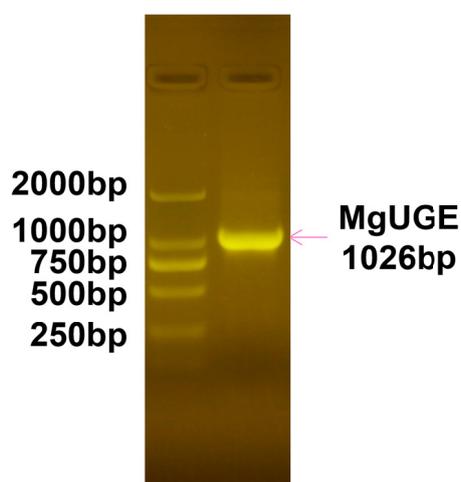
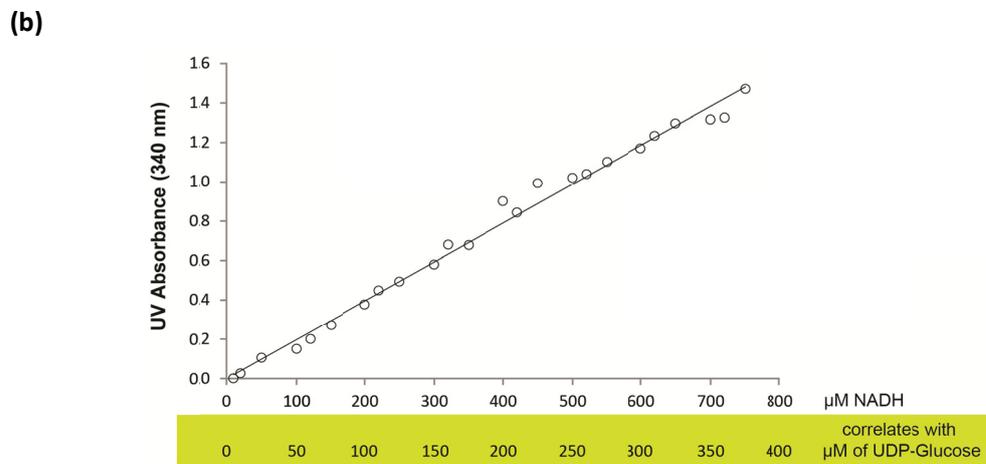
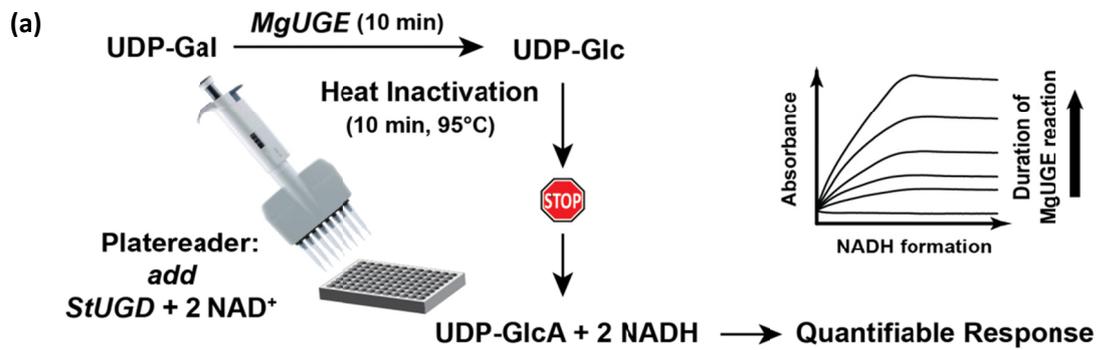

Supplementary Materials: UDP-glucose 4-epimerase and beta-1,4-galactosyltransferase from the oyster *Magallana gigas* as valuable biocatalysts for the production of galactosylated products

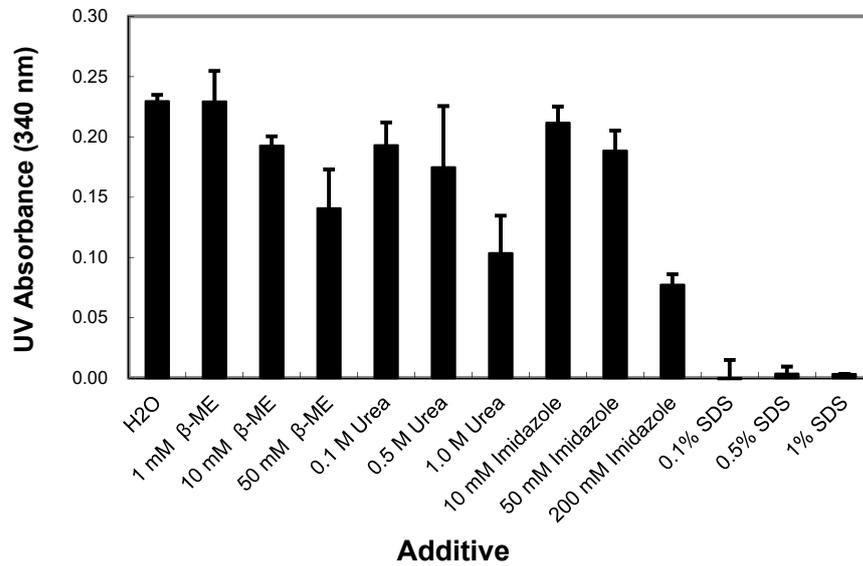
Hui-Bo Song, Meng He, Zhi-Peng Cai, Kun Huang, Sabine L. Flitsch, Li Liu and Josef Voglmeir



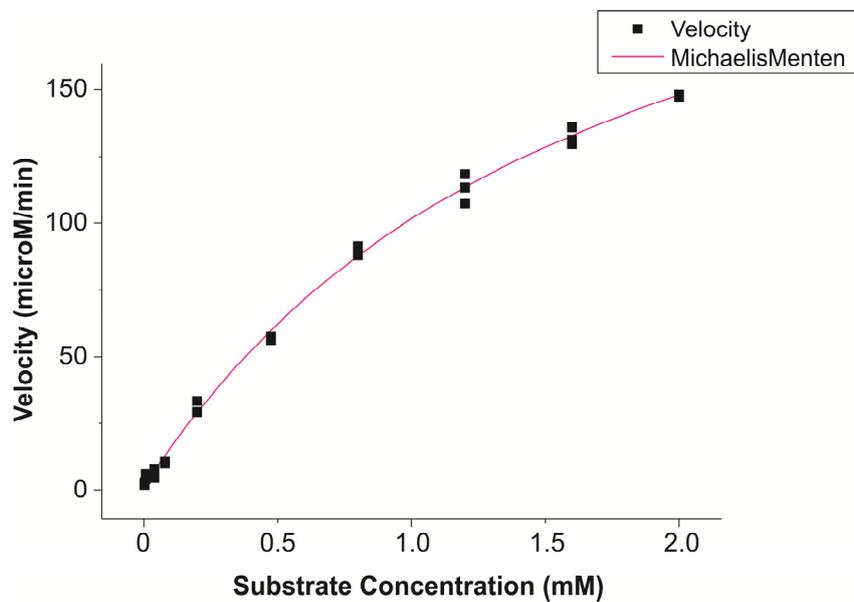
Supplementary Figure S1: Agarose gel electrophoresis of the PCR product of MgUGE.



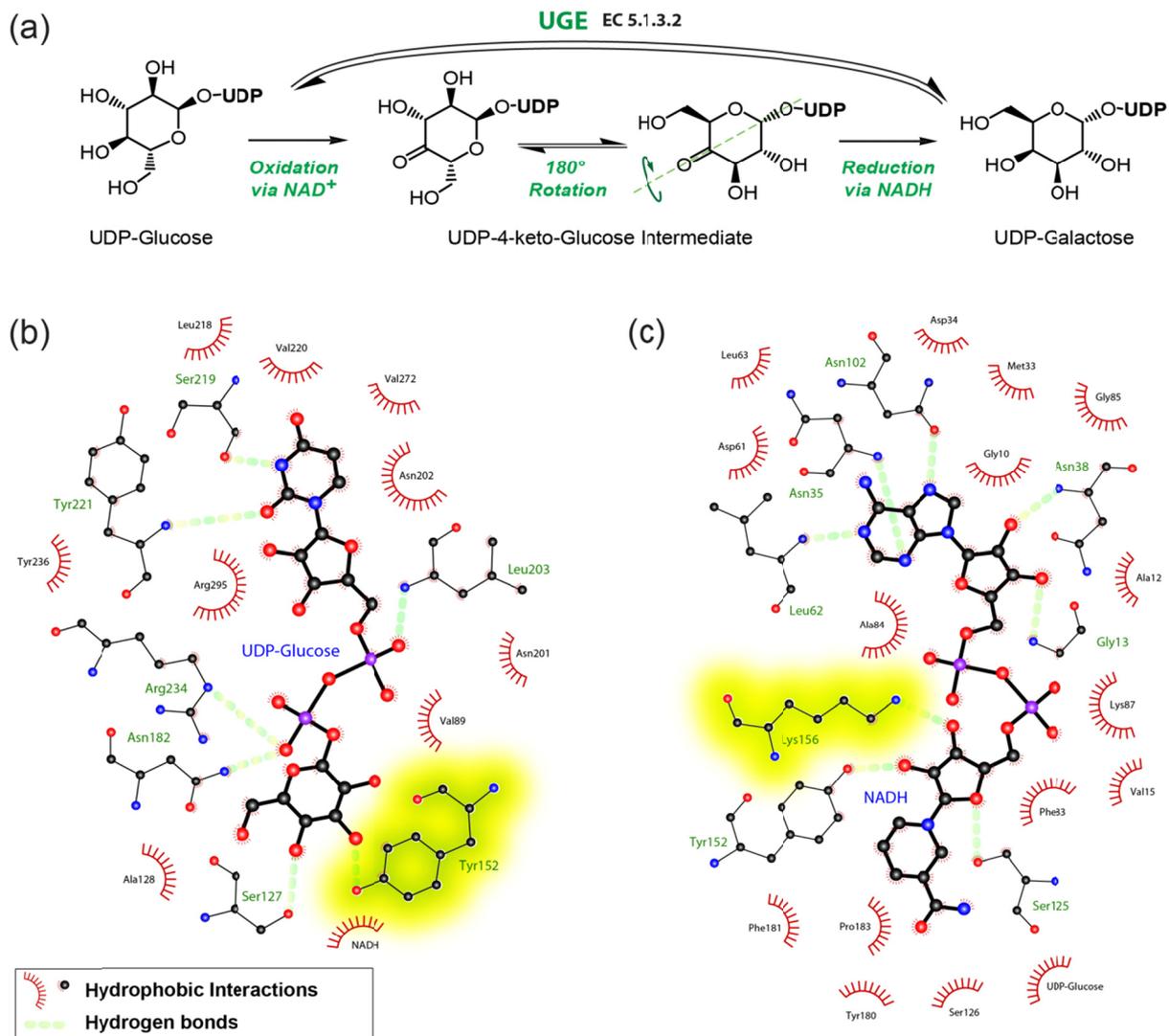
Supplementary Figure S2. (a) Schematic overview of the experimental setup for the plate reader-based spectrophotometric detection of MgUGE activity. (b) Relationship between the UV absorbance of various NADH-concentrations, and the concentration of generated UDP-Glucose.



Supplementary Figure S3: Impact of additives on the enzymatic activity of MgUGE (β -ME – 2-mercaptoethanol; SDS – sodium dodecyl sulfate).



Supplementary Figure S4: Michaelis-Menten Plot for the analysis of the kinetic parameters of MgUGE towards the substrate UDP-galactose.



Supplementary Figure S5. Catalytic mechanism and substrate recognition of UGE. (a) Reaction mechanism of the substrate interconversion within the active site of UGE; Two-dimensional illustration of the substrate-ligand interactions of MgUGE with (b) UDP-Glucose and (c) NADH based on crystallographic data of the human UGE isoform. The catalytic amino acids are highlighted in yellow.

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1
(K1QYA2) Mg MGDSCILVTTGGAGFVGSHSVIELIEAGYSVVMDNLANASM-----ESIKRVEEITGKSIPSYS
(Q14376) Hs MAEK-VLVTGGAGYIGSHTVLELLEAGYLPVVIDNFHNAFRGGGSLPESLRRVQELTGRSVEFEE
(Q9W0P5) Dm MAPPTVLVTGGAGYIGSHTVLEMLNAGYNVICVDNLCNAYSSGAKLPEALSRVQEITGKKVNFYR
(P09147) Ec MRVLVTGGSGYIGSHTCVQLLQNGHDVILLDNLCNSKRS-----VLPVIERLGGKHPFTVE-
(C8VAU8) An MPSGSVLVTGGTGYIGSFTTLALLEAGYKVVVADNLYNSSA-----EALNRIELISGKKAFAQ

60
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(Q14376) Hs MDILDQALQRLFKKYS-FMAVIHFAGLKAVGESVQKPLDYRVNLTGTIQLEIMKAHGKVNLI
(Q9W0P5) Dm VDI TDREQVRSVFQEHK-IDMVAHFAALKAVGESCRIPLYYHNNMTGTNVLLEAMADNNVFKFV
(P09147) Ec GDIRNEALMTEILHDHA-IDTVIHFAGLKAVGESVQKPLEYYDNNVNGTLRLISAMRAANVKNFI
(C8VAU8) An LDVTDEAAFDKVFEAHPDIDSVIHFAALKAVGESGKPLDYHVNVYGTICLLRSMVRHNVTNIV

124
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(Q14376) Hs FSSSATVYGN----PQYLPIDEAHP TGGCTNPYGKSYFFIEEMIRDLCAADKT-----
(Q9W0P5) Dm YSSSATVYGE----PKFLPVTEEHPTGNC TSPYGKTYFYFIEEILKDLCKSDKR-----
(P09147) Ec FSSSATVYGD----NPKIYVYESFPTGTPQS P YGKSYLMVEQILTDLQKAQPD-----
(C8VAU8) An FSSSATVYGDATRFPMIPIPEHCLG-PTNPTGNTYFAIELAITDVINAQRNNAKAGNETEAA

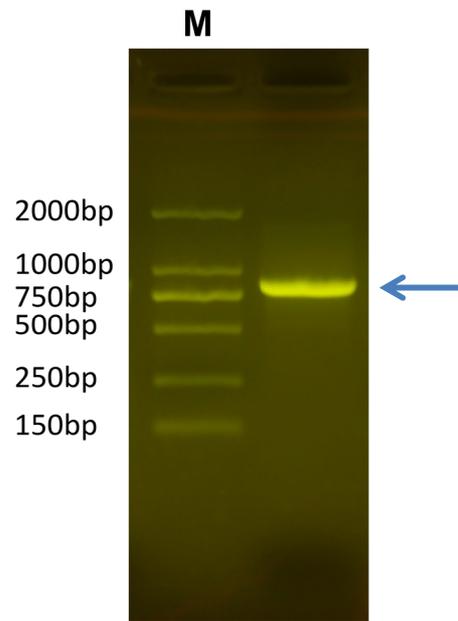
173
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(Q9W0P5) Dm -WAVVSLR YFN PVGAHISGRIGEDPNGE PNNLMPYIAQVAVGRRPSLSVYGSDFPTH DGTGVRDY
(P09147) Ec -WSIALLR YFN PVGAHPSGDMGEDPQGI PNNLMPYIAQVAVGRRDSLAI FGN DYPTEDGTGVRDY
(C8VAU8) An KWNGALLR YFN PAGAHPSGIMGEDPQGV PYNLLPLLAQVATGKREKLLVFGDDYASHDGTAIRDY

237
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(Q14376) Hs IHVVDLAKGHIAALRKLKE--QCQCRIYNLGTGTGYSVLQM VQAMEKASGKKIPYKVVARREGDV
(Q9W0P5) Dm IHIVDLAEGHVKALDKLRNIAETGFFAYNLGTGVGYSVLD MVKAF EKASGKKVNYTLVDRRSGDV
(P09147) Ec IHVMDLADGHVVA MEKLAN--KPGVHIYNLGAGVGN SVLDVVNAFSKACGKPVNYHFAPRREGDL
(C8VAU8) An IHILD LADGHLKALNYLRA--NNPGVRAWNLGTGRGSTVYEMIRAFSKAVGRDLPYEVAPRRAGDV

300
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(Q14376) Hs AACYANPSLAQEELGWTAALGLDRMCEDLWRWQKQNP SGFGTQA
(Q9W0P5) Dm ATCYADATLADKKL GWKAERGIDKMCEDTWRWQSQNPNGYANK
(P09147) Ec PAYWADASKADRELNWRVTRTLDEMAQD TWHWQSRHPQGYPD
(C8VAU8) An LNLTSNPTRANTELGWKAQRTLEQACEDLWLWTKNNPQGYRQPPAELLEQLKK

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Supplementary Figure S6: Homology alignment of MgUGE with UGE homologues from *Homo sapiens* (Hs), *Drosophila melanogaster* (Dm), *Escherichia coli* (Ec) and *Aspergillus nidulans* (An). The tyrosine and lysine residues highlighted in green are the proposed catalytic amino acids. Uniprot identifiers are shown in parentheses.



Supplementary Figure S7: Agarose gel electrophoresis of the PCR product of MgGalT7.