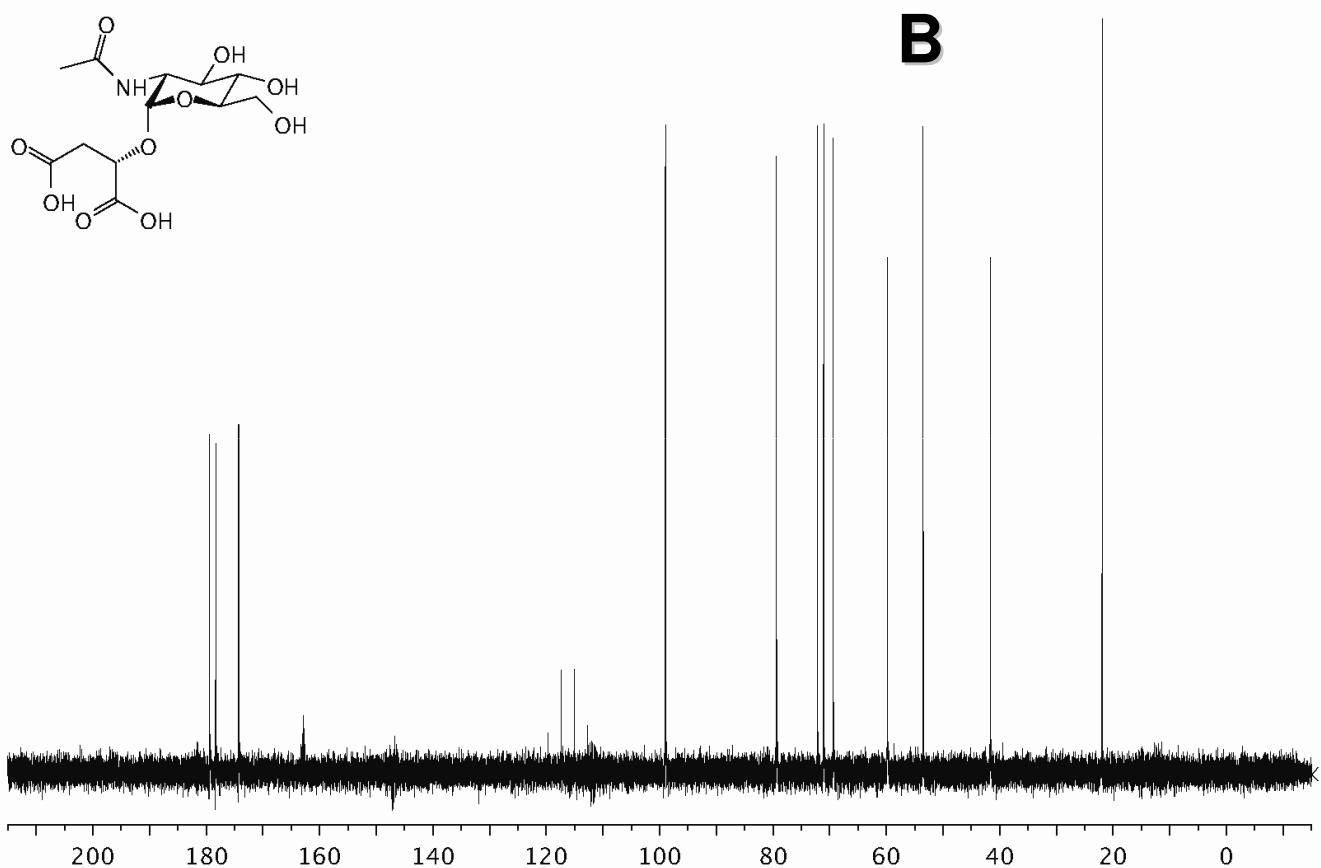


Figure S1. Preparative conversion of UDP-GlcNAc to GlcNAc-malate by *BaBshA* (ORF BA1558). Product was confirmed independently by NMR (^1H , gCOSY, and ^{13}C ; Figure S2) and high resolution mass spectrometry.

Figure S2. NMR spectra of purified GlcNAc-malate (structure given in Figure S2A) synthesized enzymatically with *BaBshA* as described in "Experimental Procedures" and Figure S1. (A) 500 MHz proton NMR spectrum and gradient COSY NMR correlation of GlcNAc-malate at pH 6-7. (B) 125 MHz carbon NMR spectrum at pH 6-7. ^1H NMR (500 MHz, D_2O , trace sodium trifluoroacetate) δ ppm; 4.89 (d, J 3.8 Hz, 1H), 4.17 (dd, J 2.6, 11.1 Hz, 1H), 3.86-3.9 (m, 2H), 3.76-3.83 (m, 2H), 3.70 (dd, J 2.3, 12.5 Hz, 1H), 3.50 (dd, J 10.0, 8.8 Hz, 1H), 2.56 (dd, J 2.5, 15.0 Hz, 1H), 2.43 (dd, J 11.1, 15.0 Hz, 1H,), 2.04 (s, 3H); ^{13}C NMR (125 MHz, D_2O , trace sodium trifluoroacetate) δ ppm; 179.4, 178.3, 174.3, 99.0, 79.4, 72.2, 71.1, 69.4, 59.8, 53.5, 41.6, 21.9.

A

B

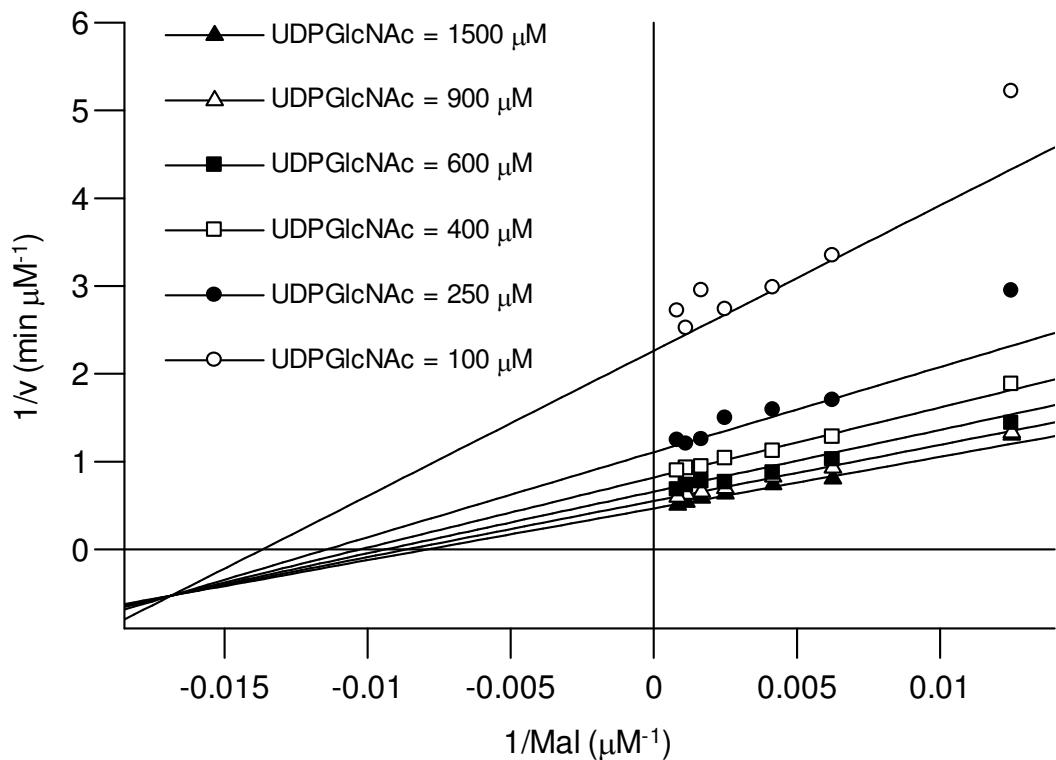


Figure S3. Initial velocity reciprocal plots of *BaBshA* activity. The [L-malate] was varied while [UDP-GlcNAc] was held constant at 1.5 mM (\blacktriangle), 0.9 mM (\triangle), 0.6 mM (\blacksquare), 0.4 mM (\square), 0.25 mM (\bullet), or 0.1 mM (\circ). Solid lines are from fits of the data to equation 2.

Table S1: Primers Used in the Study

Name	Sequence	Use
BAS1444up FWD	CACACACAGCGGCCGCAAAGCAGCAAGGTCTCC	Construction and verification of $\Delta bshA$
BAS1444up REV	CTCATAAATCCTTCGGCATACCCGGGCACTCATAATTGTTCCCCTC	
BAS1444dwn FWD	CCCGGGTATGCCAAGGATTATGAG	
BAS1444dwn REV	CACACACAGCGGCCAACATCAGATTGCTCAATACC	
BAS1444out FWD	CAGTAGGAACAATTATCGCTCC	
BAS1444out REV	CACAAACATCTGCACACG	
BAS1445up FWD	CACACACAGCGGCCGCGAGGGGAACAATTATGAGTG	Construction and verification of $\Delta bshB$
BAS1445up REV	CTGCTCGTAACACACTTCTCTCCCGGGCCAGAACCCACCTACAGAAGG	
BAS1445dwn FWD	CCCGGGAAAGAGAAAGTGTGTTACGAGCAG	
BAS1445dwn REV	TATATATAGCGGCCGCTCCACTGTAATTCTCTCAATTGC	
BAS1445out FWD	CAATCTGGTCCTCTTGGC	
BAS1445out REV	CCCATGCCTCAATGTCTG	
CoADR5f	ATAAGAATGCAGGCCGCGTGGTACGACTATTTGAGCGCATGATCTG	Construction and verification of Δcdr
CoADR5r	CCGCTCGCCTCCGGGACTTACCACTCCTCTTTTACATACTCTTACG	
CoADR3f	GTGGTTAAAGTGCCCGGGAGGGAGCGGAATAGCACTTTAAGTGC	
CoADR3r	CGCGGATCCCCTCAAATTGATGTTAGCTCCCTATATACTTTCTCC	
CDR-F	CCGTATTAGACAGGAAAGCTAACGGC	
CDR-R	GCCCCCTGGTTCGTTCATTGCTTTG	
CDRRHD1	TAATAGTGAGGATCCCCGGGACAGTCTTACCAAGAACGTATTGTAT	Construction and verification of $\Delta cdr2$
CDRRHD2	CCCGGGGGATCCTCACTATTATCCGCCTACTACAACATTTTCTG	
CDRRHD3	CGCGCGGCCGCGTACAGAGAATATTGTGTGTACTGA	
CDRRHD4	CGCGCGGCCGCTACATATACCTGTAGGGGTATAAT	
CDRRHD5	GTTCTGTGAATAAAAGAACAGGAAA	
CDRRHD6	CTTTTATACTCATAGTCGATTCTC	

Table S2: *BaBshA* (ORF BA1558) Acceptor Substrate Specificity

Substrate ^a	Specific activity (nmol/min/mg)	Relative rate
L-Malic acid	4800 ± 400	100
D-Malic acid	19 ± 4	0.4
Inositol 1-L-phosphate	14 ± 4	0.29
D-Glyceric acid	20 ± 1	0.42
Glycolic acid	22 ± 2	0.46
D,L-Isocitric acid	22 ± 3	0.46
Citric acid	22 ± 2	0.46
D-Lactic acid	17 ± 2	0.35

^a0.1 mM Acceptor substrate with 3 mM UDP-GlcNAc, using *BaBshA*

assay conditions described in "Experimental Procedures" (n = 3).