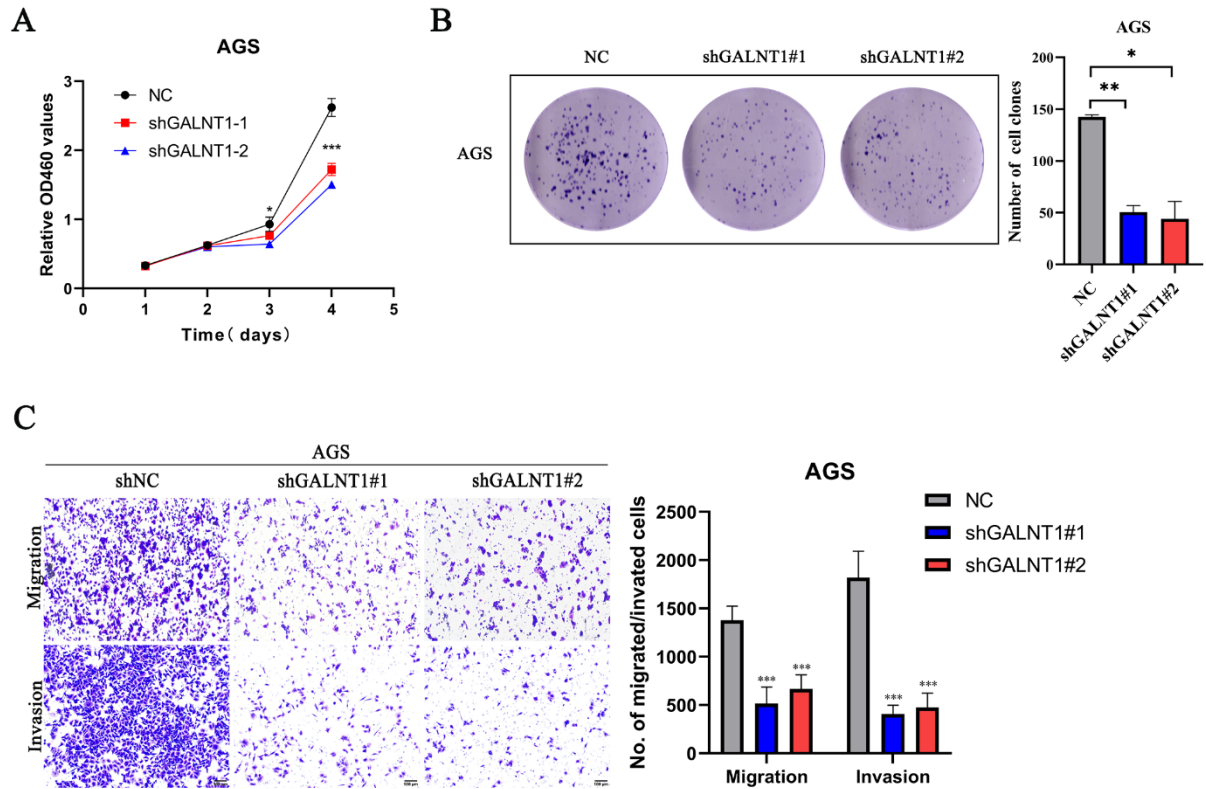
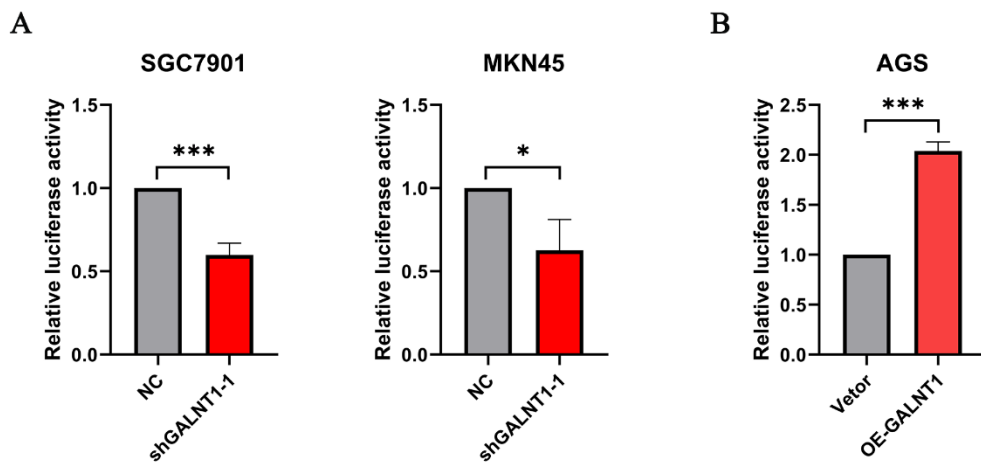


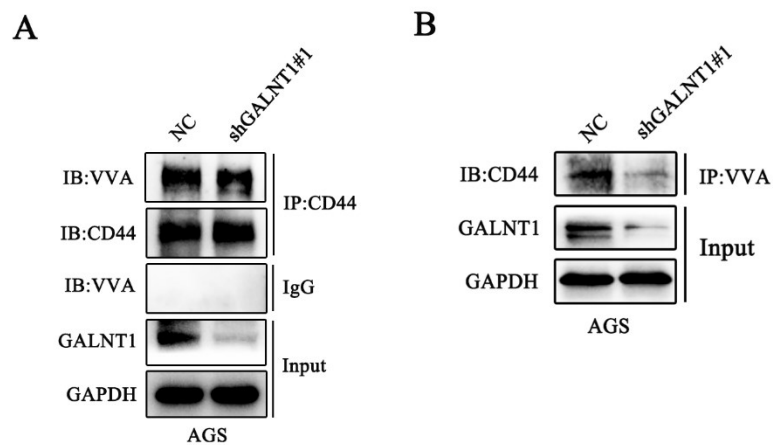
Supplementary figure S1: (A–D) Analysis of *GALNTs* expression in gastric cancer using GEPIA database. **E**, Analysis of *GALNTs* expression in gastric cancer each molecular subtype.



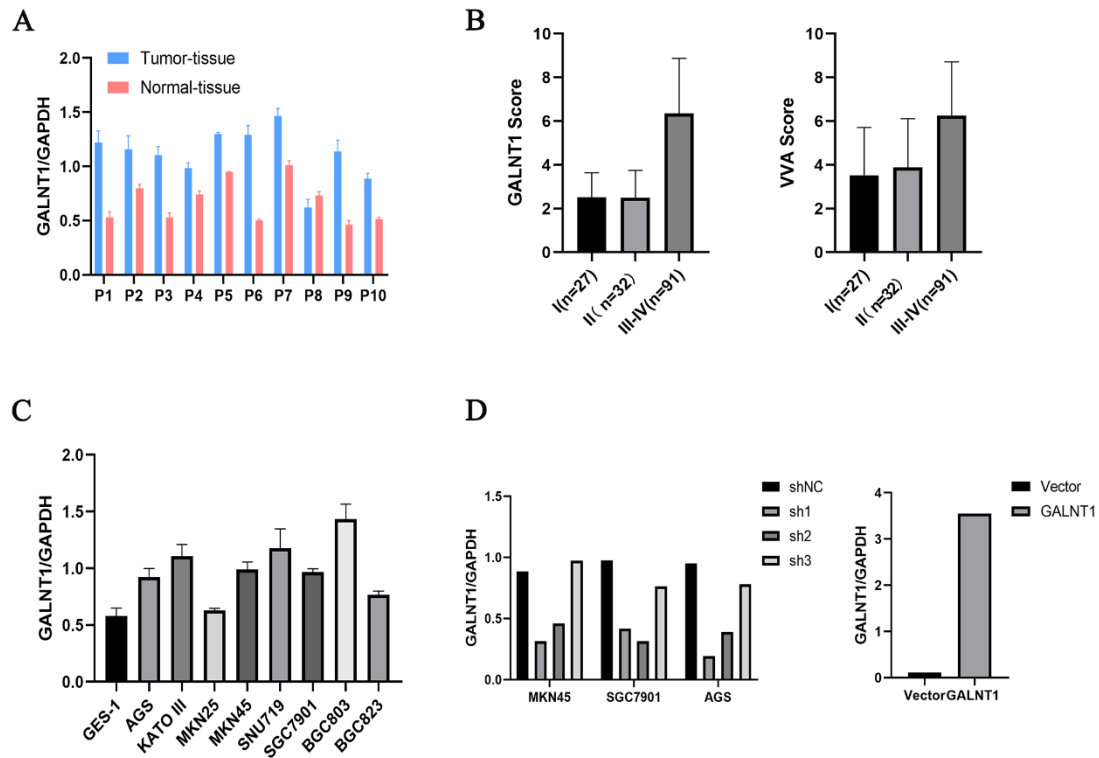
Supplementary figure S2: GALNT1 promotes the growth and metastatic ability of AGS cells and modifies O-glycosylation of CD44. A–B, AGS cells' growth abilities after *GALNT1* knockdown were determined by CCK8 assay (A) and colony formation assay (B). C, *GALNT1* knockdown decreased the abilities of migration and invasion of AGS cells.



Supplementary figure S3: TOP luciferase reporter assays were measured in GALNT1-silenced (**A**) and -overexpressing cells (**B**). Normalization was based on internal Renilla luciferase activity.



Supplementary figure S4: *GALNT1* knockdown decreased the O-glycosylation levels of CD44 by co-immunoprecipitation in AGS cells (**A-B**).



Supplementary figure S5: (A) The quantification of *GALNT1* levels in paired primary tumors and adjacent normal tissues. (B) Statistical analysis of *GALNT1* staining at different clinical stages. (C) The quantification of *GALNT1* levels in gastric cancer cell lines. (D) The quantification of the silencing efficiency of *GALNT1* in MKN45, AGS, and SGC7901 cells as well as the overexpression of *GALNT1* in AGS cells.

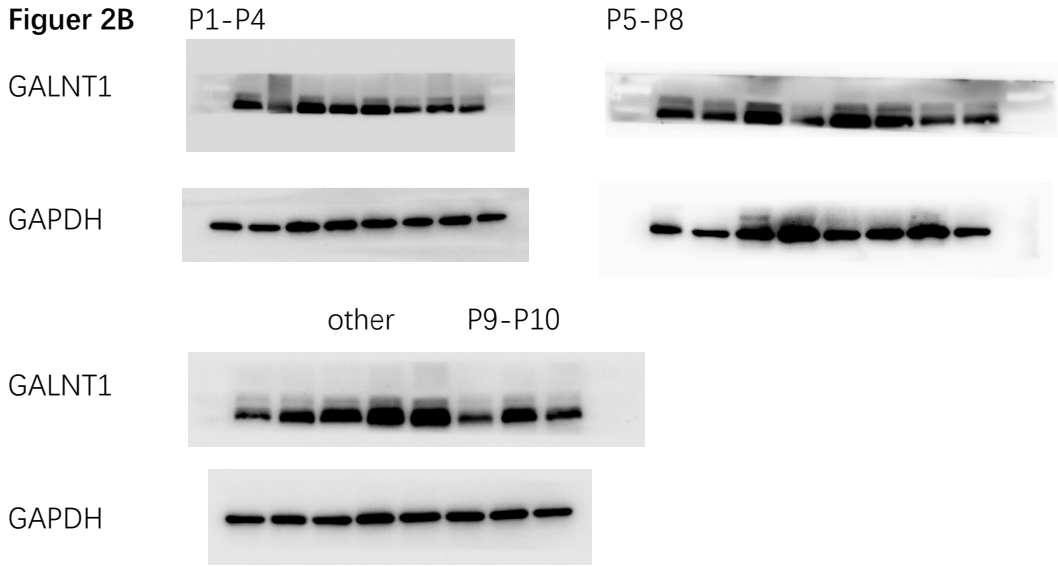
Table S1. The target sequence of sh*GALNT1* and sh*CD44*

Gene	Target sequence
sh <i>GALNT1</i> -1	5'– GCTTGGATGTTTCCAAACTTA –3'
sh <i>GALNT1</i> -2	5'– ATTGATCAGAGCTAGATTAAA –3'
sh <i>GALNT1</i> -3	5'– GACGTGAAACTGCATAGTAAT –3'
Sh <i>CD44</i> -1	5'– GGC GCAGATCGATTTGAATAT –3'
sh <i>CD44</i> -2	5'– CCAGTATGACACATATTGCTT –3'
sh <i>CD44</i> -3	5'– CCATTCAAATCCGGAAGTGCT –3'

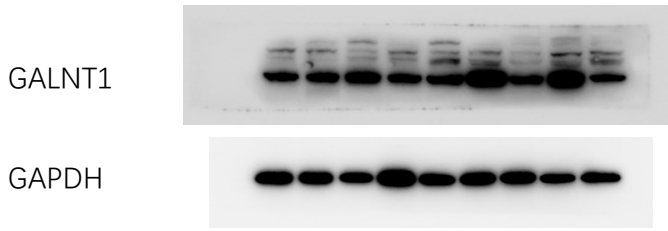
Table S2. Sequences of primers used in this study for qRT-PCR.

Gene	Sequence
<i>GALNT1</i>	F:5'- GTGAGATGATTGCACTCAACAGA -3'
	R:5'- ACTATGGACAGTTCGCAGAAGT -3'
GAPDH	F:5'- CGGAGTCAACGGATTTGGTCGT -3'
	R:5'- TCTCAGCCTTGACGGTGCCA -3'
CD44	F:5'- TCCTCACATCCAACACCTCC-3'
	R:5'- GCTGCTCACGTCATCATCAG -3'

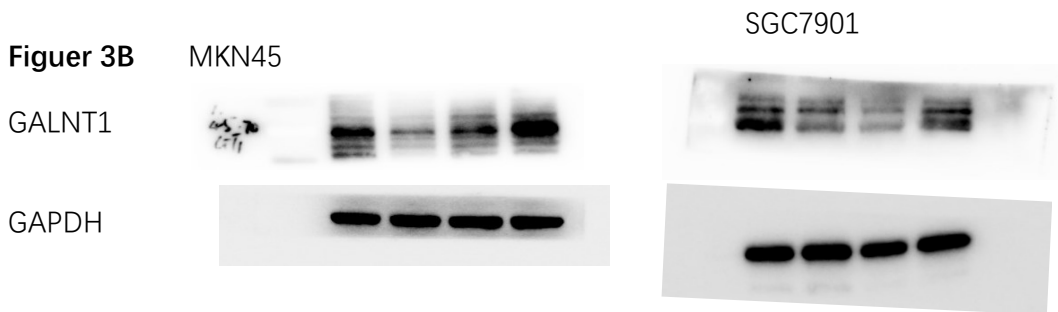
Figuer 2B



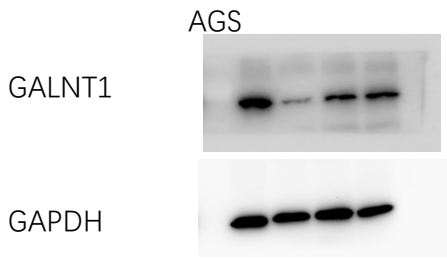
Figuer 3A



Figuer 3B



AGS



AGS-OE

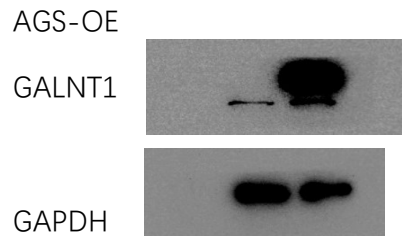
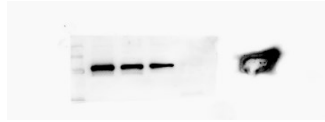


Figure 5C-F



MKN45

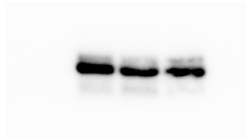


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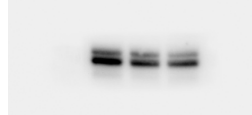


C-MYC

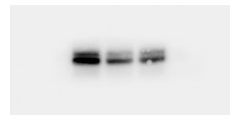
SGC7901



AGS



MKN45



AGS-OE

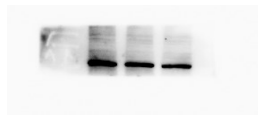


7901 p-b-catenin

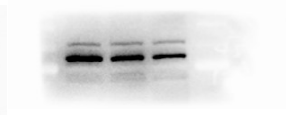
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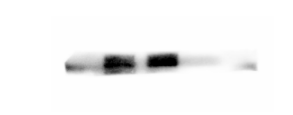
AGS



MKN45



AGS-OE

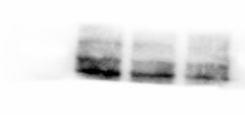


GALNT1

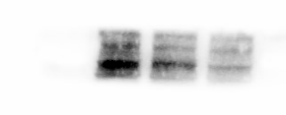
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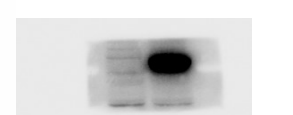
AGS



MKN45

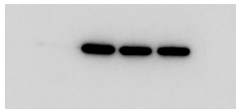


AGS-OE

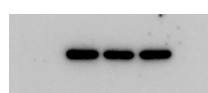


GAPDH

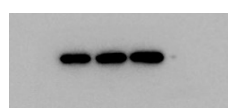
SGC7901



AGS



MKN45



AGS-OE

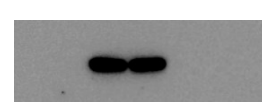


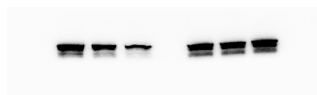
Figure 5G

MKN45

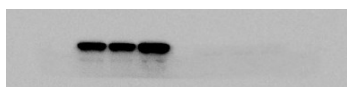
Nuclear

Cytoplasm

β -catenin



LaminB1



GAPDH

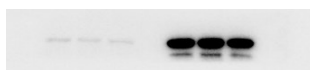


Figure 5H

SGC7901

Nuclear

Cytoplasm

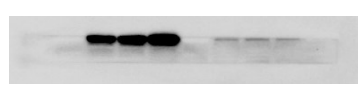
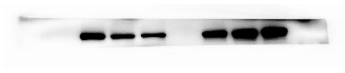


Figure 5I

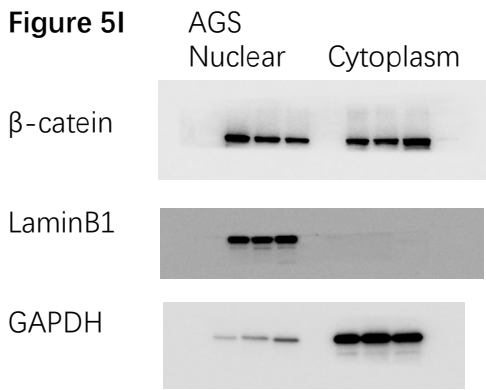


Figure 5J

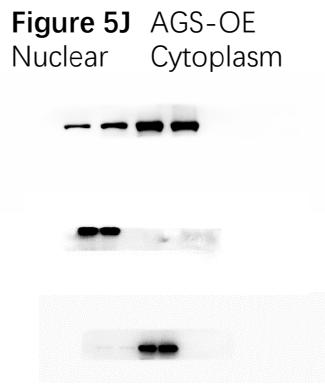
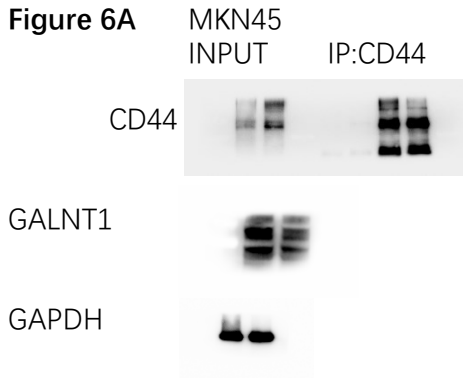


Figure 6A



IP:CD44

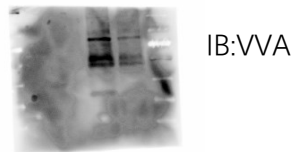
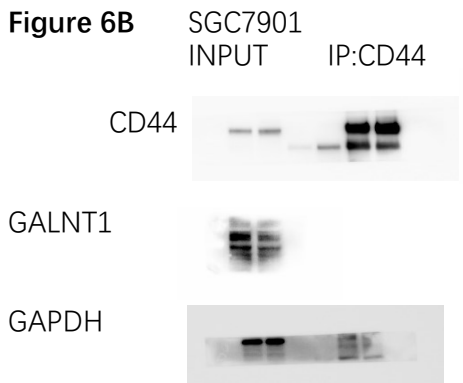


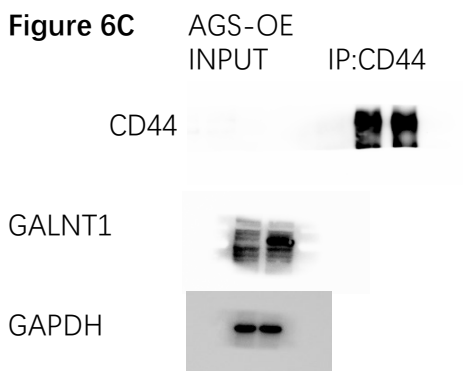
Figure 6B



IP:CD44



Figure 6C



IP:CD44

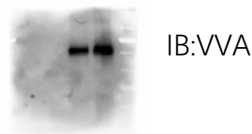


Figure 6D

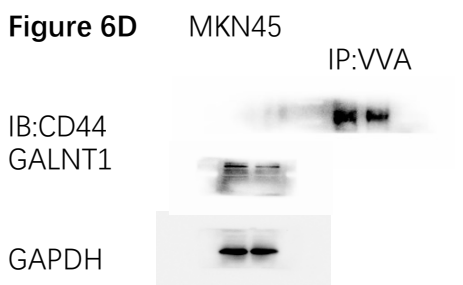


Figure 6E



