

# A Novel Mouse Skin Squamous Cell Carcinoma Allograft Model For In Vivo Pharmacological Analysis Of Immunotherapy

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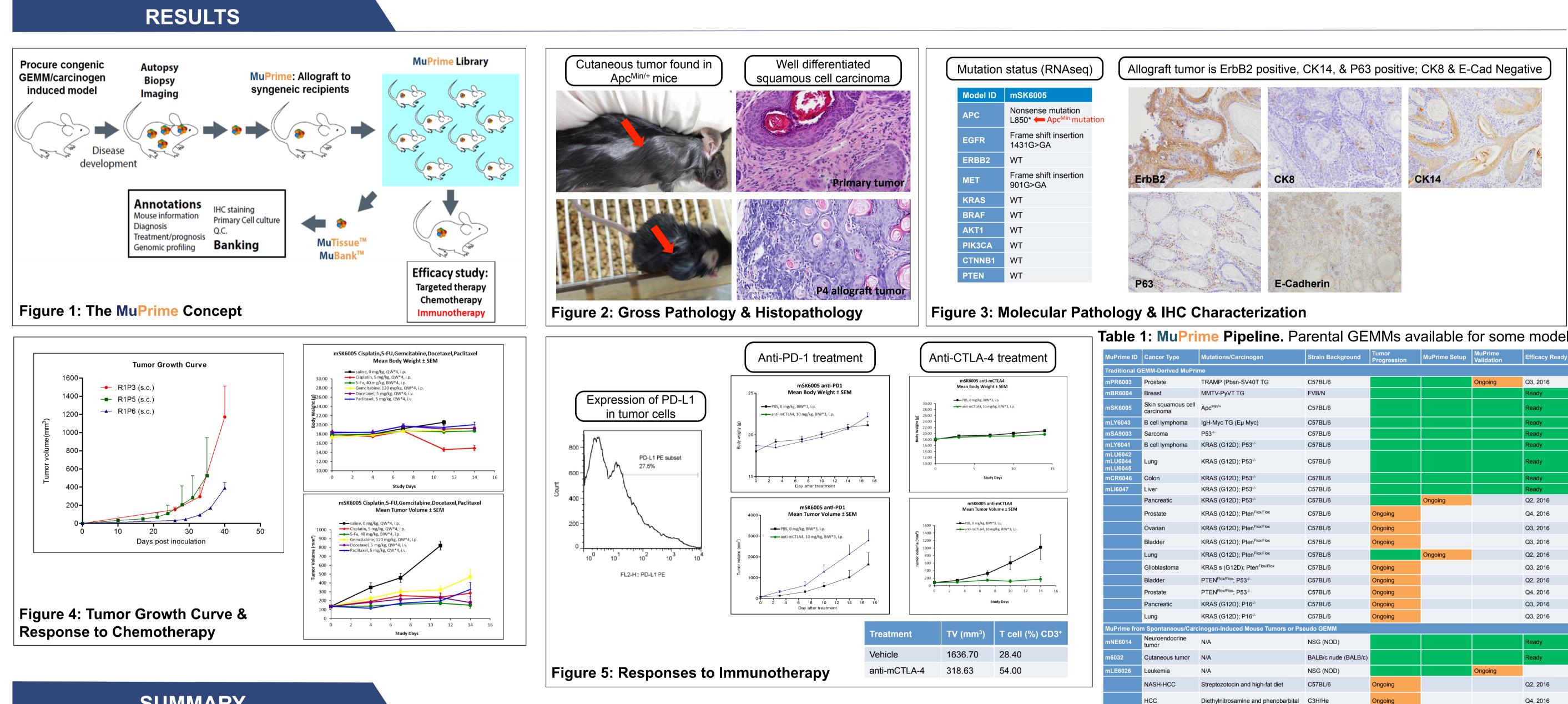
## **ABSTRACT**

The recent clinical success of novel therapeutics blocking the immune checkpoints cytotoxic T-lymphocyte associated antigen 4 (CTLA-4) and programmed cell death 1 (PD-1) has fueled an intense interest around immuno-oncology. However, the lack of relevant animal models is a major bottleneck for understanding the mechanism of action and evaluating the efficacy of such therapeutics. Syngeneic mouse tumor models, despite being widely used as experimental models for efficacy studies, are limited by the restricted number of available models responsive to current checkpoint inhibitors. Genetically engineered mouse models (GEMMs) of human cancers are effective tools for mechanistic analysis, but they are often not suitable for efficacy studies due to usually unsynchronized tumor progression. Syngeneic allografts of spontaneous mouse tumors derived from GEMMs (MuPrime<sup>®</sup>) may be used as a new type of immuno-oncology model with the following advantages: 1) their primary nature of "stem cell disease" and relevant tumor microenvironment mirror patient-derived xenograft (PDX) models; 2) the availability of diverse cancer types and oncogenic drivers deriving from a wide range of GEMMs. We set out to build a library of allografts of spontaneous mouse tumors, including those derived from GEMM, in immunocompetent hosts to support pharmacological investigation, particularly of immuno-oncology agents<sup>1</sup>

It is well documented that heterozygous Apc<sup>Min/+</sup> mice are highly susceptible to developing intestinal adenoma<sup>2,3</sup>. Female Apc<sup>Min/+</sup> mice also occasionally develop mammary squamous cell carcinomas<sup>4</sup>. Recently, we observed a spontaneous cutaneous tumor on the neck of a C57BL/6J Apc<sup>Min/+</sup> mouse. Histopathology suggests the tumor is a well-differentiated skin squamous cell carcinoma. We transplanted the tumor subcutaneously into syngeneic C57BL/6J mice. The allografts grow robustly, they are serially transplantable, and maintain the primary tumor histopathology. Transcriptome sequencing revealed that allografts maintain the original Apc<sup>Min</sup> mutation, express high levels of ErbB2, and present frameshift mutations in both c-Met and EGFR. In vivo pharmacological assessment indicates that the allograft tumors respond to 5-FU, paclitaxel, gemcitabine, docetaxel, and cisplatin chemotherapies. They are also sensitive to an anti-mouse CTLA-4 antibody; this response is associated with an increased number of tumor infiltrating lymphocytes, including T cells. Interesting, although the tumor cells express PD-L1, they are not responsive to PD-1 treatment. In summary, we show that we have established an allograft model suitable for *in vivo* efficacy analysis of immunotherapy using surrogate anti-mouse antibodies.

## References

4. Moser, A.R., Hegge, L.F., Cardiff, R.D. Genetic background affects susceptibility to mammary hyperplasia and carcinomas in Apc<sup>Min/+</sup> Mice. Cancer Research 61, 3480–3485 (2001)



## SUMMARY

- □ We observed a skin squamous carcinoma which developed in a C57BL/6 Apc<sup>Min/+</sup> mouse.
- We have established & characterized the mSK6005 MuPrin studies.
- □ mSK6005 responds well to I/O therapeutics, i.e. anti-mCTLA4 antibody, but not anti-mPD1 antibody.
- □ The response is correlated with increased tumor infiltrating T cells.
- □ To date, 12 MuPrime models have been validated for immuno-oncology efficacy studies

e allograft model, and utilized this model for *in vivo* pharmacological

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### Pipeline. Parental GEMMs available for some models

al SEMM-Derived MuPHered   TRAMP (Pbsn-SV40T TG   C57BL/6   Image: State   TRAMP (Pbsn-SV40T TG   FVB/N   Image: State   Image: State   Image: State   Image: State   FVB/N   Image: State   Image: State   Image: State   Image: State   Image: State   FVB/N   Image: State   <	Ongoing     Image: Comparison of the second se	Q3, 2016     Ready     Ready     Ready     Ready     Ready     Ready     Ready     Ready     Ready     Q2, 2016     Q4, 2016     Q3, 2016
BreastMMTV-PyVT TGFVB/NImage: Comparison of the sector of	Ongoing   Image: Comparison of the compa	ReadyReadyReadyReadyReadyReadyReadyReadyReadyQ2, 2016Q4, 2016Q3, 2016
Skin squamous cell carcinomaApcMin/+C57BL/6Interference setB cell lymphomaIgH-Myc TG (Eµ Myc)C57BL/6Interference SetInterference SetB cell lymphomaP53'-C57BL/6Interference SetInterference SetB cell lymphomaKRAS (G12D); P53'-C57BL/6Interference SetInterference SetLungKRAS (G12D); P53'-C57BL/6Interference SetInterference SetColonKRAS (G12D); P53'-C57BL/6Interference SetInterference SetVerKRAS (G12D); P53'-C57BL/6Interference SetInterference SetPancreaticKRAS (G12D); P53'-C57BL/6Interference SetOngoingProstateKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingInterference SetImage: Set		ReadyReadyReadyReadyReadyReadyReadyQ2, 2016Q4, 2016Q3, 2016
carcinomaApcCS7BL/6carcinoma <thc>carcinoma<!--</td--><td></td><td>Ready     Ready     Ready     Ready     Ready     Ready     Ready     Q2, 2016     Q4, 2016     Q3, 2016</td></thc>		Ready     Ready     Ready     Ready     Ready     Ready     Ready     Q2, 2016     Q4, 2016     Q3, 2016
SarcomaP53 <sup>-/-</sup> C57BL/6IndexIndexB cell lymphomaKRAS (G12D); P53 <sup>-/-</sup> C57BL/6IndexIndexLungKRAS (G12D); P53 <sup>-/-</sup> C57BL/6IndexIndexColonKRAS (G12D); P53 <sup>-/-</sup> C57BL/6IndexIndexLiverKRAS (G12D); P53 <sup>-/-</sup> C57BL/6IndexIndexPancreaticKRAS (G12D); P53 <sup>-/-</sup> C57BL/6IndexOngoingProstateKRAS (G12D); Ptor <sup>Flox/Flox</sup> C57BL/6OngoingIndexVarianKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingIndexIungKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6IndexOngoingIungKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6IndexIndexIungKRAS (G12D); Pten <sup>Flox/Flox</sup> KKIndex		ReadyReadyReadyReadyReadyQ2, 2016Q4, 2016Q3, 2016
B cell lymphomaKRAS (G12D); P53-/-C57BL/6Image: Constant of the sector		Ready     Ready     Ready     Ready     Q2, 2016     Q4, 2016     Q3, 2016
LungKRAS (G12D); P53'-C57BL/6InterpretationColonKRAS (G12D); P53'-C57BL/6InterpretationLiverKRAS (G12D); P53'-C57BL/6InterpretationPancreaticKRAS (G12D); P53'-C57BL/6OngoingProstateKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingPortateKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingImage: State of the		Ready     Ready     Ready     Q2, 2016     Q4, 2016     Q3, 2016
A definitionKRAS (G12D); P53-/-C57BL/6Image: Constant of the second of		Ready       Ready       Q2, 2016       Q4, 2016       Q3, 2016
LiverKRAS (G12D); P53-/-C57BL/6Image: C57BL/6OngoingPancreaticKRAS (G12D); P53-/-C57BL/6OngoingOngoingProstateKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingImage: C57BL/6OvarianKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingImage: C57BL/6BladderKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingImage: C57BL/6LungKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingImage: C57BL/6GlioblastomaKRAS s (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingImage: C57BL/6		Ready       Q2, 2016       Q4, 2016       Q3, 2016
PancreaticKRAS (G12D); P53-/-C57BL/6OngoingProstateKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingVarianKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingBladderKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingLungKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingGlioblastomaKRAS s (G12D); Pten <sup>Flox/Flox</sup> C57BL/6Ongoing		Q2, 2016 Q4, 2016 Q3, 2016
ProstateKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingOvarianKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingBladderKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingLungKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingGlioblastomaKRAS s (G12D); Pten <sup>Flox/Flox</sup> C57BL/6Ongoing		Q4, 2016 Q3, 2016
OvarianKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingBladderKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingLungKRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6OngoingGlioblastomaKRAS s (G12D); Pten <sup>Flox/Flox</sup> C57BL/6Ongoing		Q3, 2016
Bladder   KRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6   Ongoing     Lung   KRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6   Ongoing     Glioblastoma   KRAS s (G12D); Pten <sup>Flox/Flox</sup> C57BL/6   Ongoing		
Lung KRAS (G12D); Pten <sup>Flox/Flox</sup> C57BL/6 Ongoing   Glioblastoma KRAS s (G12D); Pten <sup>Flox/Flox</sup> C57BL/6 Ongoing		02 0040
Glioblastoma KRAS s (G12D); Pten <sup>Flox/Flox</sup> C57BL/6 Ongoing		Q3, 2016
Glioblastoma KRAS s (G12D); Pten <sup>Flox/Flox</sup> C57BL/6 Ongoing		Q2, 2016
		Q3, 2016
Bladder PTEN <sup>Flox/Flox</sup> ; P53 <sup>-/-</sup> C57BL/6 Ongoing		Q2, 2016
Prostate PTEN <sup>Flox/Flox</sup> ; P53 <sup>-/-</sup> C57BL/6 Ongoing		Q4, 2016
Pancreatic KRAS (G12D); P16 <sup>-/-</sup> C57BL/6 Ongoing		Q3, 2016
Lung KRAS (G12D); P16 <sup>-/-</sup> C57BL/6 Ongoing		Q3, 2016
from Spontaneous/Carcinogen-Induced Mouse Tumors or Pseudo GEMM		
Neuroendocrine tumor N/A NSG (NOD)		Ready
Cutaneous tumor N/A BALB/c nude (BALB/c)		Ready
Leukemia N/A NSG (NOD)	Ongoing	
NASH-HCC     Streptozotocin and high-fat diet     C57BL/6     Ongoing		Q2, 2016
HCC Diethylnitrosamine and phenobarbital C3H/He Ongoing		Q4, 2016
Lung Urethane BALB/c; C3H/He; A/J	Ongoing	Q2, 2016
AML NRAS (G12D); AML1-ETO* C57BL/6	Ongoing	Q2, 2016
AML NRAS (G12D); MLL-ENL* C57BL/6	Ongoing	Q2, 2016
Derived from Non-Germline GEMM using CRISPR-Cas9 <i>In Vivo</i> Gene Editing		
Bladder P53 cKO; PTEN cKO C57BL/6 Ongoing		
Breast P53 cKO; BRCA1 cKO; MMTV-Cre C57BL/6 Ongoing		
Breast P53 cKO; SMAD4 cKO; CDH1 cKO; MMTV-Cre C57BL/6 Ongoing		
Gastric P53 cKO; SMAD4 cKO; CDH1 cKO C57BL/6 Ongoing		
Lung KRAS (G12D); P53 cKO; LKB1 cKO C57BL/6 Ongoing		
Pancreatic KRAS (G12D); P53 (R172H); Pdx1-Cre C57BL/6 Ongoing		Q3, 2016
Pancreatic KRAS (G12D); SMAD5 cKO; Pdx1-Cre C57BL/6 Ongoing		
Prostate BRAF (V600E); PTEN cKO; Pbsn-Cre C57BL/6 Ongoing		Q4, 2016
Prostate PTEN cKO; Smad4 cKO; Pbsn-Cre C57BL/6 Ongoing		Q4, 2016
Ovarian KRAS (G12D); PTEN cKO C57BL/6 Ongoing		
Medullablastoma PTCH1 cKO; P53 cKO C57BL/6 Ongoing		Q3, 2016

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<sup>3.</sup> Su, L.K., et al. Multiple intestinal neoplasia caused by a mutation in the murine homolog of the APC gene. Science 256, 668-670 (1992).