

Supplementary Information

***Helicobacter pylori*-induced gastric cancer is orchestrated by MRCK β -mediated Siah2 phosphorylation**

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>SIAH2

MSRPSSTGPSANKPCSKQPPPQPQHTPSPAAPPAAATISAAGPGSSAVPAAAavisGPG
GGGGAGPVSPQHHELTSFECPCVCFDVLPPILQCQAGHLVCNCRQKLSCCPTCRGA
LTPSIRNLAMEKVASAVLFPCKYATTGCSLT LHHTEKPEHEDICEYRPYSCPCPGASCKW
QGSLEAVMSHLMHAHKSIITLQGEDIVFLATDINLPGAVDWVMMQSCFGHHFMLVLEKQ
EKYEGHQQFFAIVLLIGTRKQAENFAYRLENGNRRRLTWEATPRSIHDGVAAIMNSDC
LVFDTAIAHLFADNGNLGINVTISTCCP

>MRCK β

MSAKVRLKKLEQLLLDGPWRNESALSVENTLDVLVCLYTECSHSALRRDKYVAEFL
KPFTQLVKEMQLHREDFEIIVIGRGAFAVEAVVKMKNTERIYAMKILNKWEMLKRAETA
CFREERDVNVGDCQWITALHYAFQDENHLYLVMDYYVGGDLTLLSKFEDKLPEDMAR
FYIGEMVLAIDSIHQLHYVHRDIKPDNVLLDVNGHIRLADFGSCLKMNDDGTVQSSVAVG
TPDYISPEILQAMEDGMGKYGPECDDWWSLGVCMYEMLYGETPFYAESLVETYGKIMNH
EERFQFPShVTDVSEEAKDLIQRILCSRERRLGQNGIEDFKKHAFFEGLNWENIRNLEAP
YIPDVSSPSDTSNFDVDDDVLRNTEILPPGSHTGFSGLHLFIGFTFTTESCFSDRGS
SIMQSNTLTKDEVQRDLEHSLQMEAYERRIRRLQEKELESRKLQESTQTVQSLHGSS
RALSNNSNRDKEIKKLNEEIERLKNKIADSNRLERQLEDTVLRQEREDSTQRLRGLEKQH
RVVRQEKEELHKQLVEASERLKSQAKELKDAHQQRKLALQESELNERMAELRAQKQK
VSRQLRDKEEEMEVATQKVDAMRQEMRRAEKLKELEAQLDDAVA
EASKERKLREHSE
NFCKQMESELEALKVKQGGRGAGATLEHQQEISKIKSELEKKVLFYEEELVRREASHVL
EVKNVKKEVHDSESHQLALQKEILMLDKLEKS
KRERHNEMEEAVGTIKDKYERERAML
FDENKKLTAENEKLCFSVDKLT
AQNRQLEDELQD
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AKKESVAHWEAQIAEIIQWV
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DARGYLQALASKMTEE
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LGSRTLDPLW
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-----Predicted interaction result of the 1st two proteins:-----

The probability for interacting of the 1st two proteins: 0.9943

The probability for non-interacting of the 1st two proteins: 0.0057

Decision: SIAH2 and MRCK β can interact

Fig. S1. Result of LR_PPI analysis showing the probability of Siah2-MRCK β interaction.

A

Siah2 sites	Predicted score
<u>S⁶</u>	0.996
<u>T²⁷⁵</u>	0.986
<u>T²⁷⁹</u>	0.931
<u>S²⁸²</u>	0.992

B

1 MSRPSS6TGPSANKPCSKQPPPQPQHTPSAAPPAAATISAAGPGSSAVPA 50
 51 AAAVISGPGGGGAGPVSPQHHELTSLFECPVCFDYVLPPILQCQAGHLV 100
 101 CNQCRQKLSCCPTCRGALTPSIRNLAMEKVASAVLFPCKYATTGCSLT LH 150
 151 HTEKPEHEDICEYRPYSCP CPGASCKWQGSLEAVMSHLMHAHKSI TT LQG 200
 201 EDIVFLATDINLPGAVDWVMMQSCFGHHFMLVLEKQE KYEGHQ QFFAIVL 250
 251 LIGTRKQAENFAYRLELNGNRRRLT²⁷⁵WEAT279PRS282IHDGVAAIMNSDCLVFD³⁰⁰
 301 TAI AHLFADNGNLGINVTISTCCP³²⁴



Fig. S2. Identification of novel phosphorylatable residues. (A) Novel Siah2 phosphorylatable residues along with their predicted scores as identified using NetPhos 3.0. (B) Sequence and multi-domain schematic depiction of human Siah2 protein. Identified highly probable phosphorylatable amino acids are underlined in the sequence and represented by vertical lines in the domain structure.

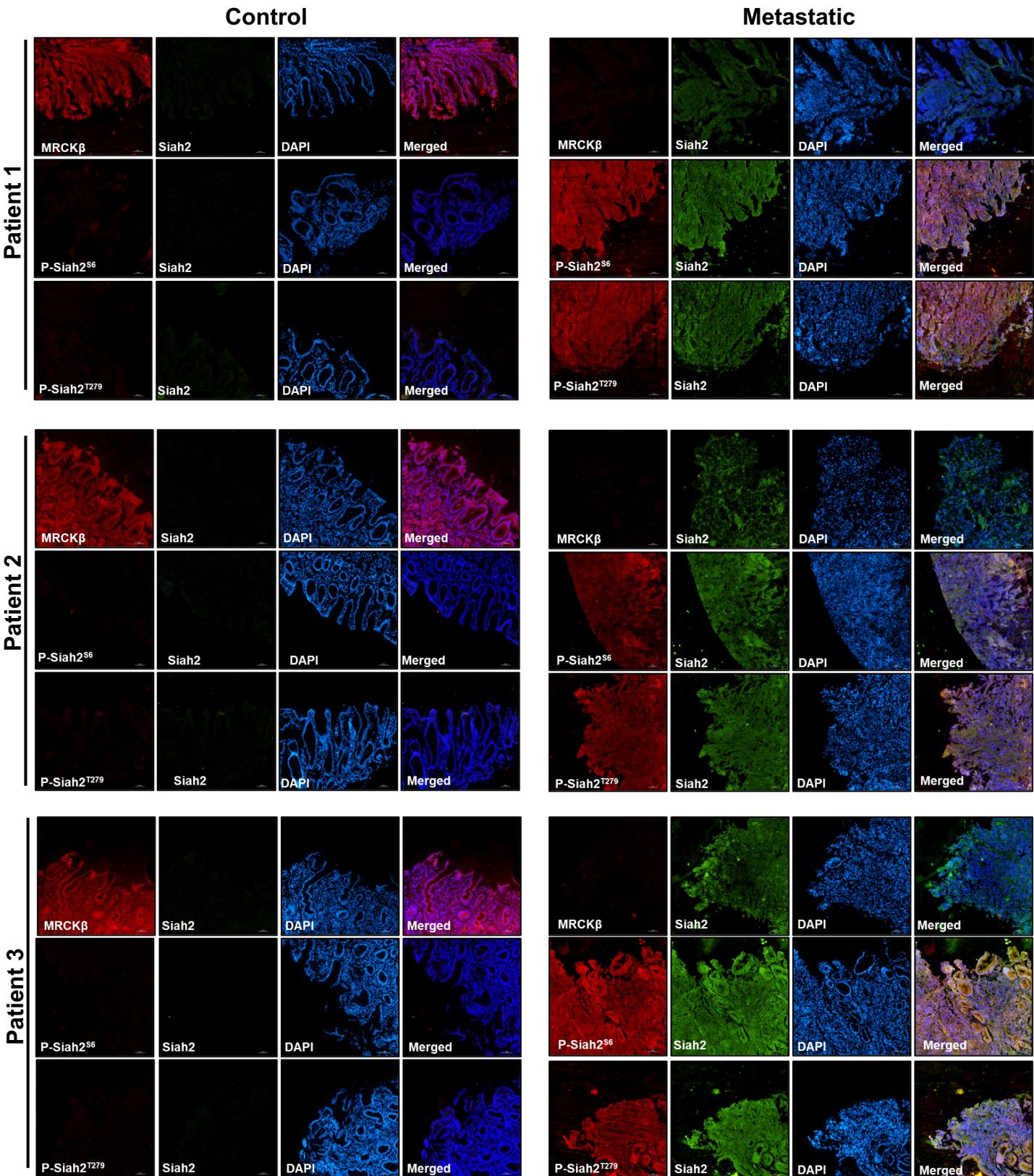


Fig. S3i. Human gastric biopsy tissues exhibit enhanced Siah2 phosphorylation and MRCK β downregulation. Immunofluorescence microscopy images showing the status of Siah2, P-S⁶-Siah2, P-T²⁷⁹-Siah2 and MRCK β in Patient #1-3. Objective: 20X and scale bar represents 50 μ m.

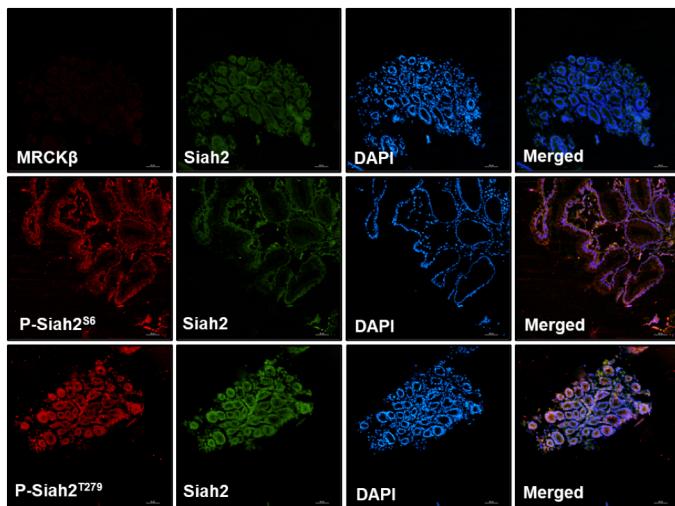
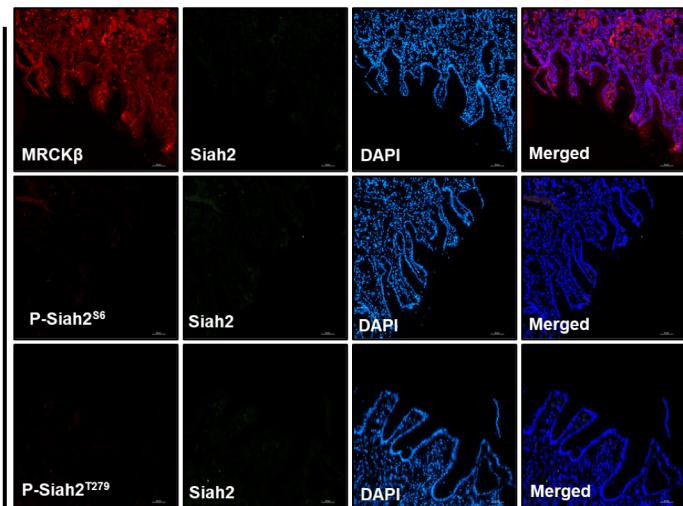
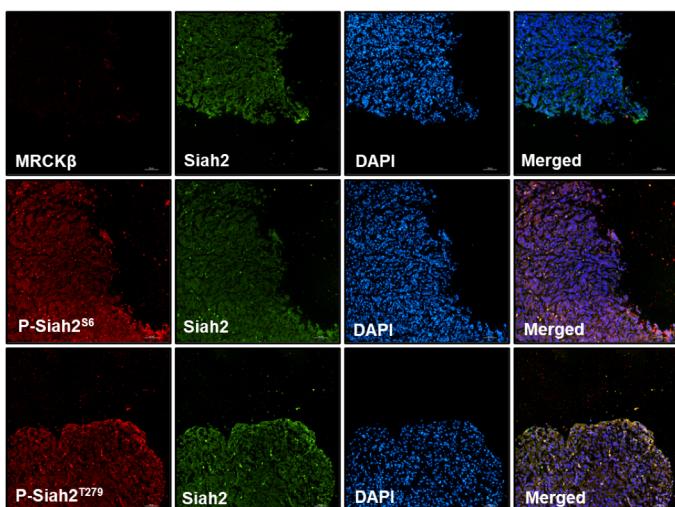
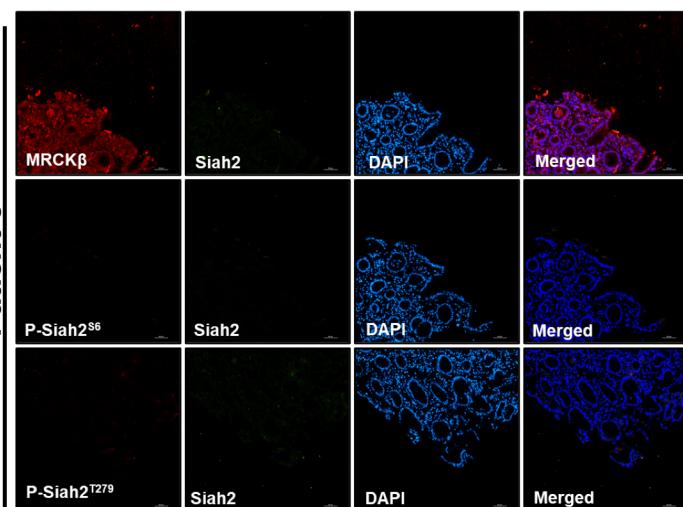
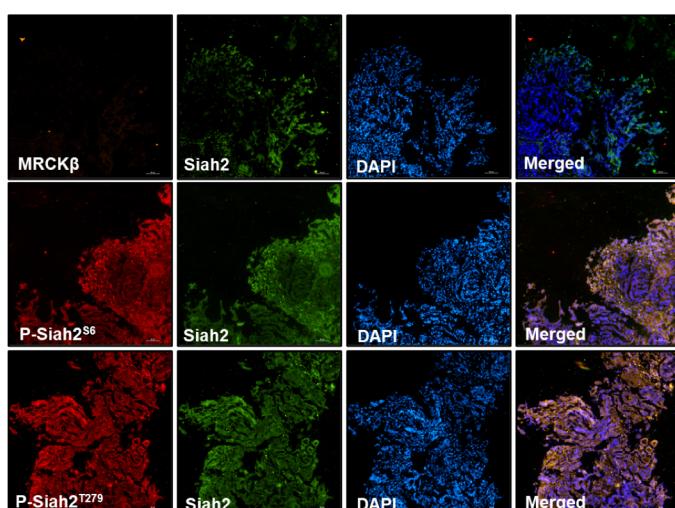
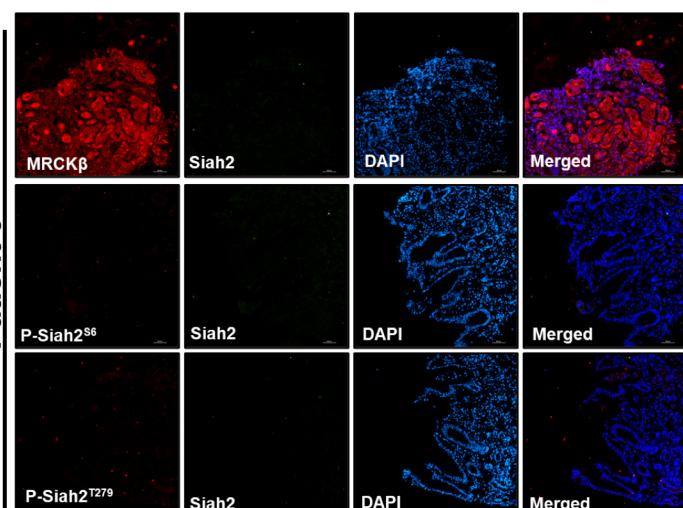
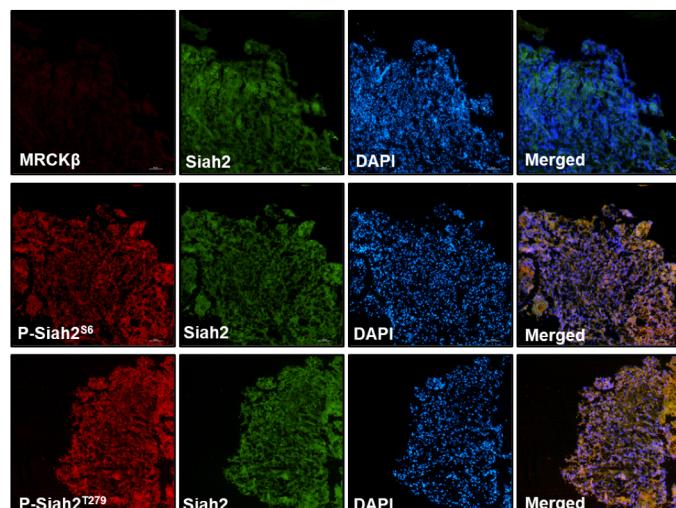
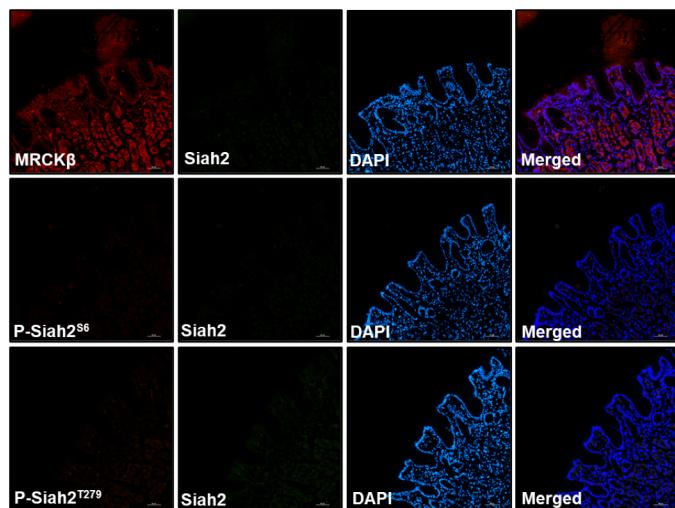
Control**Metastatic****Patient 4****Patient 5****Patient 6**

Fig. S3ii. Human gastric biopsy tissues exhibit enhanced Siah2 phosphorylation and MRCK β downregulation. Immunofluorescence microscopy images showing the status of Siah2, P-S 6 -Siah2, P-T 279 -Siah2 and MRCK β in Patient #4-6. Objective: 20X and scale bar represents 50 μ m.

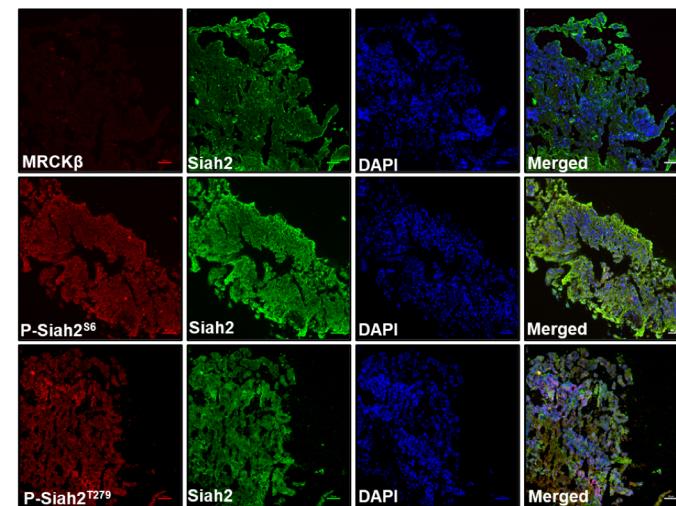
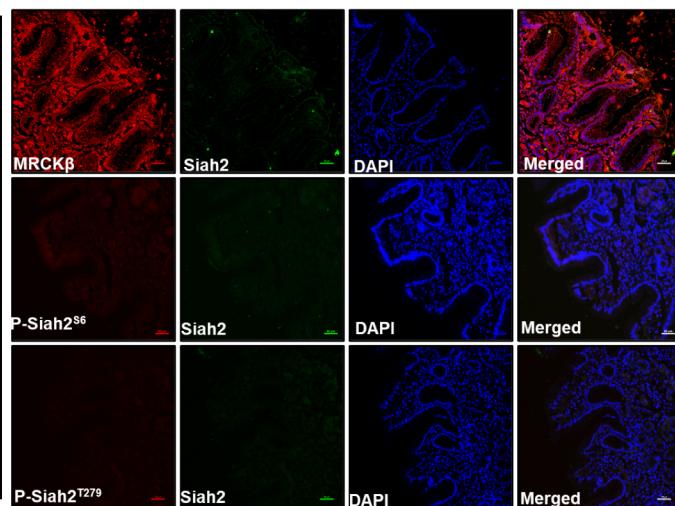
Control

Metastatic

Patient 7



Patient 8



Patient 9

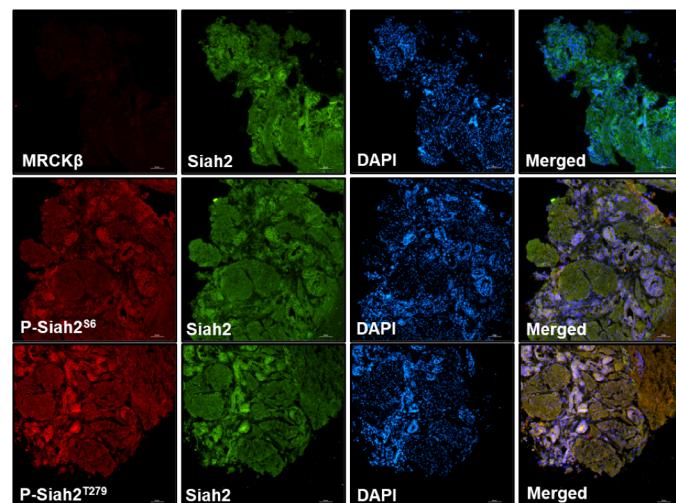
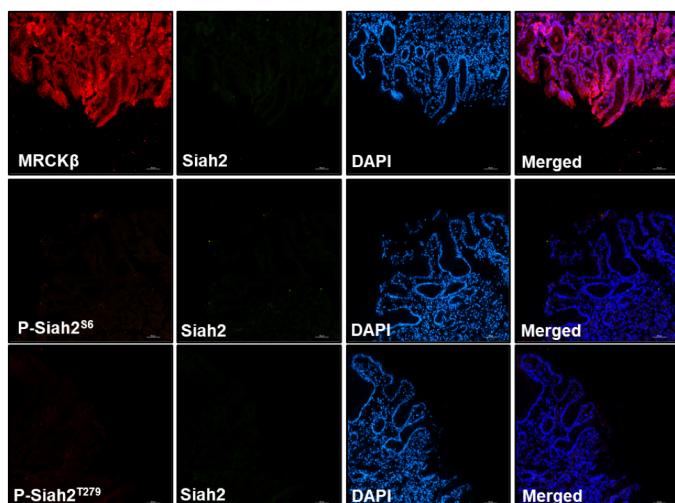


Fig. S3iii. Human gastric biopsy tissues exhibit enhanced Siah2 phosphorylation and MRCK β downregulation. Immunofluorescence microscopy images showing the status of Siah2, P-S⁶-Siah2, P-T²⁷⁹-Siah2 and MRCK β in Patient #7-9. Objective: 20X and scale bar represents 50 μ m.

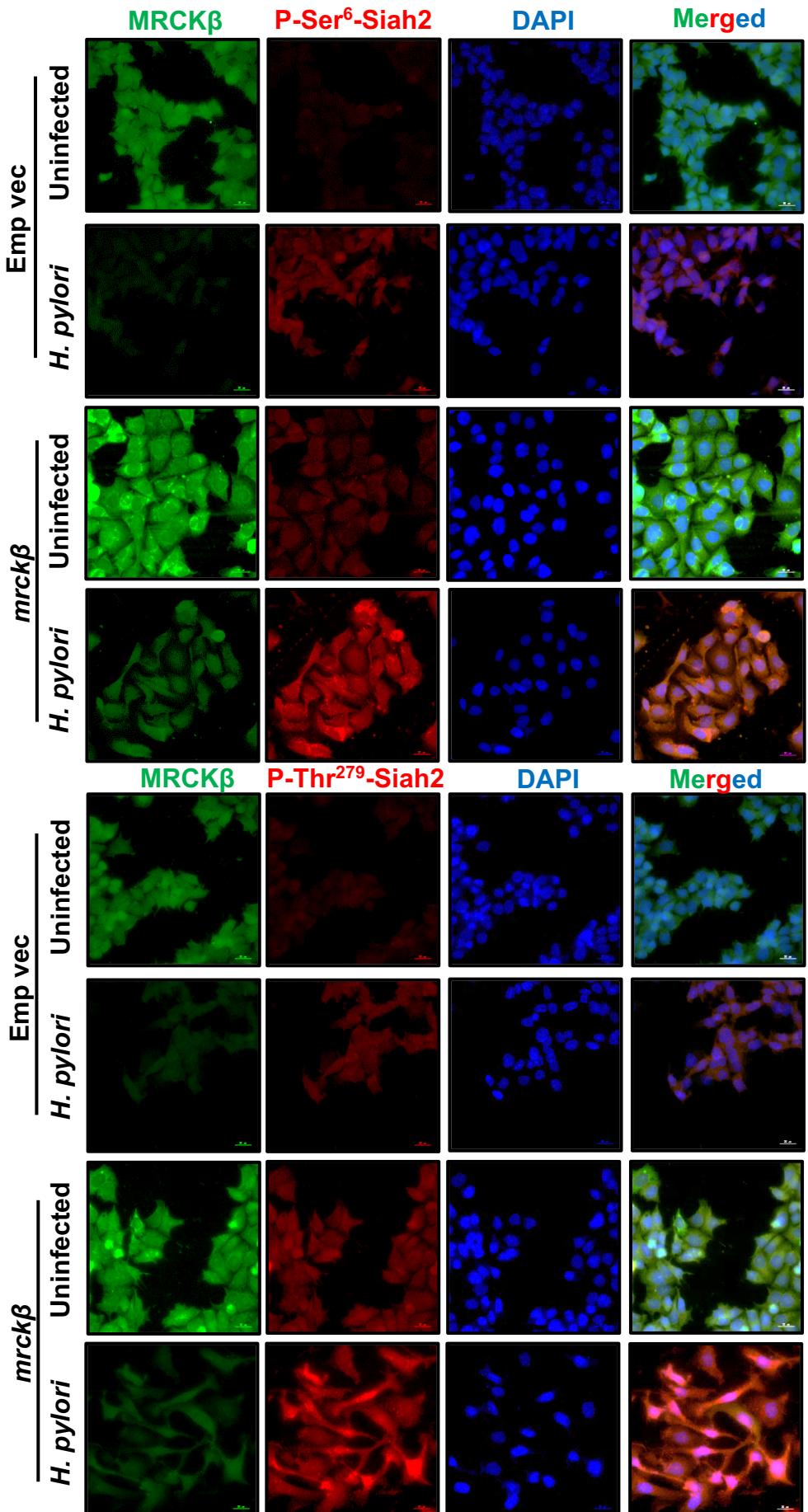


Fig. S4. Phosphorylated Siah2 is enhanced upon *mrck* β overexpression.
 Immunofluorescence microscopy images of AGS cells overexpressing the empty vector or *mrck* β and infected with 200 MOI *H. pylori* for 12 h showing enhanced P-S⁶-Siah2 and P-T²⁷⁹-Siah2. Objective: 60X and scale bar represents 20 μ m.