Supplementary material:

Table 1: Tissue distribution and sites of origin for the pattern recognition molecules of the lectin pathway.

Protein	Tissue distribution	References
MBL	Liver	[1-3]
M-ficolin	Peripheral blood leukocytes, lung	[4-7]
L-ficolin	Liver	[6, 8]
H-ficolin	Liver (hepatocytes, bile duct epithelial cells), lung	[9, 10]
CL-L1	Liver	[11]

In addition to the references above describing tissue distribution and sites of origin for the lectin pathway proteins, a Swedish company has specialized in RNA and immunohistochemical analyses of all human proteins. These are displayed online at www.proteinatlas.org [12]. Below is a summary of the pattern recognition molecules of the lectin pathway proteins based on their distribution according to www.proteinatlas.org. The tissues are listed according to expression from highest to lowest.

Table 2: Tissue distribution of the pattern recognition molecules according to the Proteinatlas.org.

Protein (gene)	RNA	IHC
MBL (mbl2)	Liver	Highly specific cytoplasmic expression in liver.
M-ficolin (fcn1)	Bone marrow, appendix, spleen, urinary bladder, lung	A subset of leukocytes, macrophages in lung, cells in spleen and bone marrow.
L-ficolin (fcn2)	Liver	Pending analyses.
H-ficolin (fcn3)	Lung, liver	Fractions of cells in the gastrointestinal tract, thyroid gland and hepatocytes showed distinct granular cytoplasmic staining. The adrenal gland was strongly stained.
CL-L1 (colec10)	Liver	Normal liver tissue showed moderate to strong, often granular, cytoplasmic staining.

IHC (Immunohistochemical staining)

Figure 1: MBL levels in patients with acute liver failure at day 1 and day 3 in relation to infection status. The box plots represent median and 25–75 percentiles. Upper and lower lines are the upper and lower adjacent values.

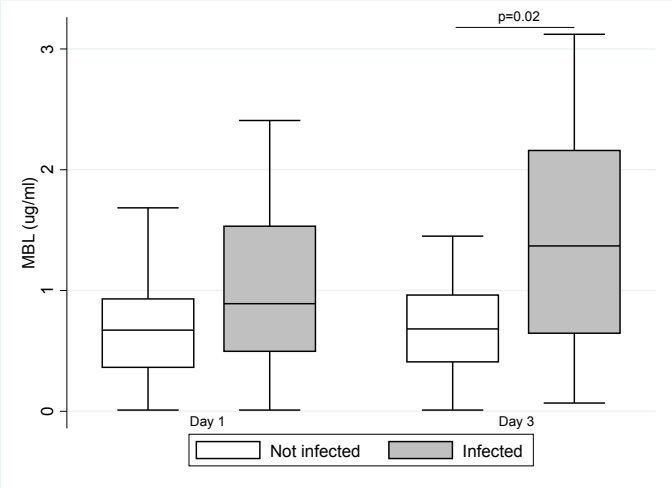
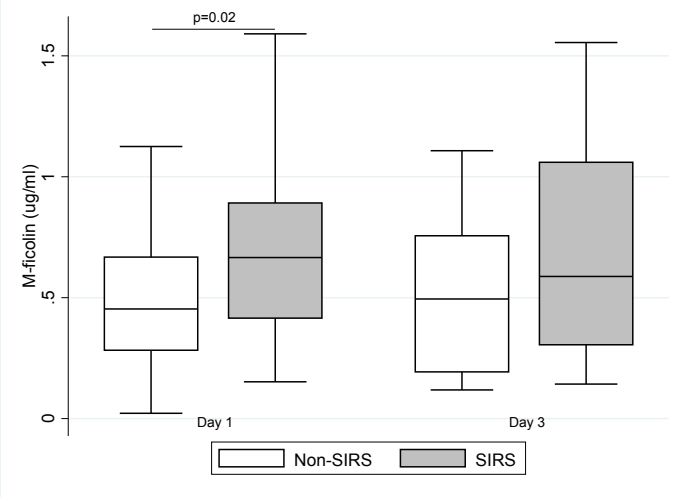


Figure 2: M-ficolin levels in patients with acute liver failure at day 1 and day 3 in relation to the SIRS criteria.



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