1	Correlation between green tea polyphenols regulating intestinal
2	bacteriophage and flora diversity in SPF mice
3	Sashuang Dong <sup>1</sup> , Zilong Xin <sup>2</sup> , Wencan He <sup>3</sup> , Yuling Zhang <sup>1</sup> , Jiaqi Xiong <sup>1</sup> , Jie
4	Wang <sup>1</sup> , Zhenlin Liao <sup>1</sup> , Li Wang <sup>1</sup> , Qingping Zhong <sup>1</sup> , Hong Wei <sup>3,4*</sup> , Xiang Fang <sup>1*</sup>
5	Affiliation:
6	<sup>1</sup> College of Food Science, South China Agricultural University, Guangzhou,
7	Guangdong, 510630, P. R. China
8	<sup>2</sup> College of Engineering, South China Agricultural University, Guangzhou,
9	510630, P. R. China
10	<sup>3</sup> Precision Medicine Institute, The First Affiliated Hospital, Sun Yat-sen
11	University, Guangzhou, Guangdong, 510030, P. R. China
12	<sup>4</sup> Department of Laboratory Animal Science, College of Basic Medicine Science,
13	Army Medical University, Chongqing, 400037, P. R. China
14	
15	* Corresponding Author:
16	Hong Wei
17	Email:weihong63528@163.com
18	Xiang Fang
19	Email: <u>fxiang@scau.edu.cn</u>
20	
21	





23 Supplementary Fig.1 Chemical structures of major tea polyphenols. A: Catechin;





25 gallate (ECG); E: Epicatechin (EC)







29 Supplementary Fig. 3 Relative abundance of main bacteriophages at the family level



31



33 species level



35 Supplementary Fig. 5 Virome diversity correlates with mirobiome diversity in two

36 groups. (Spearman correlation coefficient = 0.77, p = 0.0053, R<sup>2</sup>= 0.5929)



Fig.6 Relative Supplementary abundance Lactobacillus of and Lactobacillus prophage. The statistical significantly correlations were displayed as \**p*< 0.05, \*\**p*< 0.01. 

Ingredient	kcal/g	g/kg	kcal/g
Com Starch	4	397.5	1590
Casein	4	200	800
Maltodextrin	4	132	528
Sucrose	4	100	400
Soybean Oil	9	70	630
Cellulose	0	50	0
Mineral MixAIN-93G-MX	1.6	35	56
VItamin MixAIN-93-VM	3.9	10	39
L-Cystine	4	3	12
Choline Bitartrate	0	2.5	0
Total		1000	4055

## Supplementary table 1 AIN-93G feed formulation of mice chow food

53

## 54 Supplementary table 2 The result statistics of denovo virus identification

	Sample	Total_base(Mb)	Total_num	Max_len	Min_len	N50	GC
	Novel.viral.contig.final	4.57	2131	112567	300	5666	44.38%
55							
56							
57							
58							
59							
60							
61							
62							
63							
64							
65							

	Sample	Phages	Other_virus	Unclassified
	PFD.1	75.97%	10.80%	13.23%
	PFD.2	47.46%	38.65%	13.89%
	PFD.3	59.52%	1.45%	39.03%
	PFD.4	89.43%	1.60%	8.97%
	PFD.5	41.76%	0.29%	57.96%
	PFD.6	35.81%	2.80%	61.40%
	TPD.1	61.83%	0.52%	37.65%
	TPD.2	86.96%	2.41%	10.63%
	TPD.3	76.62%	0.99%	22.38%
	TPD.4	79.10%	8.96%	11.94%
	TPD.5	61.61%	0.23%	38.16%
	TPD.6	93.15%	0.91%	5.94%
67				
68				
69				
70				
71				
72				
73				
74				

	Sample	DNA	RNA	Unclassified	
	PFD.1	86.75%	0.02%	13.23%	
	PFD.2	86.09%	0.02%	13.89%	
	PFD.3	60.97%	0.00%	39.03%	
	PFD.4	91.03%	0.00%	8.97%	
	PFD.5	42.04%	0.00%	57.96%	
	PFD.6	38.60%	0.00%	61.40%	
	TPD.1	62.35%	0.00%	37.65%	
	TPD.2	89.37%	0.00%	10.63%	
	TPD.3	77.62%	0.00%	22.38%	
	TPD.4	88.06%	0.00%	11.94%	
	TPD.5	61.84%	0.00%	38.16%	
	TPD.6	94.06%	0.00%	5.94%	
76					
17					
78					
79					
30					
31					
32					
33					