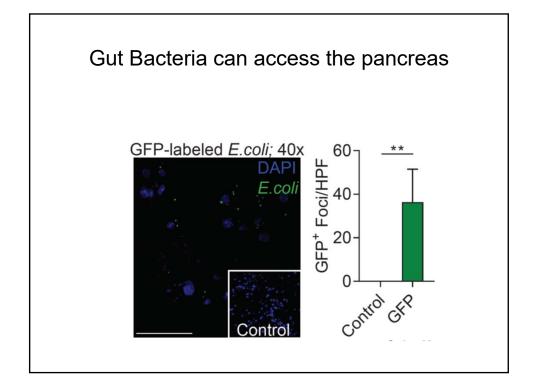


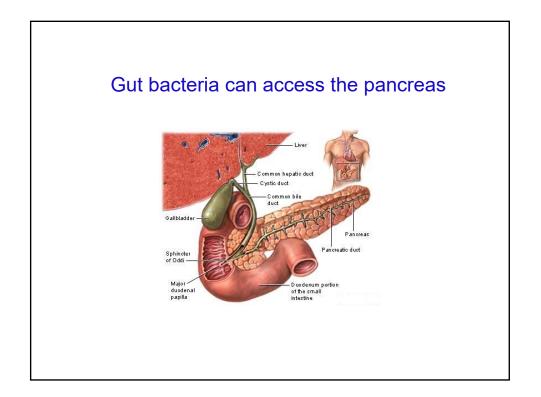
## **Hypothesis**

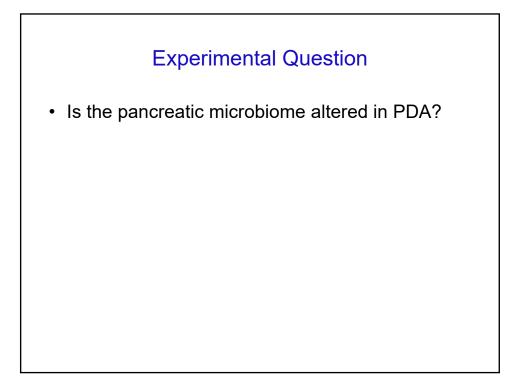
In genetically susceptible hosts the microbiome alters the risk of pancreatic carcinogenesis and can promote aggressive disease biology

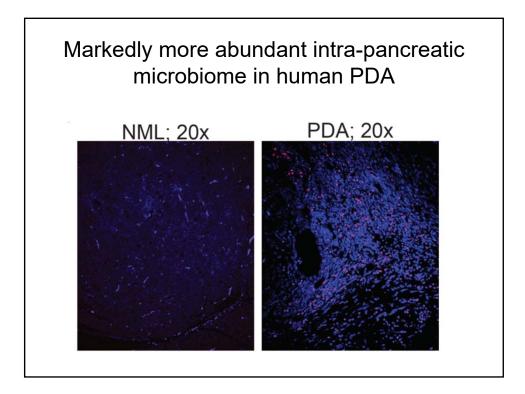
## Question

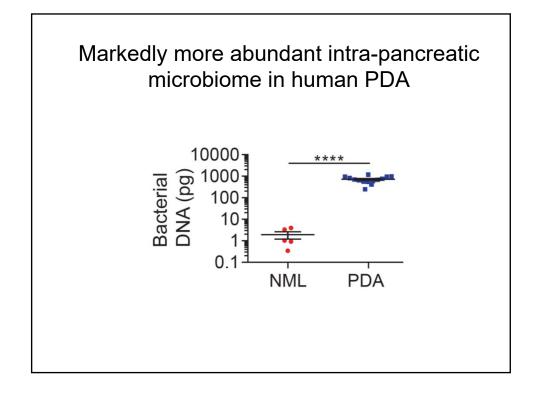
Can Gut Bacteria Access the Pancreas?

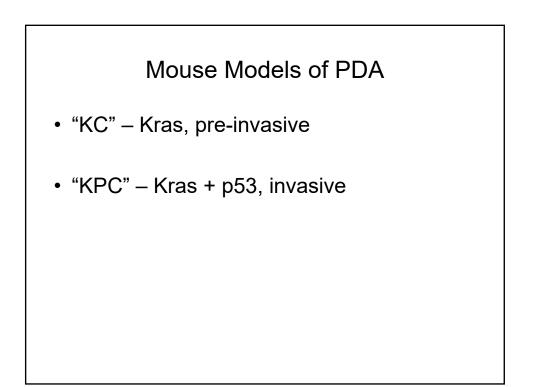


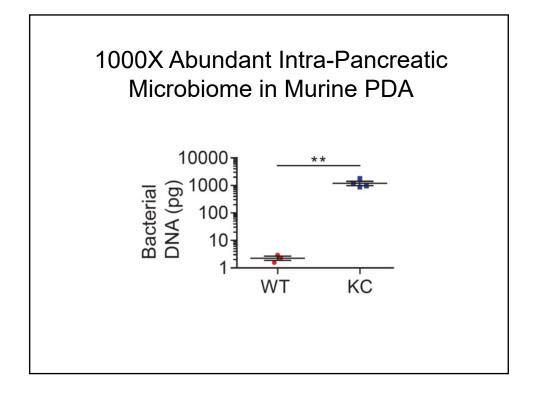


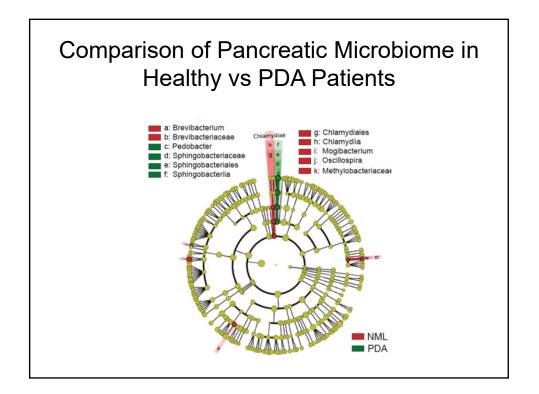


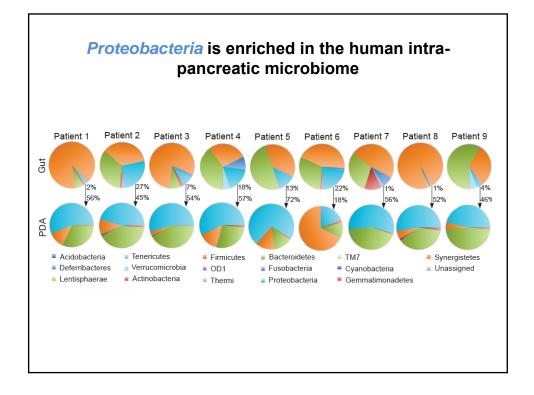


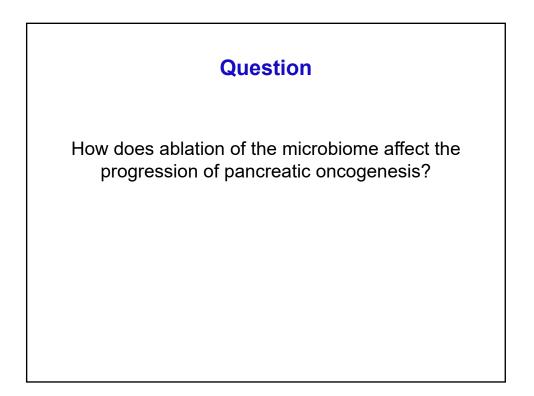


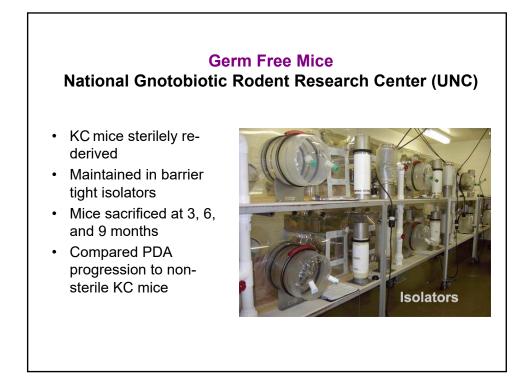


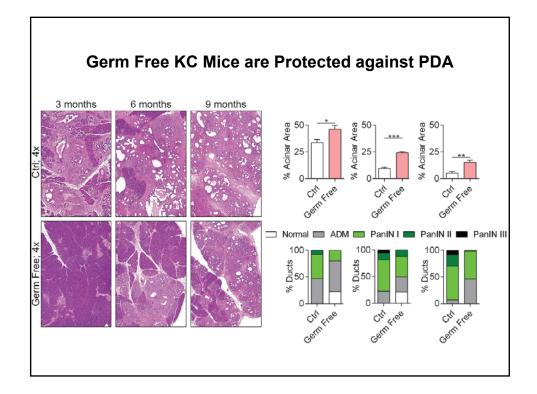


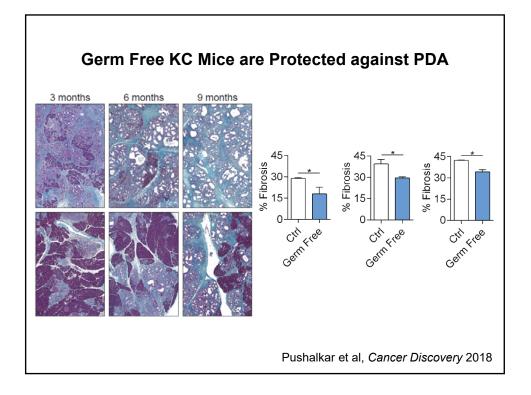


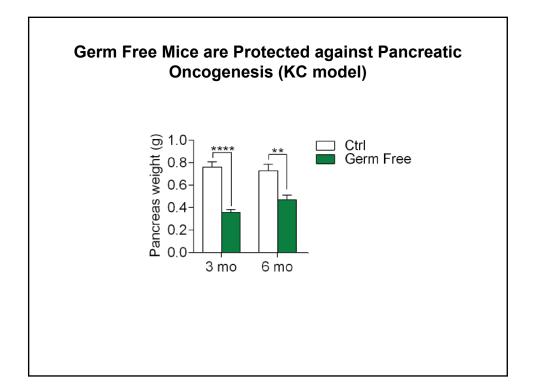


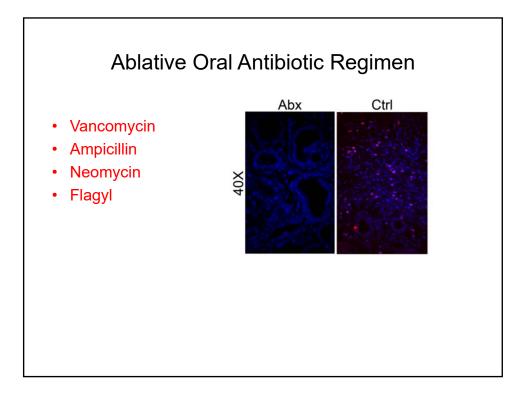


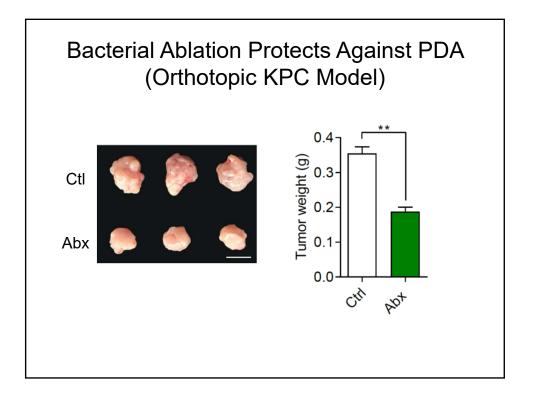


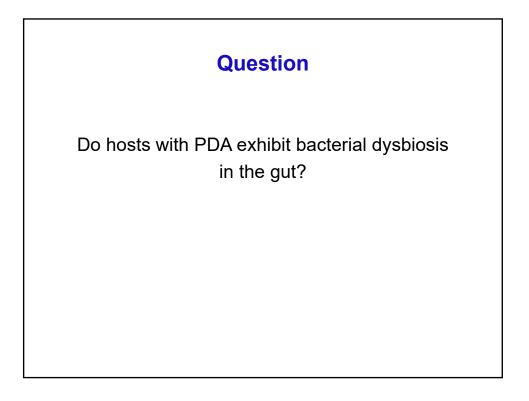


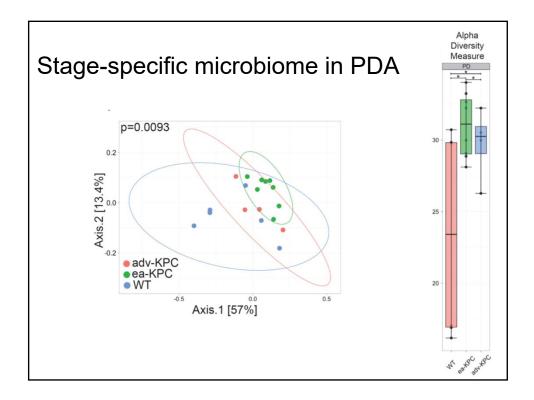


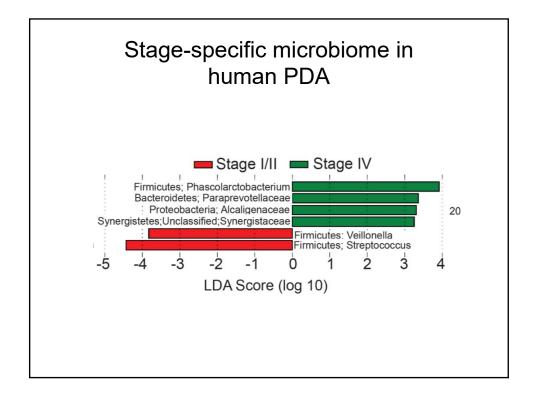


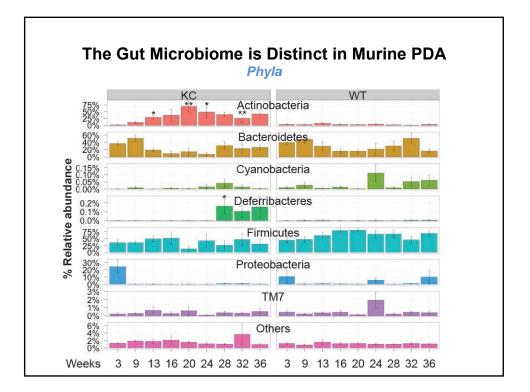


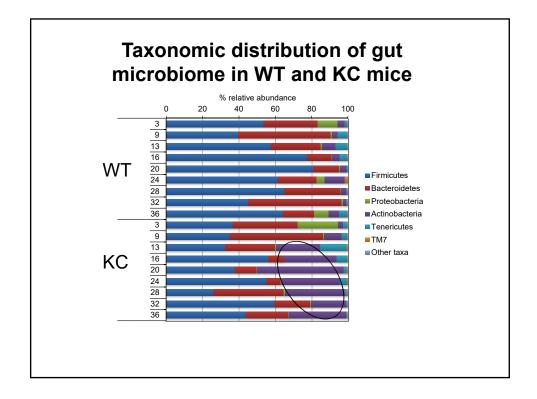


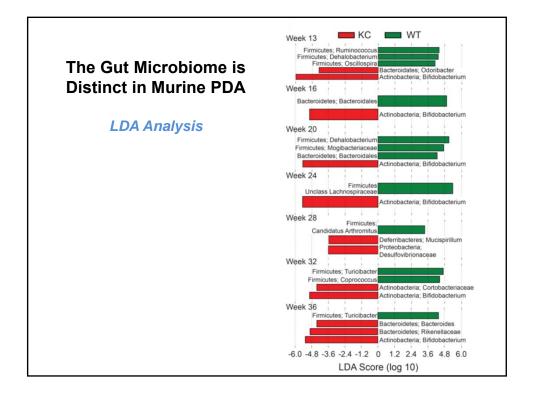






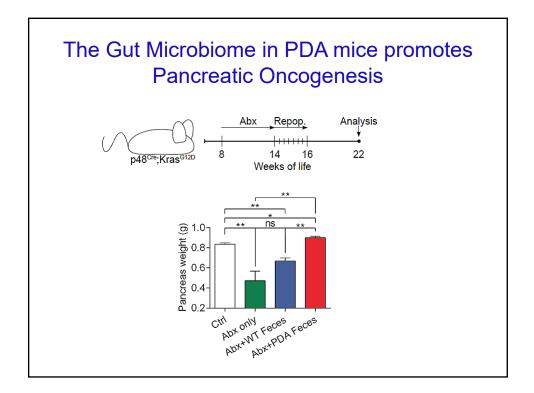


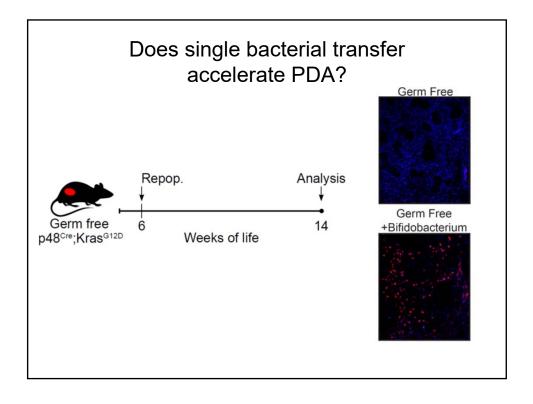


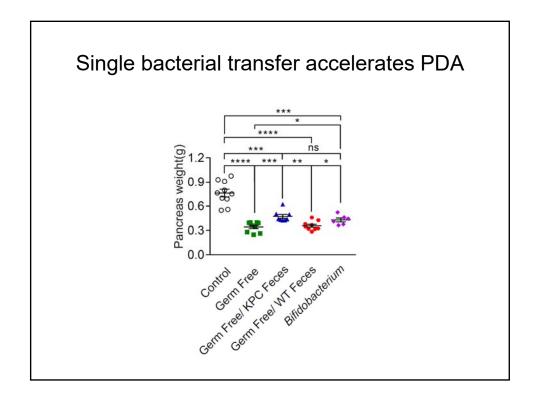


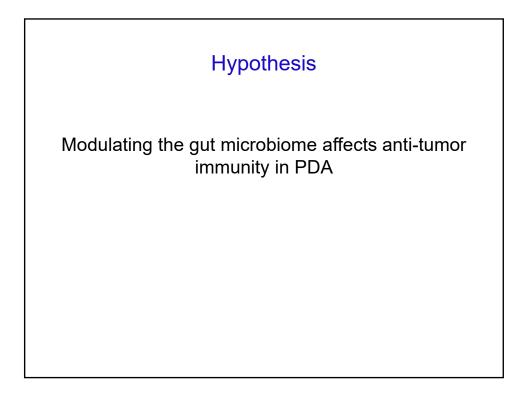
## Question

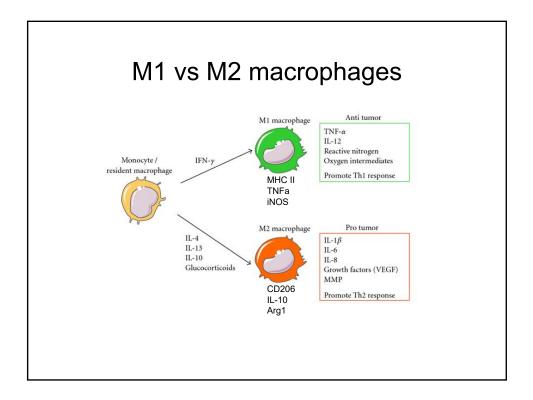
Does direct gut repopulation with "Good" or "Bad" Microbes alter PDA Progression?

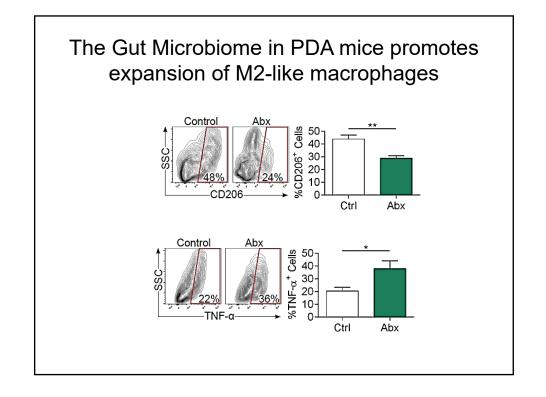


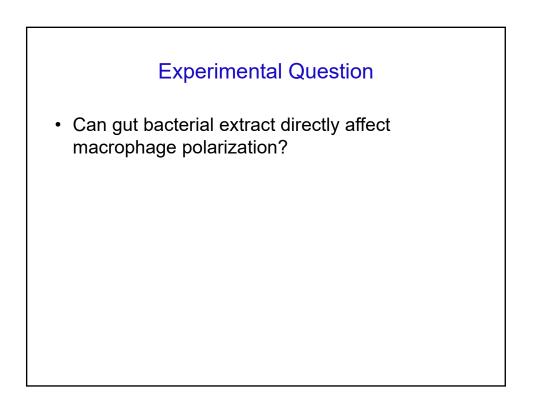


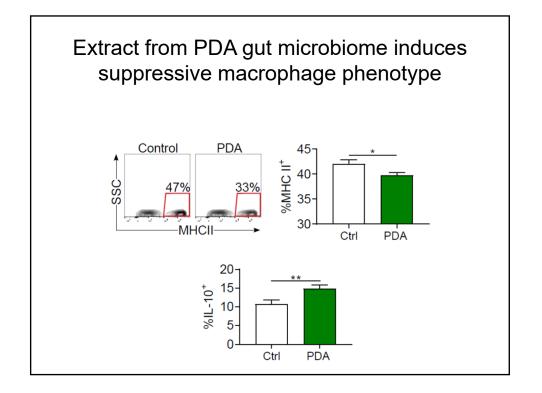


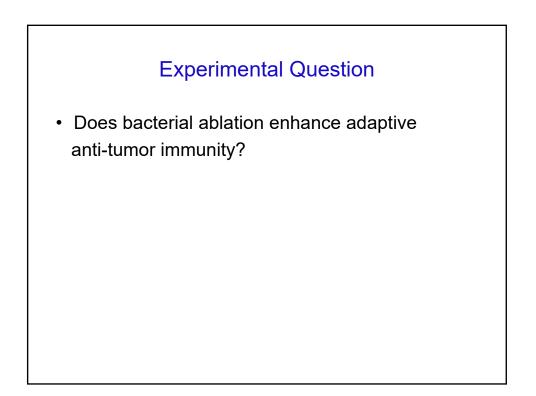


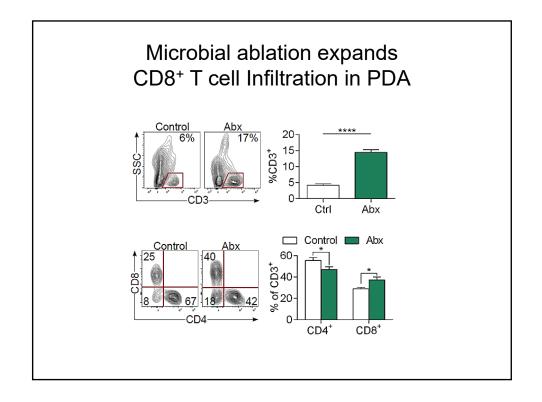


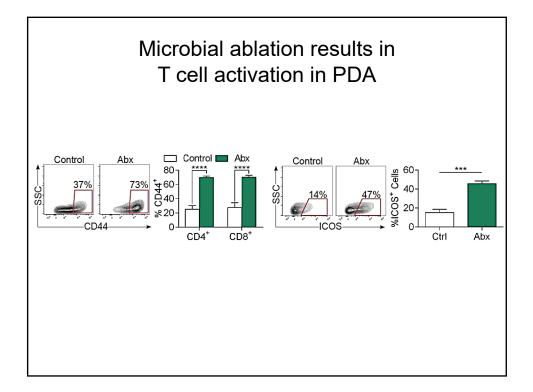


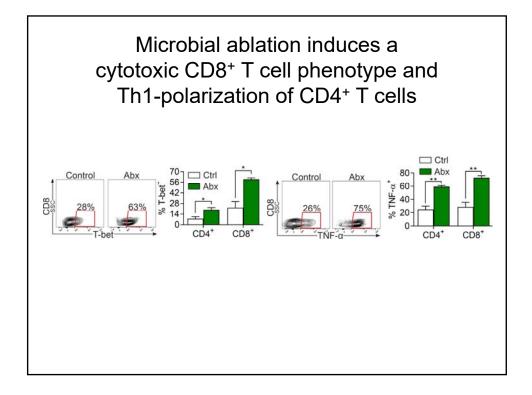


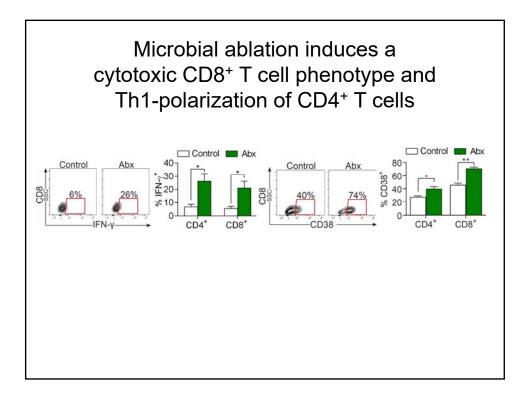


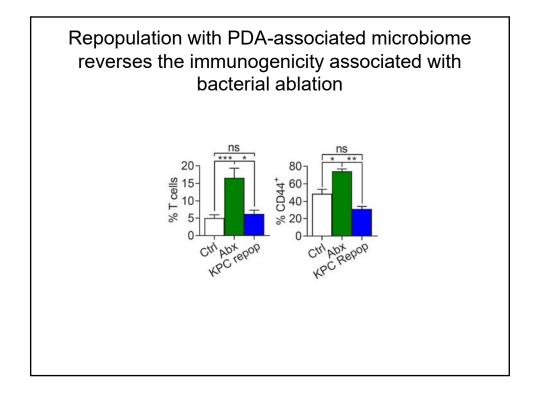


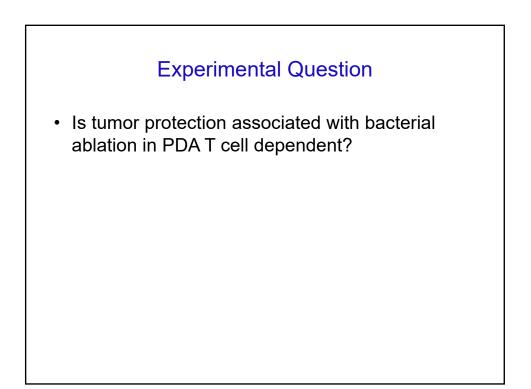


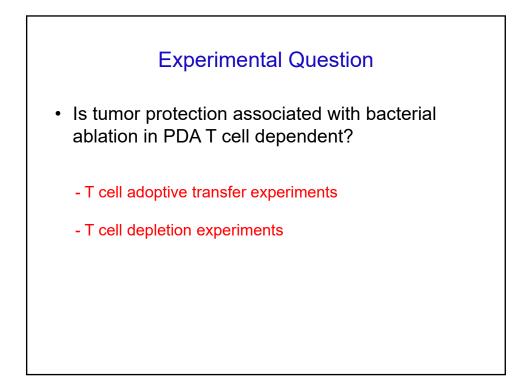


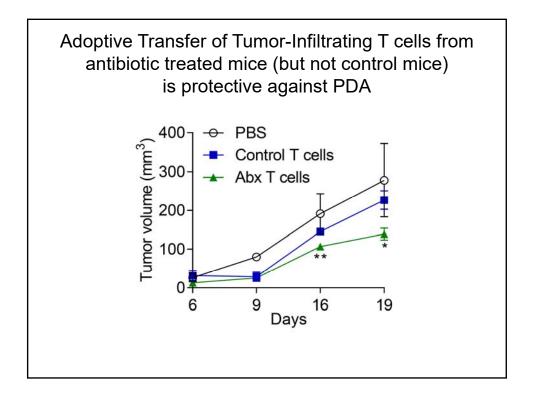


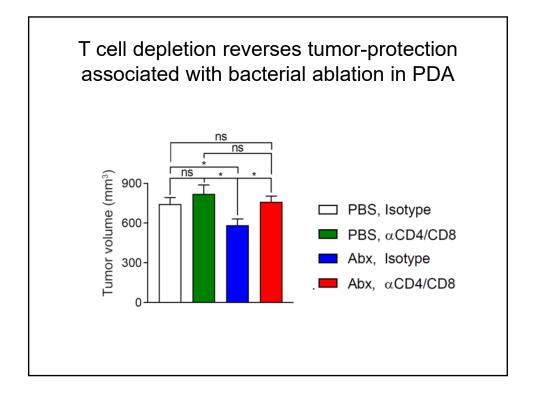


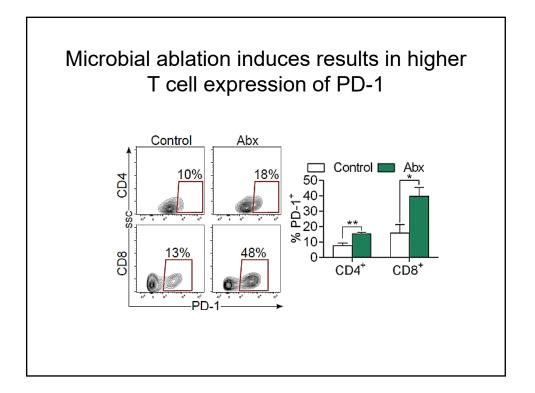


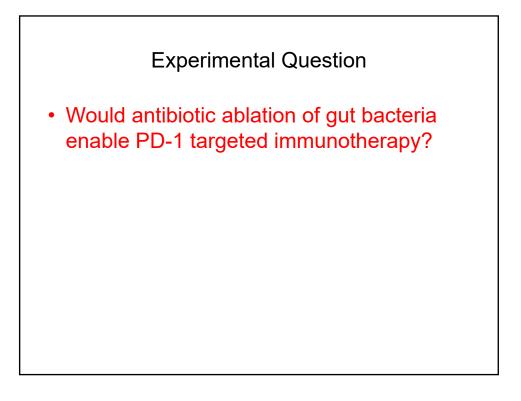


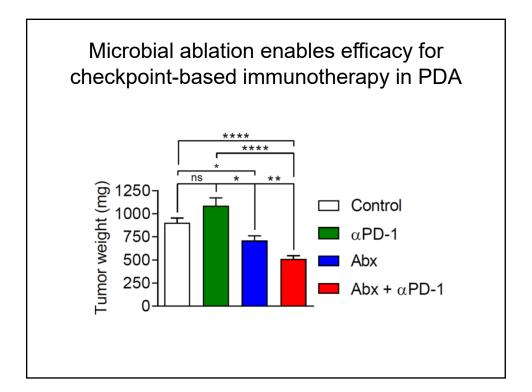


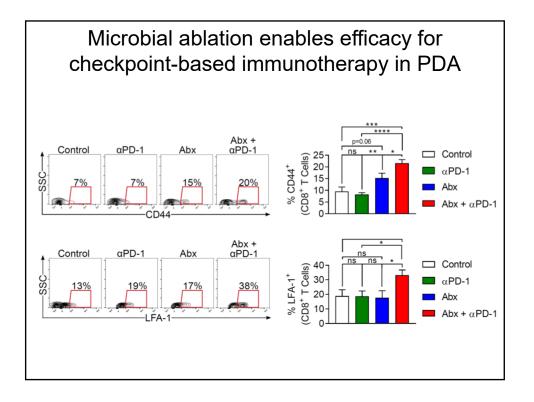


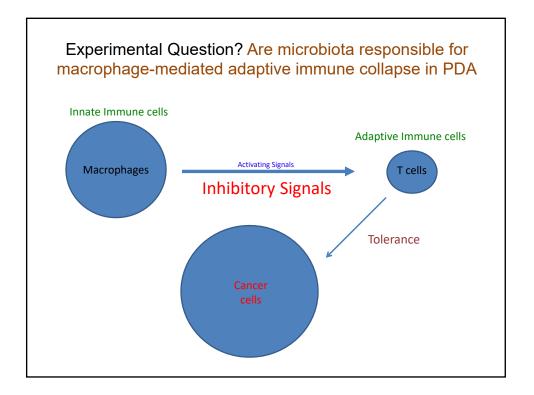


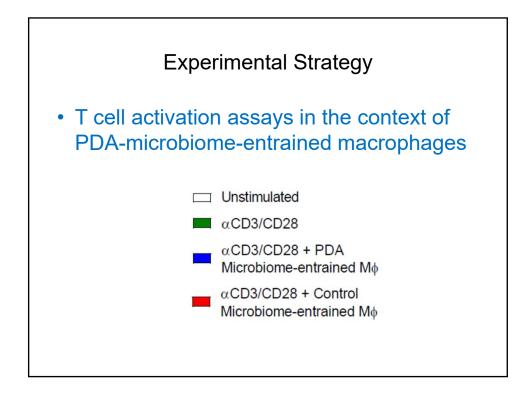


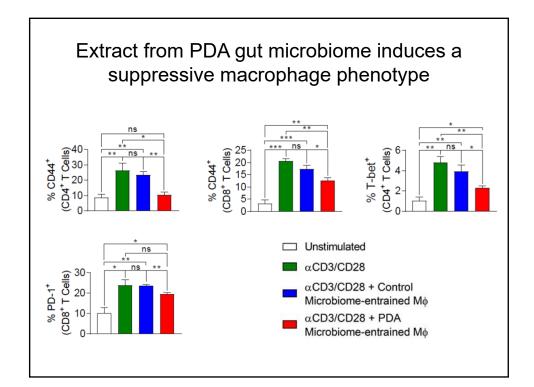


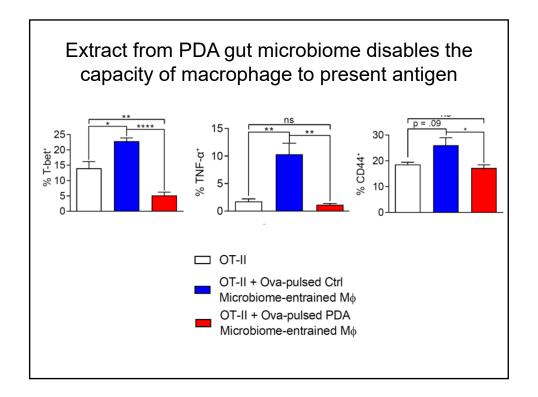


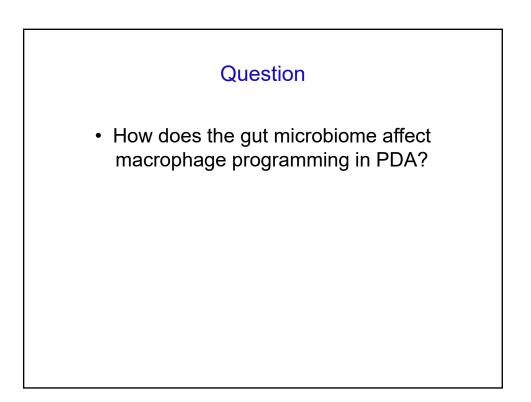


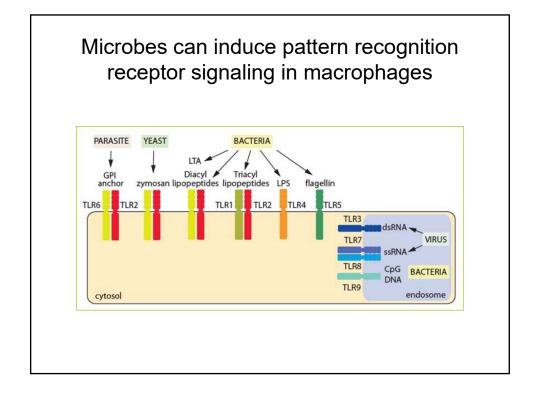


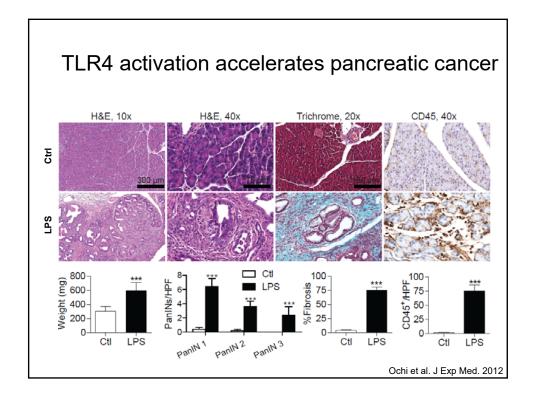


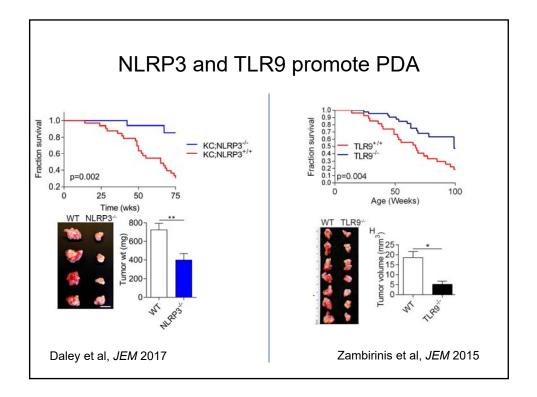


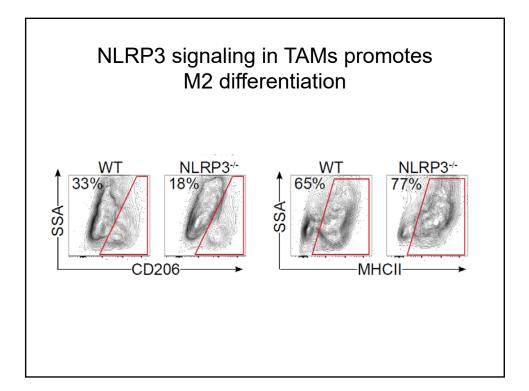


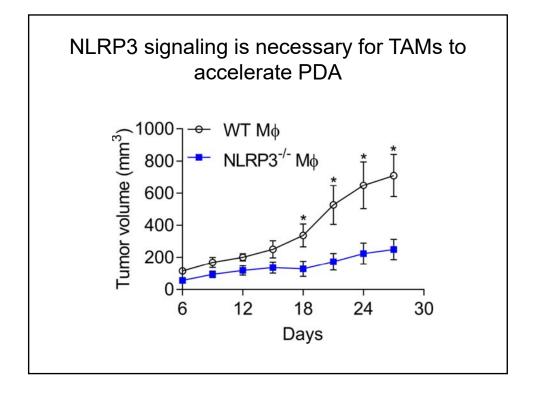


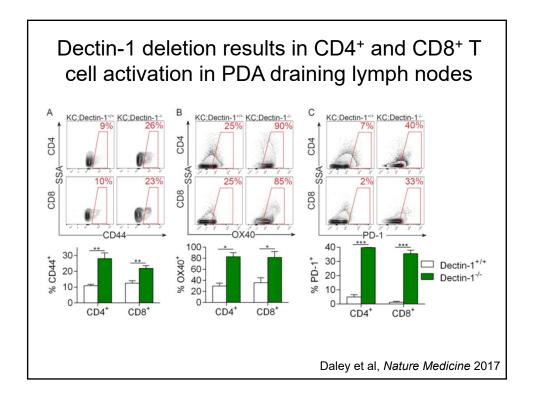


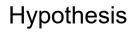




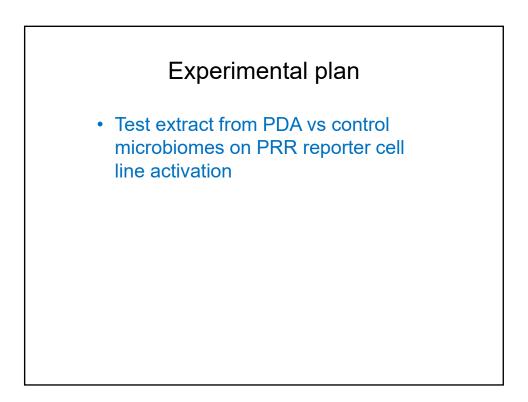


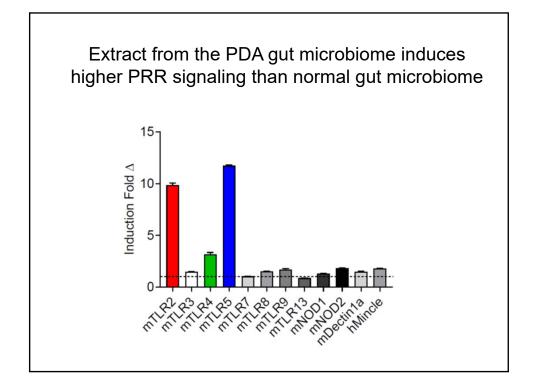


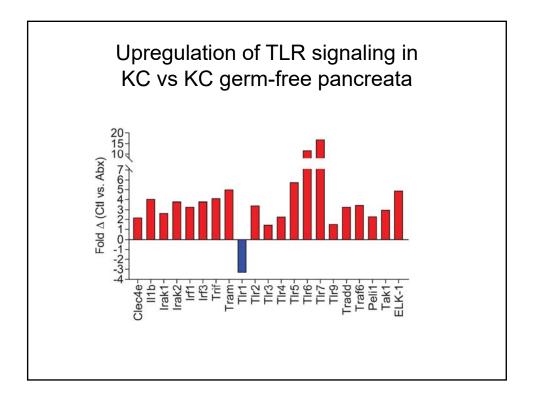


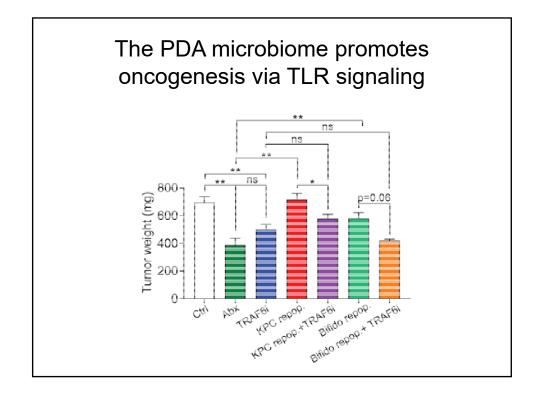


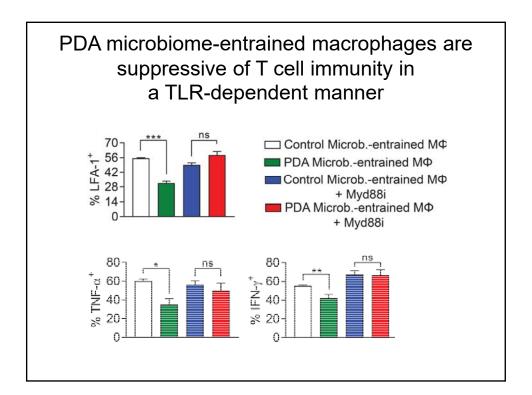
• The PDA microbiome induces crippling immune-suppression in PDA via diverse pattern recognition receptor (PRR) activation

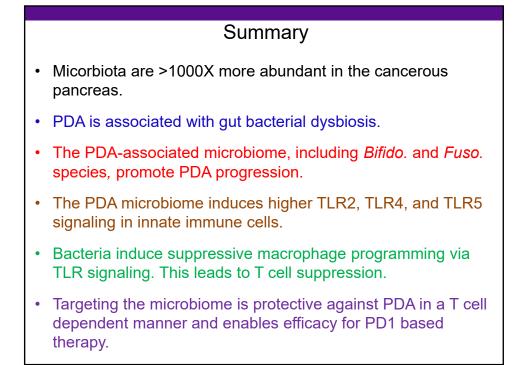


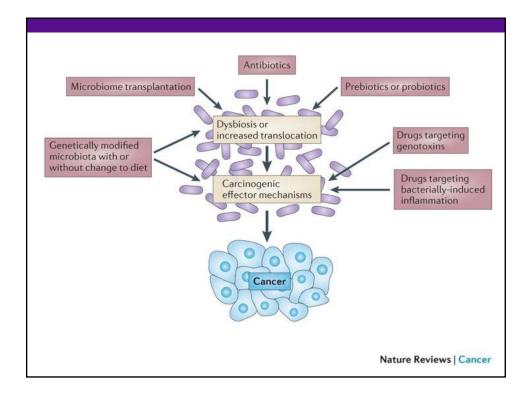


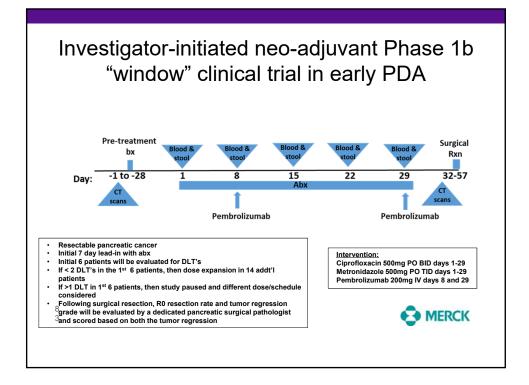


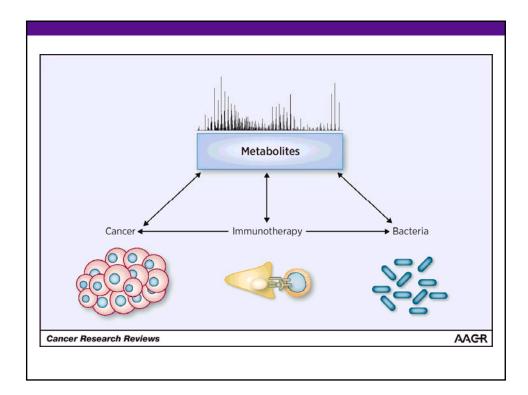


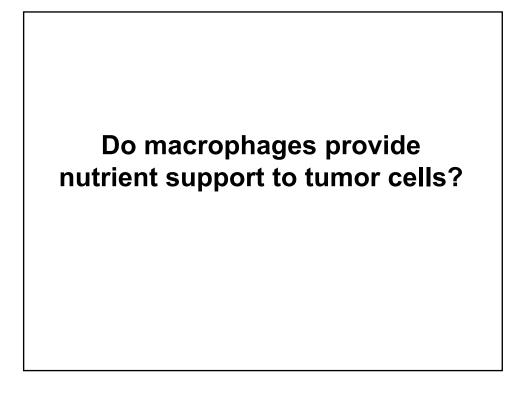


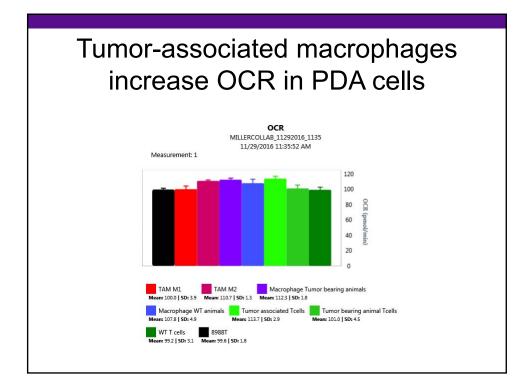


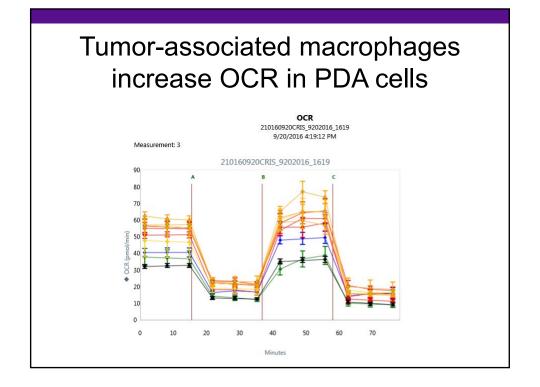


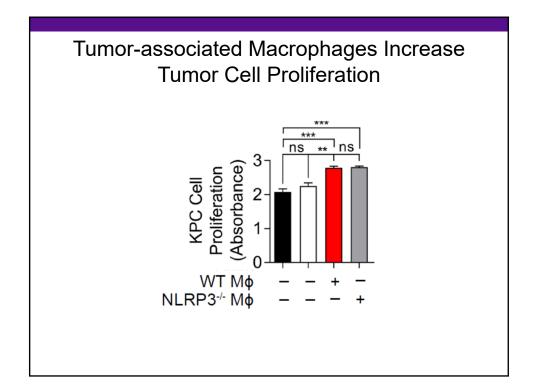


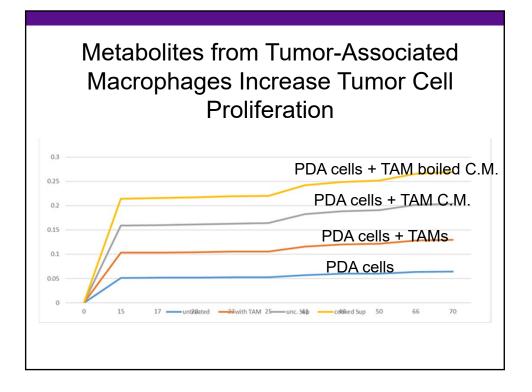












TAMs Produce High Levels of Urea Cycle Intermediates				
	TAM CM vs PDAC CM		TAM CM vs T-cell media	
Metabolite	Fold Change	p-value	Fold Change	p-valu
ADP	8.51	0.0051	1.43	0.036
dAMP	6.12	0.0296	7.24	0.026
ornithine	1.74	0.0038	6.62	0.003
sarcosine (Alanine)	1.62	0.0010	1.20	0.02
Urea	1.49	0.0032	1.44	0.00
alanine	1.44	0.0009	1.24	0.00
L-arginino-succinate	1.23	0.0092	1.70	0.00
arginine	0.86	0.0022	0.77	0.000
creatine	0.85	0.0128	0.73	0.004
dimethylglycine	0.53	0.0592	0.64	0.017
fructose-6-phosphate	0.43	0.0024	0.68	0.068
hypoxanthine	0.39	0.0002	0.39	0.019
inosine	0.33	0.0011	0.40	0.004
D-sedoheptulose-1-7-phosphate	0.30	0.0015	0.51	0.032
	0.16	0.0036	0.44	0.008

