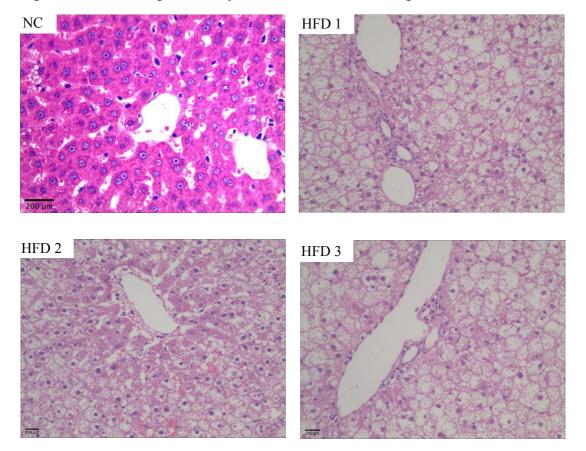
## **Supplementary Materials**



Fgiure S1:Histological analysis of HFD-induced hepatic steatosis.

the HE-stained liver tissue(400X)

Parameters	NC	HFD 1	HFD 2	HFD 3
Steatosis (0-2)	0	2	1	2
Lobular inflammation (0-3)	0	1	2	1
Ballooning degeneration (0-2)	0	2	1	2
Total NAS(0-8)	0	5	4	5

Feeding Method	Time	e Samp le ID	0.97 (The operational taxonomic units(OTUs) that reached a 97% nucleotide similarity level)							
			OTU	ace	chao	coverage	shannon	simpson		
		C011	613	705 (677,747)	708 (672,765)	0.996432	4.57 (4.55,4.59)	0.0337 (0.0327,0.0347)		
		C012	536	647 (614,695)	631 (596,687)	0.995405	4.4 (4.38,4.42)	0.0276 (0.027,0.0282)		
		C014	659	748 (720,787)	776 (733,845)	0.996755	4.56 (4.54,4.58)	0.0302 (0.0295,0.031)		
		C015	666	766 (736,808)	783 (741,848)	0.99577	4.71 (4.69,4.73)	0.0225 (0.0219,0.023)		
		C06	685	798 (766,844)	786 (750,841)	0.996595	4.51 (4.49,4.52)	0.03 (0.0293,0.0306)		
		C07	605	714 (681,760)	736 (688,812)	0.995905	4.33 (4.3,4.35)	0.052 (0.0505,0.0536)		
		H013	800	895 (866,935)	897 (862,951)	0.99832	4.7 (4.69,4.71)	0.0228 (0.0224,0.0231)		
		H016	630	718 (690,757)	711 (680,760)	0.99876	3.91 (3.9,3.93)	0.0507 (0.0501,0.0513)		
LIED	OW	H019	696	823 (788,871)	829 (784,898)	0.997879	3.56 (3.55,3.58)	0.0847 (0.0835,0.0859)		
HFD	OW	H024	738	879 (840,932)	897 (843,979)	0.997984	4.2 (4.19,4.21)	0.0464 (0.0457,0.0471)		
		H08	640	729 (702,769)	713 (686,757)	0.997989	4.04 (4.03,4.06)	0.0545 (0.0536,0.0555)		
		H09	799	870 (847,903)	867 (840,911)	0.999264	4.32 (4.31,4.33)	0.0356 (0.0352,0.036)		
		M017	645	762 (728,809)	784 (736,858)	0.995393	4.33 (4.31,4.35)	0.0439 (0.0427,0.0452)		
		M02	670	811 (772,864)	825 (773,904)	0.994945	4.61 (4.59,4.63)	0.0256 (0.025,0.0263)		
		M020	610	773 (729,833)	751 (705,819)	0.994687	4.01 (3.99,4.04)	0.0529 (0.0517,0.0542)		
		M022	598	723 (687,774)	735 (687,809)	0.994609	4.48 (4.46,4.5)	0.0272 (0.0265,0.0278)		
		M023	564	678 (645,725)	678 (638,741)	0.992933	4.38 (4.35,4.4)	0.0353 (0.034,0.0366)		
		M05	663	753 (726,792)	777 (736,843)	0.996721	4.32 (4.3,4.33)	0.0457 (0.0445,0.0469)		

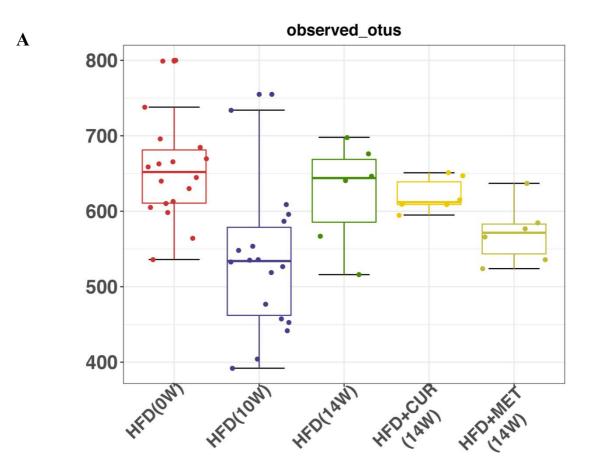
Table S2: The community coverage index of each fece sample.

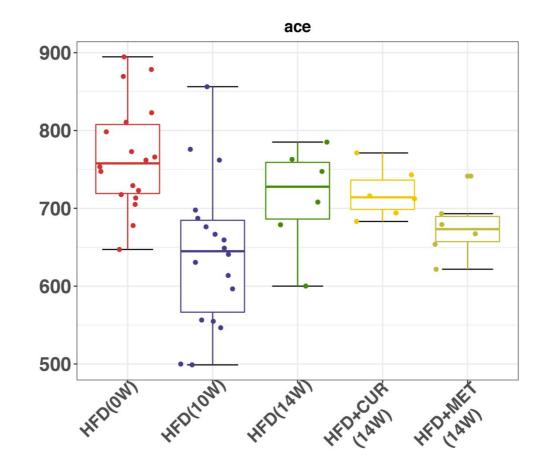
				222			0.07	
		11710	500	698	693	0.998062	3.67	0.0777
		H713	596	(667,742)	(658,748)		(3.65,3.69)	(0.0766,0.0789)
		H716	587	687	682 (646 729)	0.998541	3.96	0.0524
		11/10	007	(656,732)	(646,738)	0.996041	(3.94,3.97)	(0.0517,0.0532)
		H719	519	631 (597,679)	628 (588,691)	0.997873	3.45 (3.44,3.47)	0.0877 (0.0863,0.089)
		11/15	015	. , ,			. , ,	
		H724	554	614 (593,646)	603 (582,639)	0.999112	3.73 (3.72,3.74)	0.0941 (0.0927,0.0956)
			001	776	776	0.999434	4.65	0.0208
		H78	734	(760,801)		0.999434	4.05	(0.0205,0.021)
			101	856	857	0.998728	4.66	0.0232
		H79	755	(826,899)		0.330720	(4.66,4.67)	(0.0229,0.0236)
				649	664		3.98	0.0763
		M717	533	(614,700)		0.996636	(3.95,4)	(0.0742,0.0784)
				547	556		3.33	0.1335
		M72	457	(517,590)		0.997361	(3.31,3.35)	(0.1308,0.1362)
				597	650	0.996316	3.72	0.0921
		M720	477	(559,652)	(583,759)		(3.69,3.74)	(0.09,0.0943)
HFD	10W	M722	536	641	671	0.996929	4.05	0.0601
				(608,688)	(620,753)		(4.03,4.07)	(0.0585,0.0617)
		M723	609	762	775	0.994961	3.95	0.0997
				(719,821)	(719,859)		(3.93,3.98)	(0.0968,0.1026)
		M75	404	500	515	0.997715	3.31	0.1255
				(468,549)	(470,591)	0.997715	(3.29,3.33)	(0.1227,0.1282)
				676	685		3.4	0.1609
		C711	548	(639,730)	(636,762)	0.996546	(3.37,3.42)	(0.1573,0.1644)
				557	556	0.998207	3.91	0.0414
		C712	453	(523,606)	(516,623)	0.330207	(3.9,3.93)	(0.0408,0.0419)
		C714		667	704	0.998352	3.9	0.0615
			535	(627,724)	(641,804)	0.000002	(3.89,3.92)	(0.0604,0.0627)
				499	497	0.996645	3.76	0.053
		C715	392	(463,552)	(455,567)		(3.75,3.78)	(0.0519,0.0541)
				555	571	0.997692	3.55	0.0795
		C76	442	(519,608)	(520,655)		(3.53,3.58)	(0.0781,0.0808)
				659	631	0.996501	3.92	0.0742
		C77	527	(620,715)			(3.9,3.94)	(0.0723,0.0762)
		L101	670	747	744	0.998713	4.39	0.0379
		H121	676	(724,781)	(717,788)	0.000000	(4.38,4.4)	(0.0372,0.0387)
		U101	600	785	802	0.998692	4.81	0.0148
		H121	698	(758,825)	(762,865)	0.007010	(4.8,4.82)	(0.0146,0.0149)
HFD	14W	H121	647	763	741	0.997812	4.14	0.0471
		11121	047	(129,810)	(707,793)		(4.12,4.15)	(0.0463,0.048)

1								
				600	571	0.998396	3.84	0.0727
		H122	516	(573,641)	(549,608)		(3.82,3.85)	(0.0713,0.0742)
				708	704	0.999044	4.57	0.0217
		H128	641	(686,742)	(678,747)		(4.56,4.58)	(0.0214,0.0219)
				679	677	0.994987	4.41	0.0257
		H129	567	(646,726)	(637,739)		(4.39,4.43)	(0.0252,0.0263)
		M121 7	566	667	695	0.996125	4.1	0.066
				(636,712)	(647,772)		(4.08,4.12)	(0.0641,0.068)
		MICO	50.4	622	615	0.996002	3.65	0.1324
		M122	524	(592,665)	(581,670)		(3.62,3.68)	(0.1287,0.136)
		M122		679	685		4.68	0.0297
		0	585	(650,722)		0.995785	(4.66,4.71)	(0.0286,0.0308)
HFD+MET	14W	M122		654	663		4.08	0.0578
		2	536	(619,704)		0.994444	(4.06,4.11)	(0.0561,0.0594)
		M122 3	637	742	751	0.996033	4.48	0.0355
				(710,786)			(4.46,4.5)	(0.0345,0.0365)
			577	693	706	0.995431	4.22	0.0599
		M125		(659,742)			(4.19,4.24)	(0.0579,0.0619)
	14W	C121	595	712	742	0.996522	4.14	0.0693
				(677,762)			(4.12,4.16)	(0.0674,0.0713)
				694 729		4.85	0.0156	
		C121	609	(667,734)	(682,806)	0.997252	(4.84,4.86)	(0.0153,0.0159)
		C121	615	716	730	0.995364	4.91	0.0142
HFD+CUR				(685,760)			(4.89,4.93)	(0.0139,0.0145)
			609	683	687	0.997038	5.06	0.0109
				(659,719)				(0.0107,0.0111)
		C126		743	733	0.996218	4.77	0.0237
				(715,783)			(4.75,4.78)	(0.0229,0.0245)
				771	784		4.52	0.044
		C127	647	(736,820)		0.994986	(4.5,4.54)	(0.0424,0.0455)
The two num	1		dad k			· 1 1	(1.0, 1.0 1)	(0.0121,0.0100)

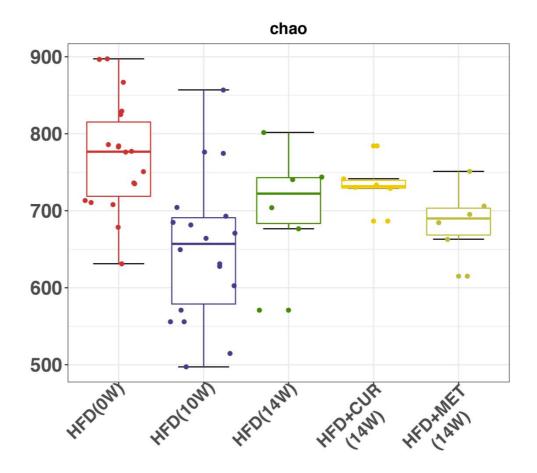
The two numbers surrounded by parentheses are minimal and maximal degrees output by multiple analyses.

Figure S2: The Alpha diversity of each group. (A) The observer OTUs of each group. (B) The ace index of each group. (C) The chao index of each group. (D) The shannon index of each group. (E)The simpson index of each group.

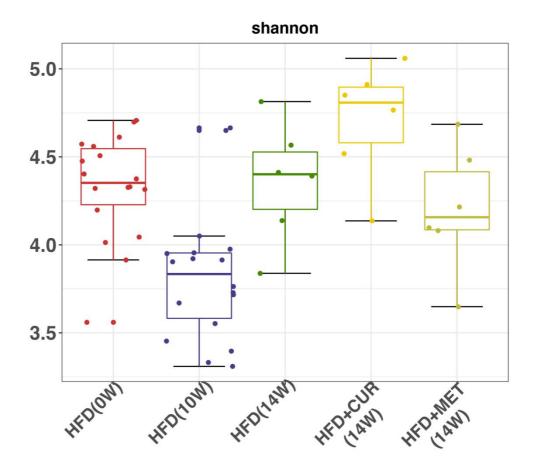




С



B



E

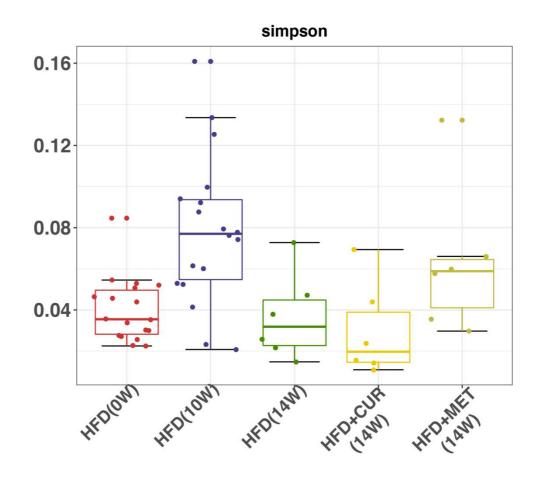
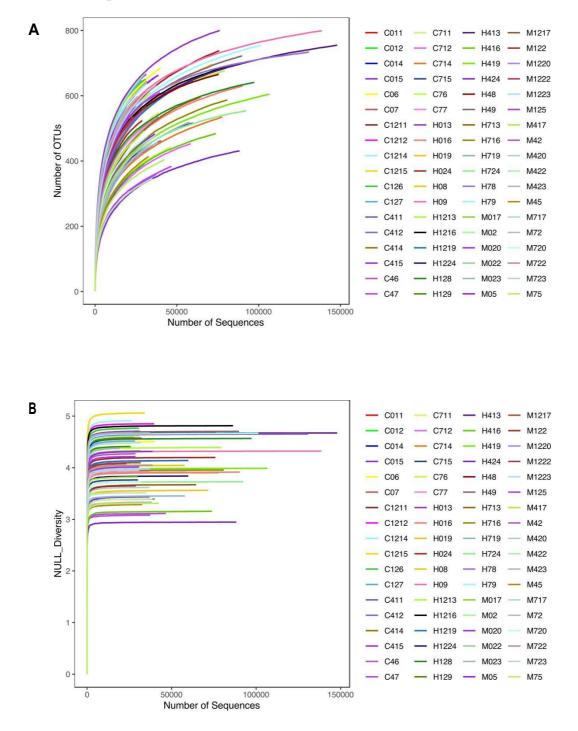


Figure S3: The rarefaction curves and Shannon curves of each fece sample. (A) The rarefaction curves of each fece sample. (B) The Shannon curves of each fece sample.



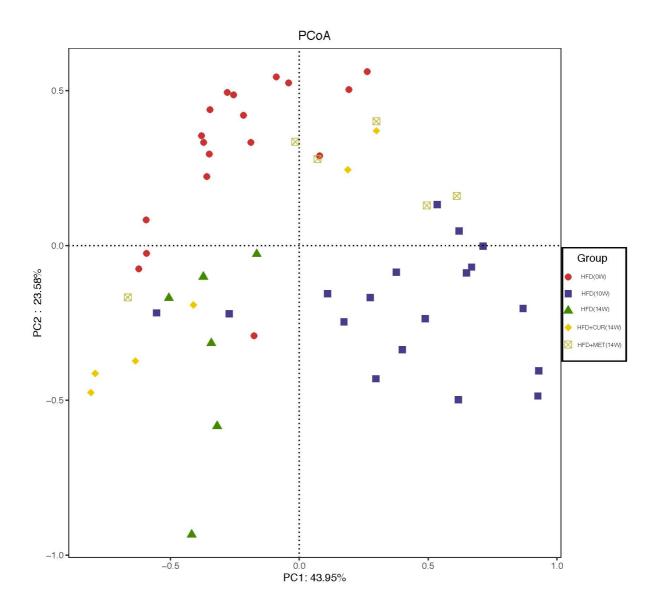
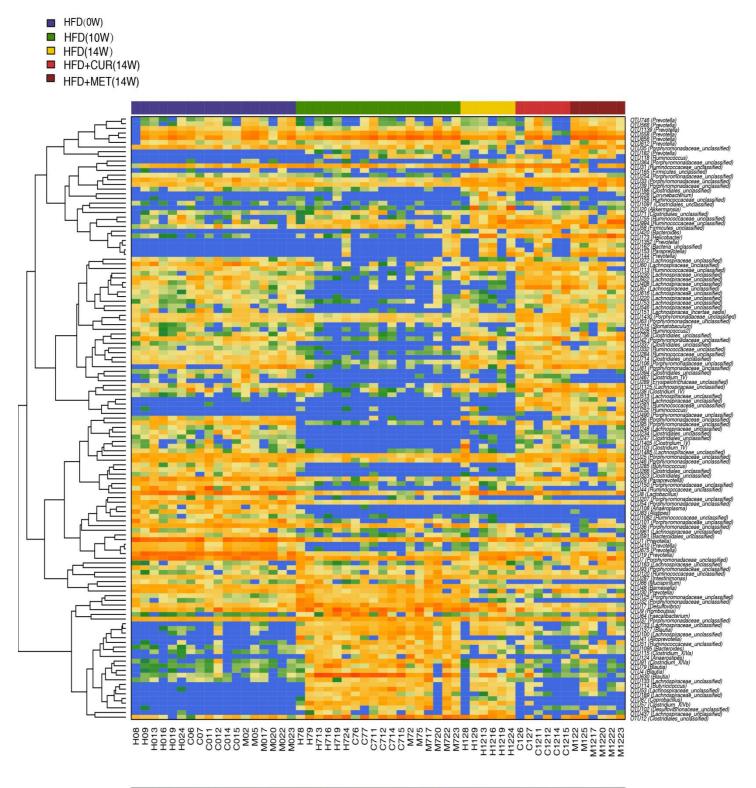


Figure S4: PCoA score plot based on weighted Unifrac metrics.

## Figure S5: Heatmap of the abundance the OTUs in the HFD(0W), HFD (10W), HFD (14W), HFD+CUR(14W) and HFD+MET(14W) groups.





Relative abundance of community (%)