

Gut Dysbiosis and Increased Intestinal Permeability Drive microRNAs, NLRP-3 Inflammasome and Liver Fibrosis in a Nutritional Model of Non-Alcoholic Steatohepatitis in Adult Male Sprague Dawley Rats [Erratum]

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This error was introduced by the Editorial staff during the publication process. The correct Table 5 is shown below.

The authors have advised that the colour coding from Table 5 was missed on page 363 of the published paper.

Table 5 Correlations between NASH markers and microbiota composition

Variables [#]	Inflammatory Response		Intestinal Permeability		Microbiota composition
	Index TNF- α /IL-10 (pg/mg)	Index IL-1 β /IL-10 (pg/mg)	Ocln	Fllr (JAM-1)	
NAFLD scoring	0.751**	0.832**	-0.719**	-0.525*	0.694**
Index TNF- α /IL-10 (pg/mg)		0.729**	-0.463*	-0.208	0.234*
Index IL-1 β /IL-10 (pg/mg)			-0.457*	-0.361	0.522*
Ocln				0.722**	0.090
Fllr (JAM-1)					0.146*

Notes: [#]Variables was represented for Spearman's correlation coefficient. Moderate ($0.3 < r < 0.6$ - blue) and strong ($0.6 < r < 0.9$ - green) correlation of Spearman's. **Correlation is significant at the 0.01 level. *Correlation is significant at the 0.05 level.

Abbreviations: IL, interleukin; JAM, junctional adhesion molecule; NAFLD, non-alcoholic fatty liver disease; NASH, non-alcoholic steatohepatitis; Ocln, occluding; TNF, tumor necrosis factor.

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