

Establishment of a Patient-derived Xenograft Model of Pancreatic Grade 3 Well-differentiated Neuroendocrine Tumor

Yosuke Kasai^{1, 2}, Grace E. Kim^{2, 3}, Byron Hann², David B. Donner^{1, 2}, Robert S. Warren^{1, 2}, Emily K. Bergsland^{2, 4}, Eric K. Nakakura^{1, 2}
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Abstract

Abstract

Introduction

Methods

Result 1

Result 2

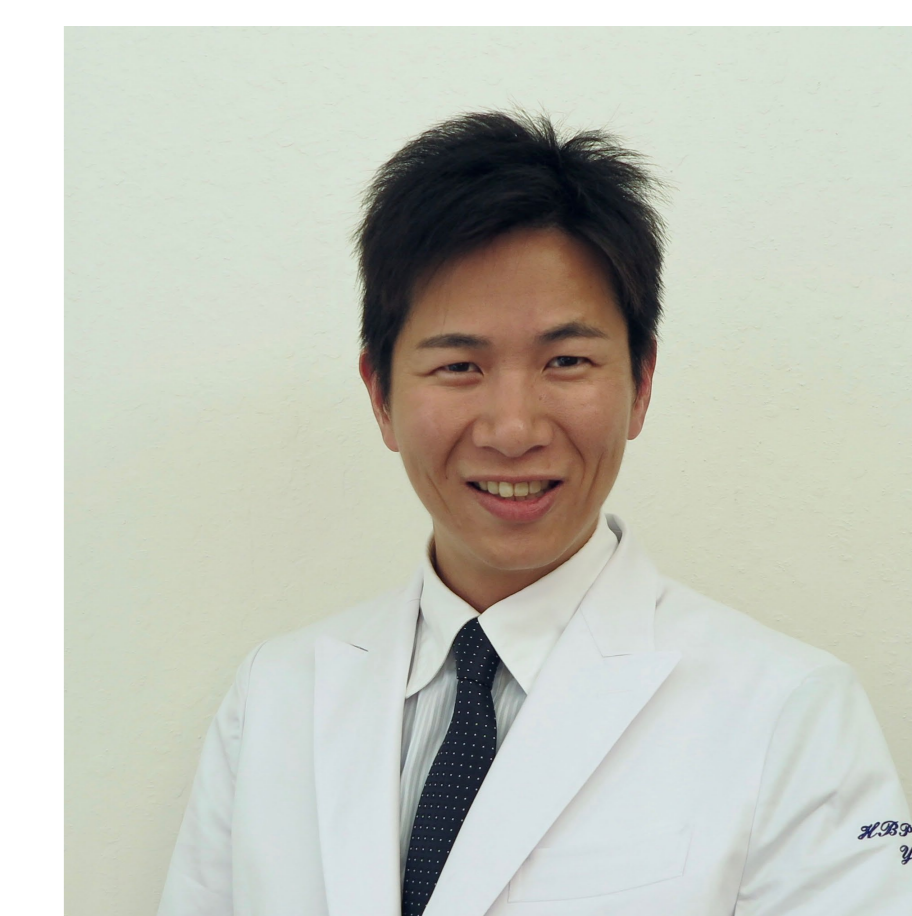
Conclusion

Background: Recently, a new pathologic category of pancreatic neuroendocrine tumors (PNETs), grade 3 well-differentiated-PNET (G3 WD-PNET), was recognized by the WHO with an intermediate prognosis compared to G2 WD-PNET and G3 poorly differentiated neuroendocrine carcinoma. The molecular and clinical features of G3 WD-PNET are ill-defined, and no standard treatment exists. We developed a patient-derived xenograft (PDX) model of G3 WD-PNET as a platform to study its biology and to inform prospective trials.

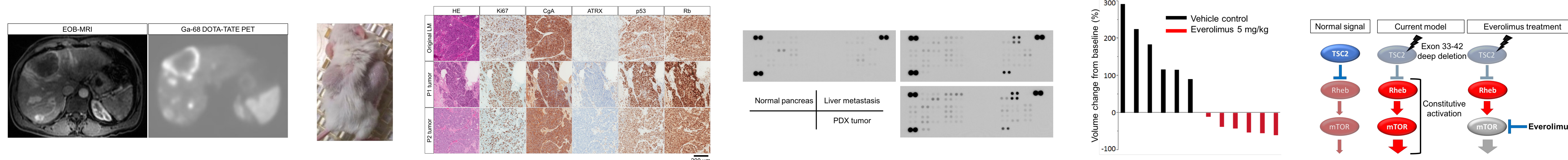
Methods: Surgically resected liver metastasis tissue from a patient with a G3 WD-PNET was implanted s.c. into NSG mice. Pathologic examination of the original patient tumor revealed a WD-PNET with 33 mitotic figures/10 hpf, Ki67 index 47%, and loss of ATRX, indicating a G3 WD-PNET. PDX tumors were passed into the next generation when the tumor volume reached 1,000-2,000 mm³. The morphologic and molecular features of the PDX tumors were compared to the original tumor. Mice bearing PDX tumors were treated with control vehicle or everolimus (5 mg/kg/day p.o.) for 28 days.

Results: Tumors grew successfully in all mice (100% take rate). Chromogranin A was diffusely positive, and the mitotic figures and Ki67 index were comparable with those of the original patient tumor. The patient and PDX tumor showed similar phospho-kinase activity, including phosphorylation of mTOR and p53. These findings were consistent with targeted sequencing results of the patient tumor showing mutations in p53 and TSC2, a negative regulator of mTOR. Everolimus significantly inhibited G3 WD-PNET PDX growth compared to control vehicle (median volume change -48% versus 150%, Wilcoxon $P = 0.005$), showing the importance of the mTOR pathway for tumor growth.

Conclusion: We successfully established a G3 WD-PNET PDX model. This model will facilitate studies of the underlying biology of G3 WD-PNET and help inform prospective clinical trials, which are desperately needed for this new disease.



Yosuke Kasai, MD, PhD



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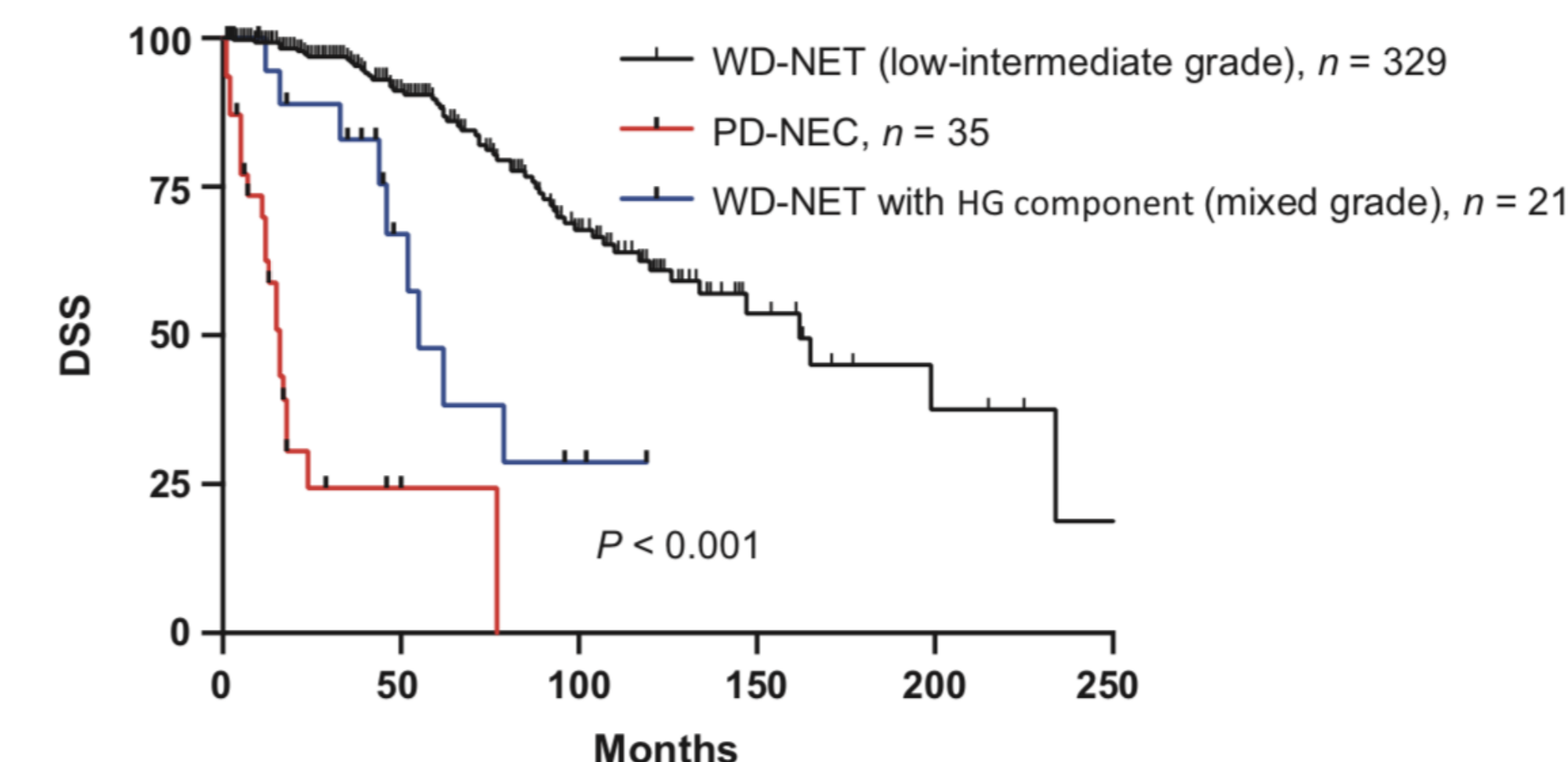
WHO2010 classification

Classification/Grade		Ki67 index	MI /10HPF
PNET	G1	≤ 2%	< 2
	G2	3-20%	2-20
PNEC	G3	> 20%	> 20
	Small cell Large cell		

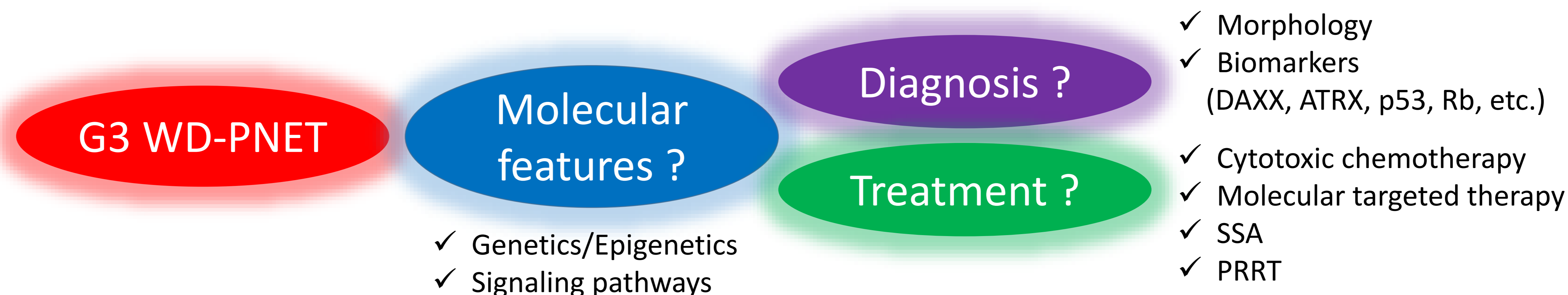


WHO2017 classification

Differentiation	Classification /Grade	Ki67 index	MI /10HPF
Well-differentiated	PNET G1	< 3%	< 2
	PNET G2	3-20%	2-20
Poorly-differentiated	PNET G3	> 20%	> 20
	PNEC G3	> 20%	> 20
	Small cell Large cell		



Tang LH, et al. Clin Cancer Res. 2016



Objective

- 1) Establish a PDX model of G3 WD-PNET
- 2) Investigate the biology and identify new treatments for G3 WD-PNET with this model

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Methods

Patient: 47 y.o. man

Diagnosis:

- ✓ Pancreatic tail neuroendocrine tumor (6.4 cm)
- ✓ Multiple liver metastases (the largest tumor 6.9 cm in S4)
- ✓ Bone metastasis (Th9)

Hormone secretion: None

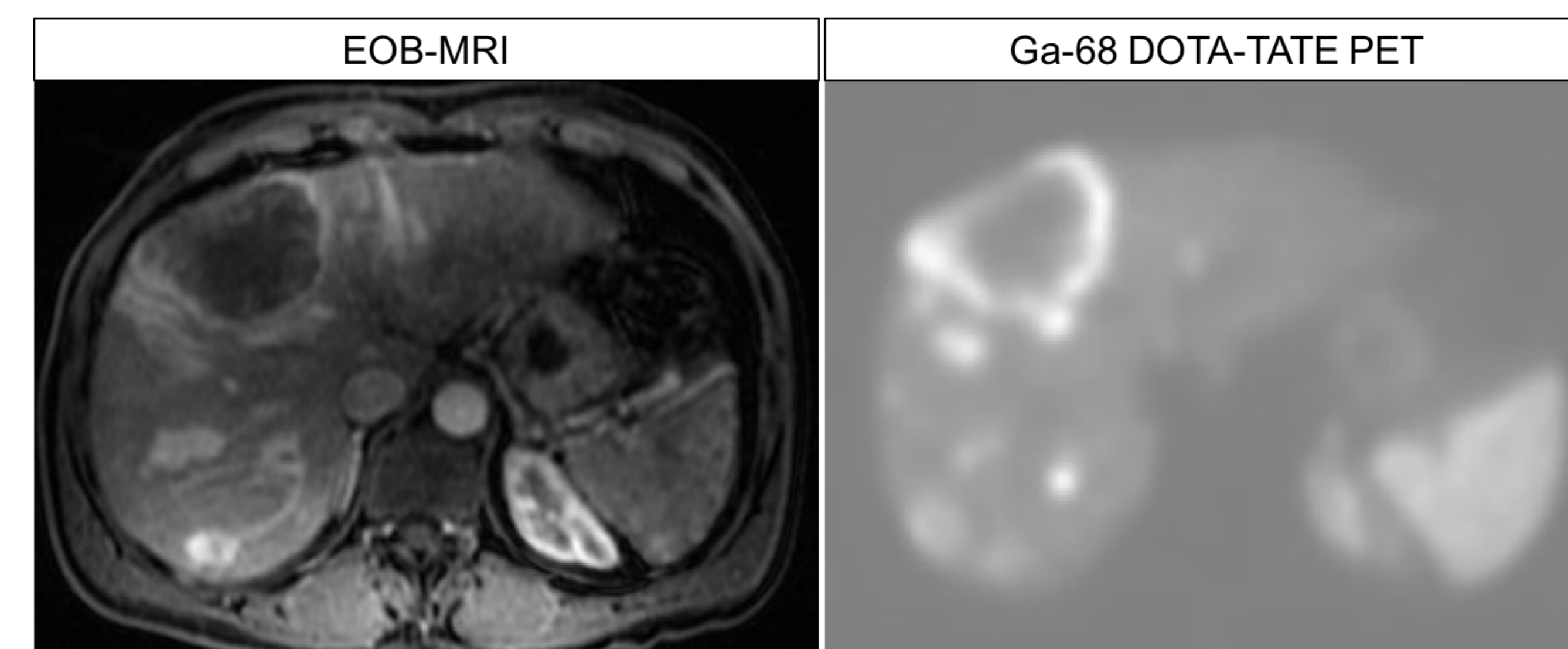
Preoperative treatment:

- ✓ Radiation for Th9
- ✓ Capecitabine/Temozolamide (6 cycles)

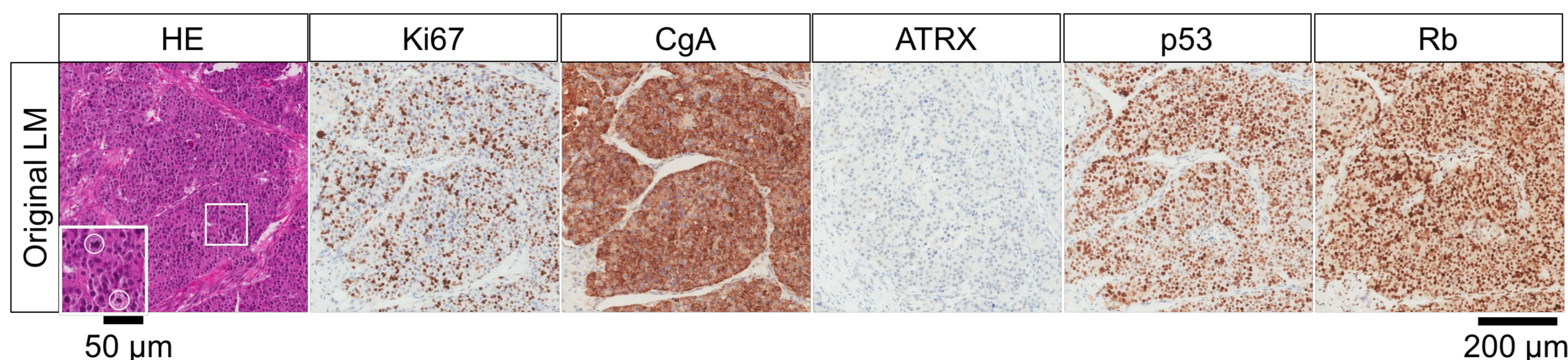
Surgical procedure:

- ✓ Distal pancreatectomy
- ✓ Debulking surgery of liver metastases (including S4 tumor)

Preoperative imaging of the S4 liver metastasis



Pathological findings of S4 liver metastasis: Well-differentiated morphology with 33 mitoses/10 HPFs (white circles indicate mitosis). Immunohistochemical stain result in tumor cells: Ki67 index of 47%, diffuse positive chromogranin A (CgA), loss of ATRX, strong nuclear staining of p53, and intact nuclear staining of Rb.



Targeted gene sequencing of the primary PNET

Gene	Variant
<i>CDKN1B</i>	Deep deletion
<i>MEN1</i>	p.W441fs
<i>TP53</i>	p.G266V
<i>ATM</i>	p.E2272K
<i>TSC2</i>	Exon 33-42 deep deletion

The fragmented tissue of the S4 liver metastasis was immediately implanted subcutaneously into male NSG mice (P0). PDX tumors were passed into the next generation when the tumor volume reached 1,000-2,000 mm³.

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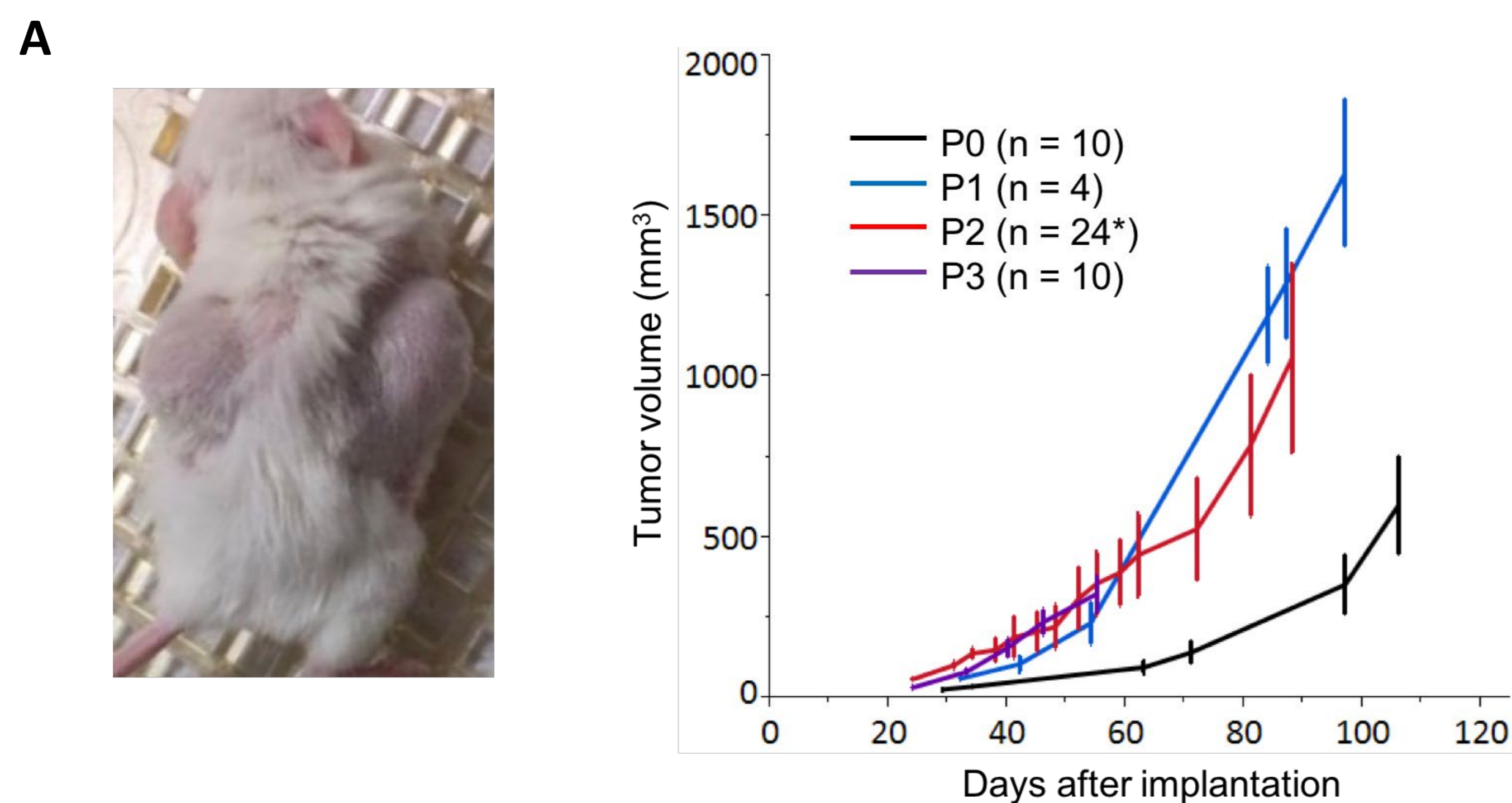
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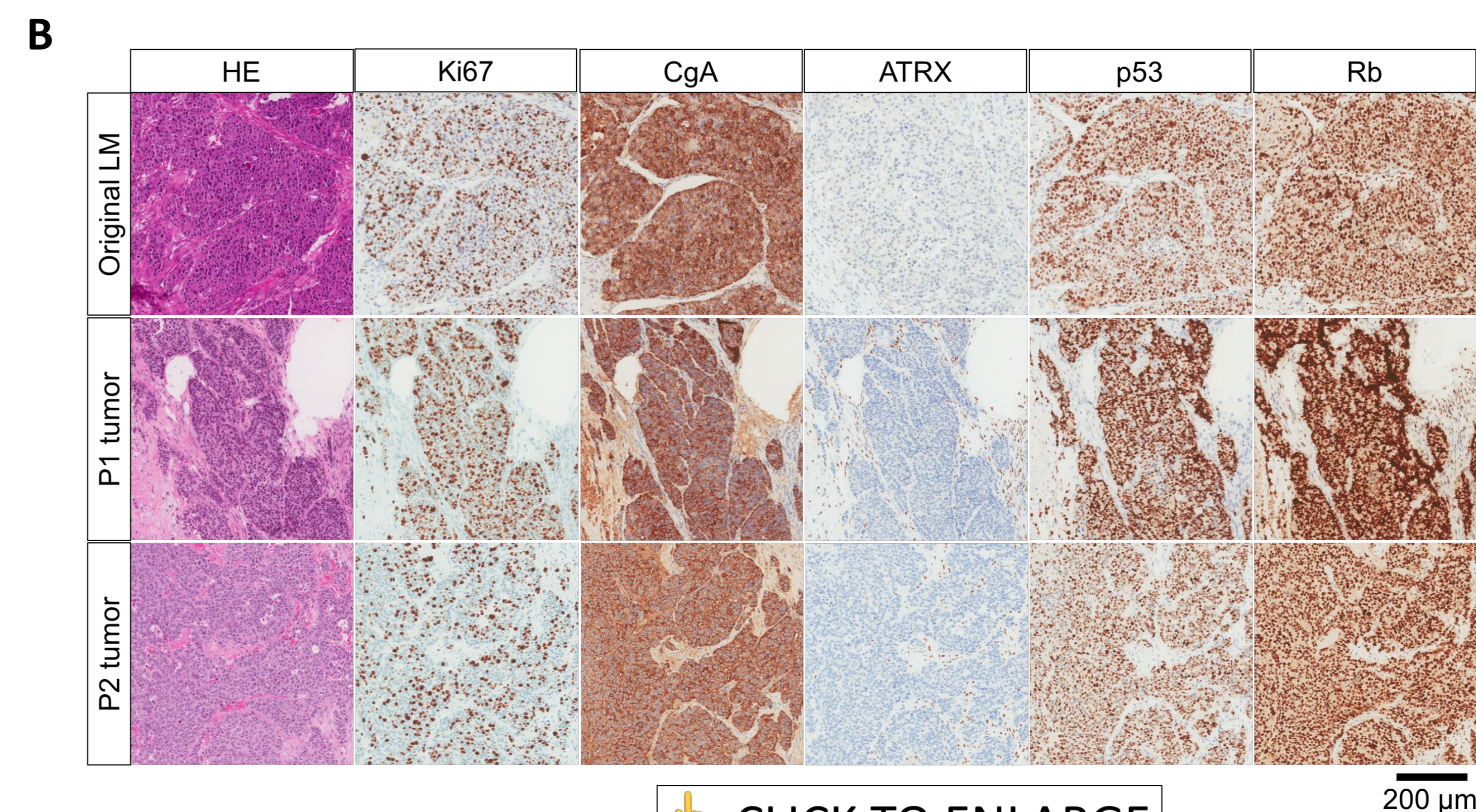
Result 1. PDX tumors exhibit similar molecular features as the original tumor

Fig. 1. (A) All implanted tumors successfully grew as of P3 (100% take rate). **(B)** PDX tumors exhibited similar morphology, proliferation, and staining patterns of CgA, ATRX, p53, and Rb as the original liver metastasis. **(C, D)** Phospho-kinase array showed a similar active kinase profile of the PDX tumor as the original liver metastasis, including enhanced phosphorylation of mTOR and p53 (S392 and S46) and attenuated phosphorylation of p27 (encoded by *CDKN1B* gene).

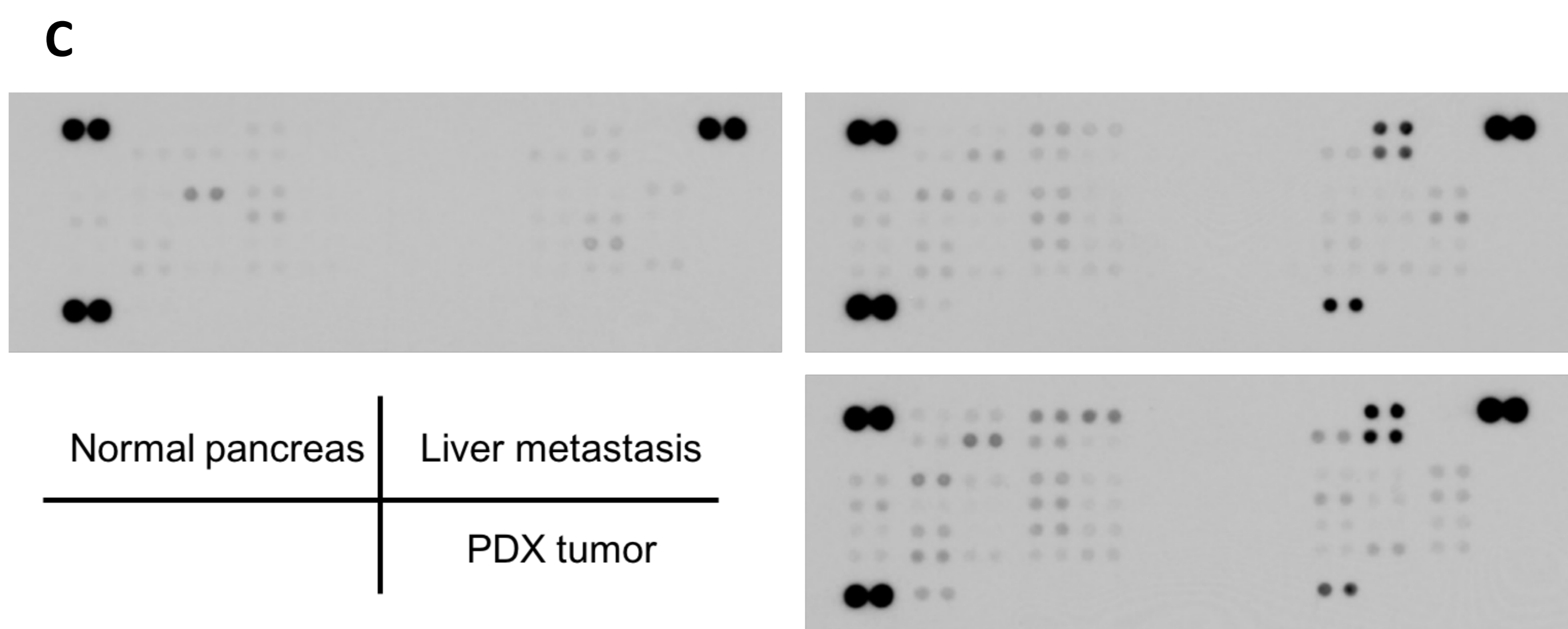
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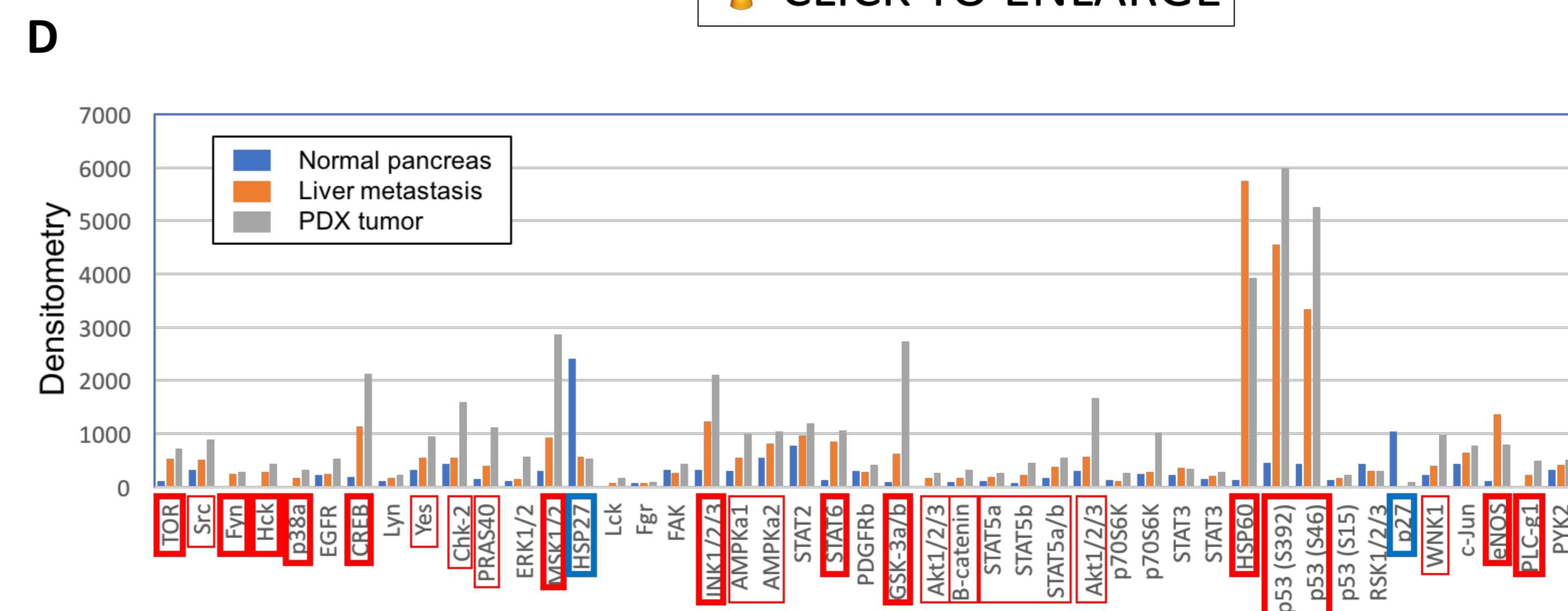
*20 of 24 P2 tumors were used for drug sensitivity test after day 34, and were excluded from the growth curve thereafter.



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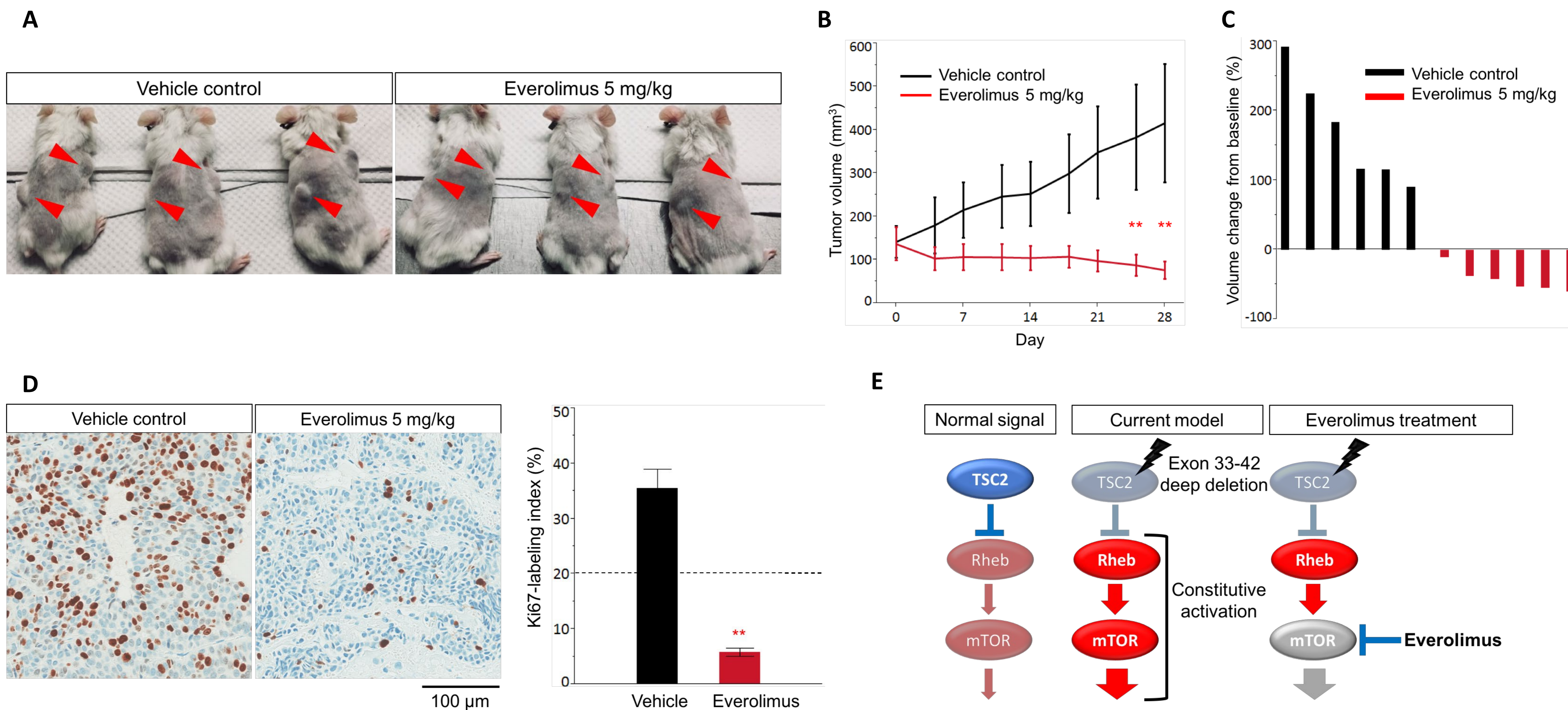
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Result 2. Everolimus suppressed PDX tumor growth.

Fig. 2. PDX tumor-bearing mice were treated once daily with everolimus (5 mg/kg) or vehicle p.o. for 28 days. **(A)** Gross appearance of PDX tumors 28 days after treatment initiation. **(B)** Growth curves of the 2 groups. **(C)** Waterfall plots of each tumor volume on day 28 compared to that on day 0. The median volume changes were 150% (vehicle) and -48% (everolimus), Wilcoxon $P < 0.05$. **(D)** Ki67 positive proliferating tumor cells significantly decreased in the PDX tumors treated by everolimus. **(E)** A possible mechanism for the activation of mTOR signaling and the effect of everolimus in the current PDX model. Data were expressed as the mean and the standard error, and were compared using the Wilcoxon's test. ** $P < 0.05$.



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Conclusion

- ✓ We successfully established a G3 WD-PNET PDX model, which faithfully retains the molecular features of the original patient tumor.
- ✓ This model will facilitate studies of the underlying biology of G3 WD-PNET and help inform prospective clinical trials.

References

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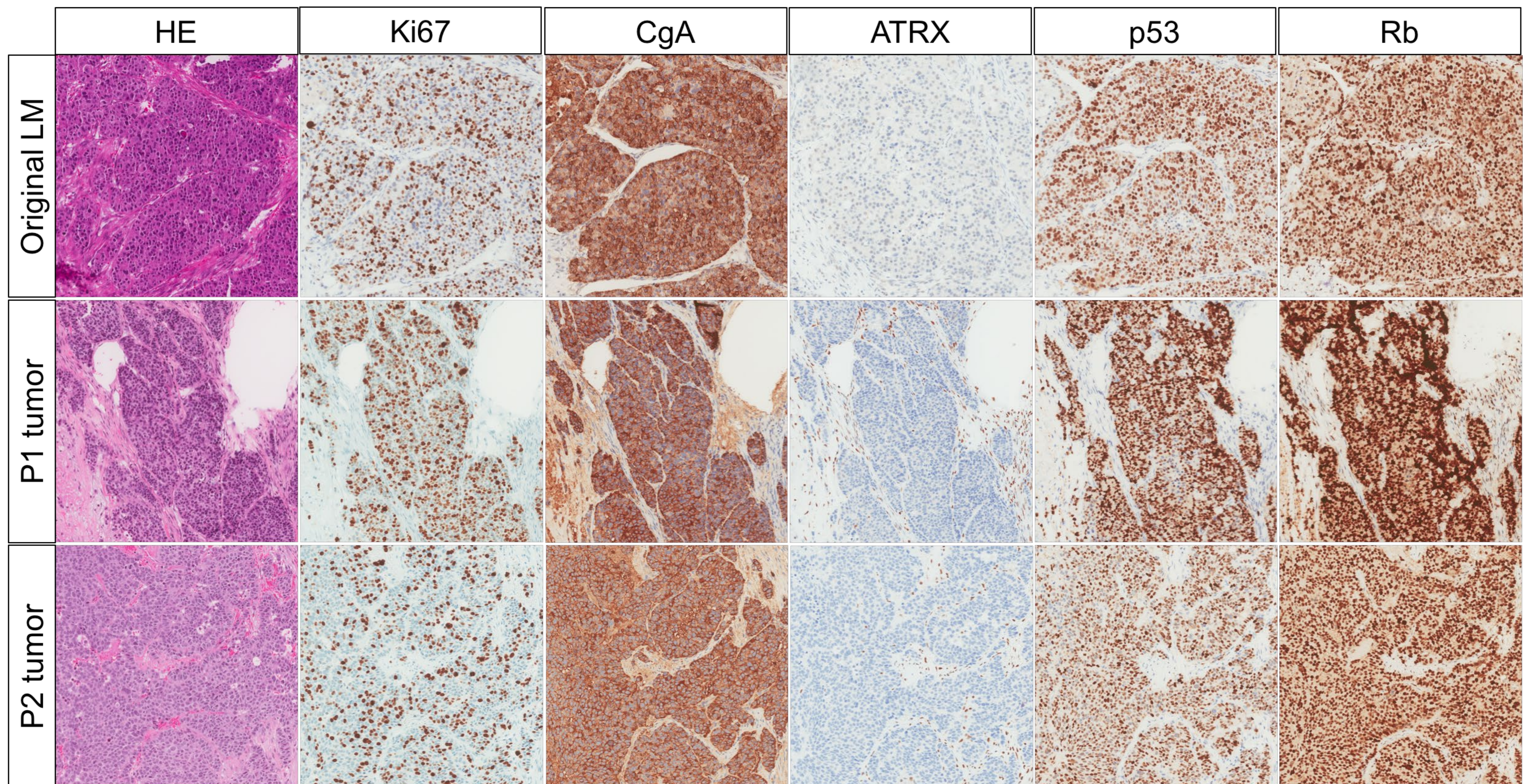
Acknowledgement

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Fig. 1B. PDX tumors exhibited similar morphology, proliferation, and staining patterns of CgA, ATRX, p53, and Rb as the original liver metastasis.



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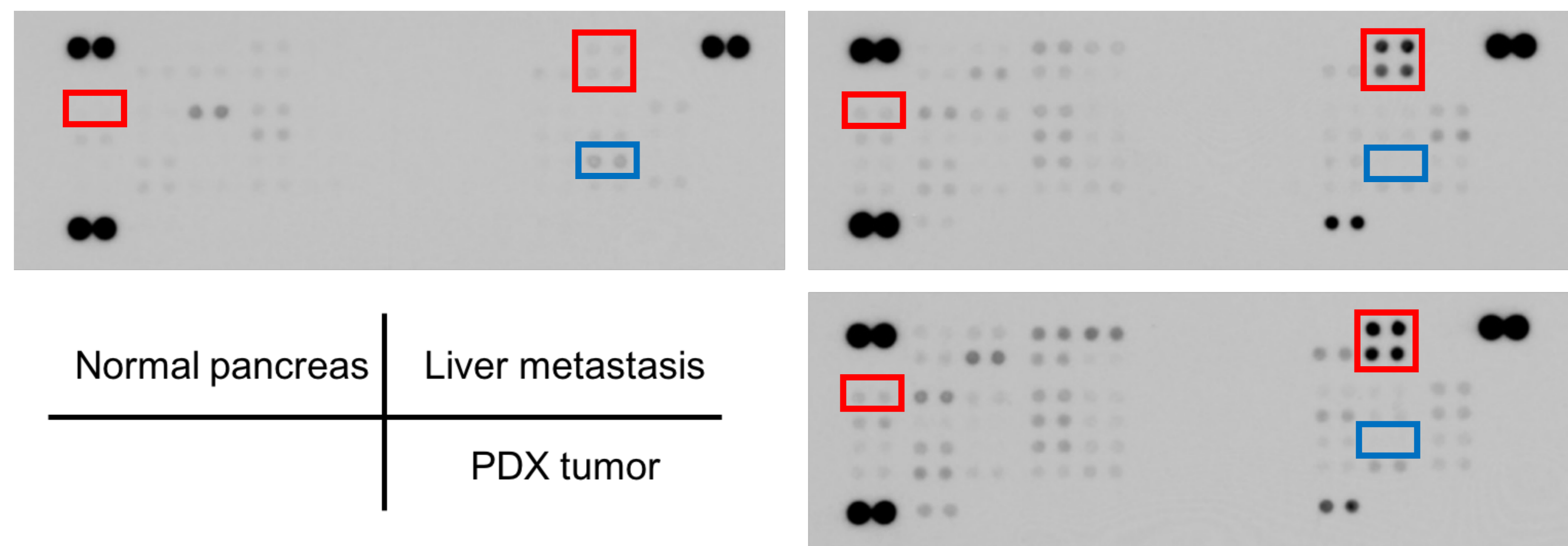
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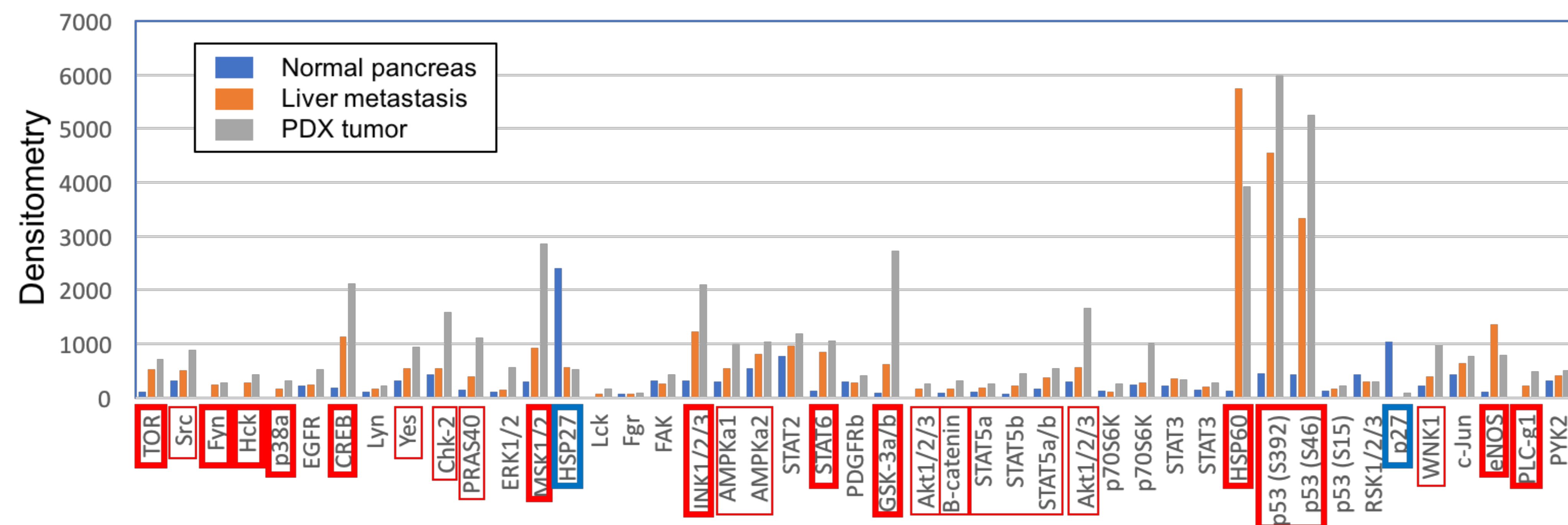
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Fig. 1C, D. Phospho-kinase array showed a similar active kinase profile of the PDX tumor as the original liver metastasis, including enhanced phosphorylation of mTOR and p53 (S392 and S46) and attenuated phosphorylation of p27 (encoded by *CDKN1B* gene).



Ref	p38α	ERK1/2	JNK1/2/3	GSK-3α/β		p53 (S392)		Ref
	EGFR	MSK1/2	AMPKα1	Akt1/2/3	Akt1/2/3	p53 (S46)		
TOR	CREB	HSP27	AMPKα2	β-Catenin	p70S6K	p53 (S15)	c-Jun	
Src	Lyn	Lck	STAT2	STAT5a	p70S6K	RSK1/2/3	eNOS	
Fyn	Yes	Fgr	STAT6	STAT5b	STAT3	p27	PLC-γ1	
Hck	Chk-2	FAK	PDGFRβ	STAT5a/b	STAT3	WNK1	PYK2	
Ref	PRAS40			NC	HSP60			NC

Proteome Profiler™ Array Human Phospho-Kinase Array Kit (ARY003B)
 from R&D SYSTEMS, Minneapolis, MN



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