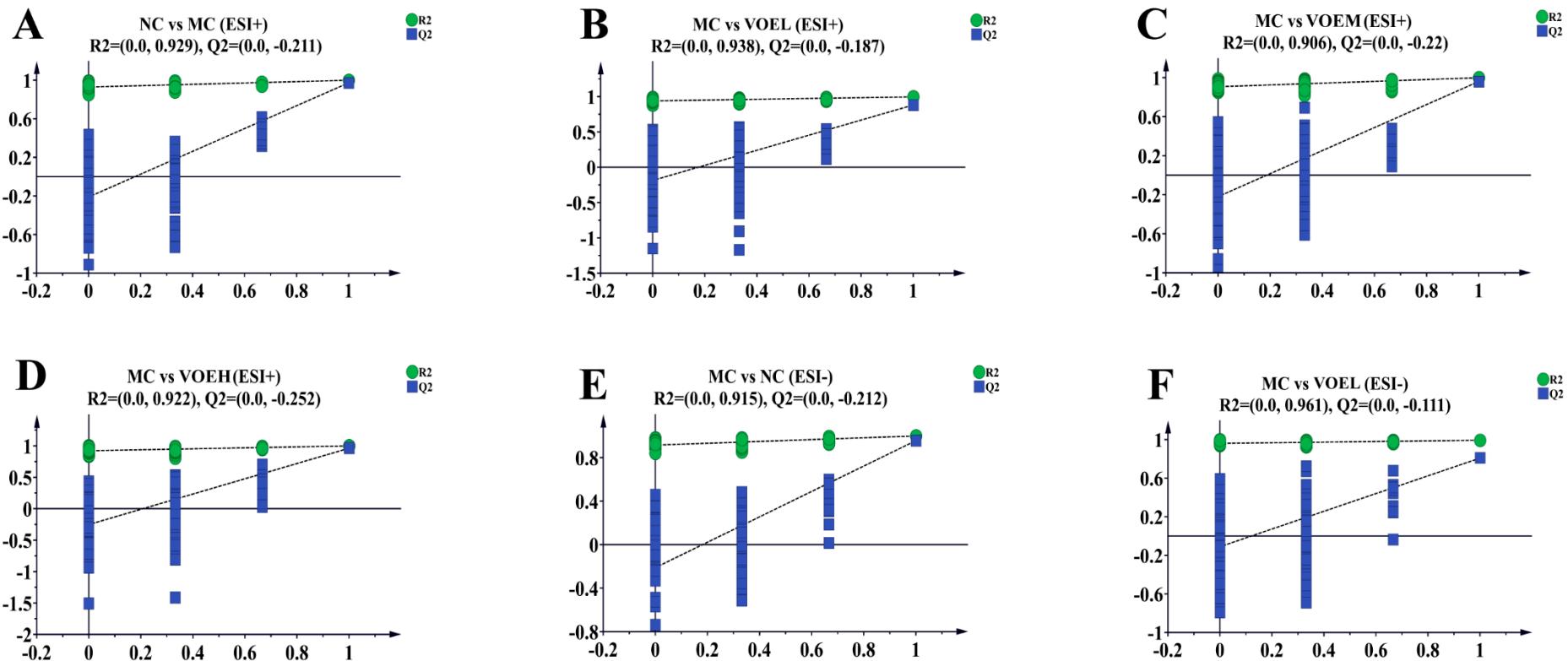
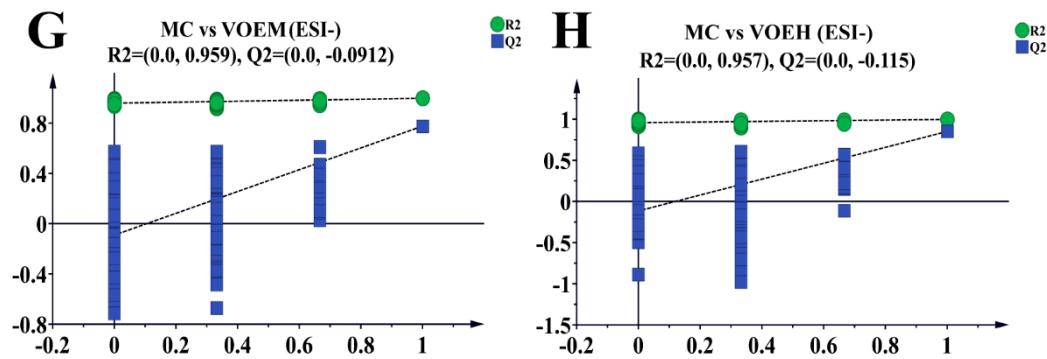


**Figure S1.** Effects of VOE on the  $\alpha$  diversity of gut microbiota in immunosuppressed mice. (A) Chao 1, (B) ACE, (C) Shannon and (D) Simpson indexes. Values with different letters of the same indicator are significantly different ( $p < 0.05$ ).





**Figure S2.** The permutation of OPLS-DA (A), (C), (E), (G) in positive ion mode and (B), (D), (F), (H) in negative ion mode in feces samples of mice

**Table S1 Fecal differential metabolites in NC, MC, and VOEH groups**

No.	Compound	MC vs NC		MC vs VOEH		HMDB ID/ KEGG ID	RT	Mass
		log <sub>2</sub> FC	Change	log <sub>2</sub> FC	Change			
1	7-Sulfocholic acid	2.27	up	2.27	up	HMDB0002421	9.70	488.24
2	3-Pyridylacetic acid	1.92	up	1.92	up	HMDB0001538	1.63	137.05
3	Cyclopassifloic acid B	2.31	up	1.49	up	HMDB0038388	21.12	580.40
4	Urobilinogen	-9.23	down	-8.85	down	C05791	10.74	590.31
5	Formamidopyrimidine nucleoside triphosphate	1.62	up	1.62	up	C05922	10.39	562.98
6	Ganglioside GD1a (d18:1/22:0)	1.94	up	1.94	up	HMDB0011799	10.59	1892.05
7	Cer (d18:1/18:0)	2.25	up	1.48	up	HMDB0000829	21.89	1092.73
8	L-655,240	2.85	up	3.52	up	HMDB0341170	25.10	373.13
9	N-arachidonoyl tyrosine	2.40	up	1.64	up	HMDB0062331	17.30	513.31
10	N-docosahexaenoyl phenylalanine	2.17	up	1.36	up	HMDB0062333	24.85	475.31
11	N-oleoyl tyrosine	2.46	up	1.07	up	HMDB0062337	24.27	445.32
12	PA (22:6/20:5)	1.93	up	1.15	up	HMDB0115418	9.38	766.46
13	PA (20:0/22:4)	1.95	up	1.22	up	HMDB0115085	22.32	824.59
14	Pantetheine	2.22	up	1.57	up	C00831	6.67	300.11
15	PC (15:0/0:0)	1.63	up	1.19	up	HMDB0010381	18.39	481.32
16	Rhizoctinic A	1.96	up	1.16	up	C17944	4.89	397.13
17	Siroheme	1.96	up	1.96	up	C00748	10.82	914.22
18	Trichostatin A	2.22	up	1.44	up	HMDB0259177	9.41	302.16