

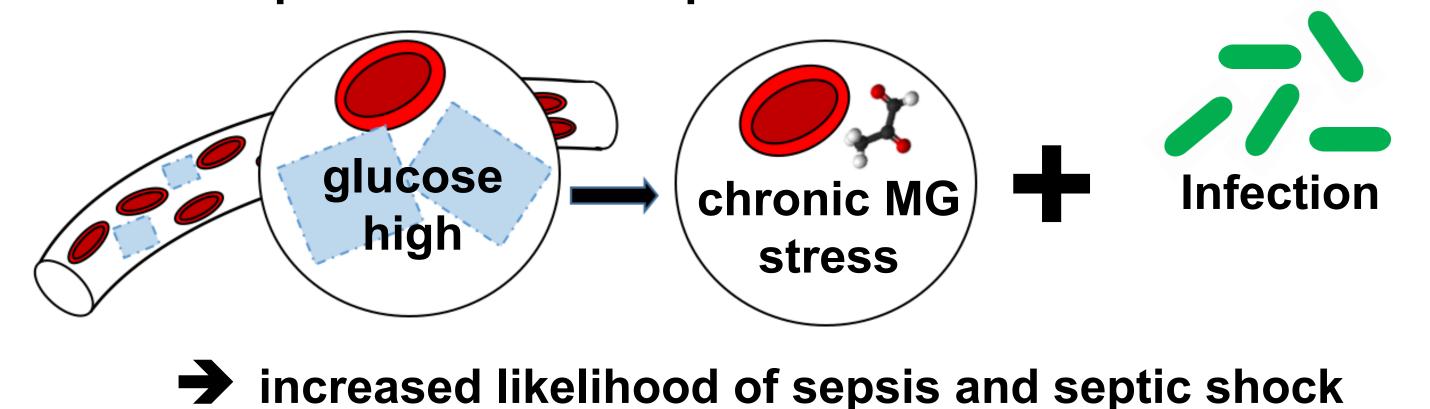
1. Which organs and which cell entities are the main sources of

Outlook

Sepsis as a late complication of Diabetes mellitus

MG-derived carbonyl stress in sepsis?

- Endogenic vs. exogenic (e.g. bacterial synthesis of MG) sources?
- Which mechanisms are responsible for the fast increase of MG plasma levels in sepsis?
- 2. Does MG-derived carbonyl stress influence the course of the disease in sepsis?
 - Which mechanisms are responsible for MG-caused damages?
 - Is it possible to modify MG-derived carbonyl stress therapeutically?



Hypothesis: Sepsis \rightarrow endogenic MG-formation $\hat{T} + MG$ -detoxification $\sqrt[]{} \rightarrow$ mortality \hat{T} (causal)

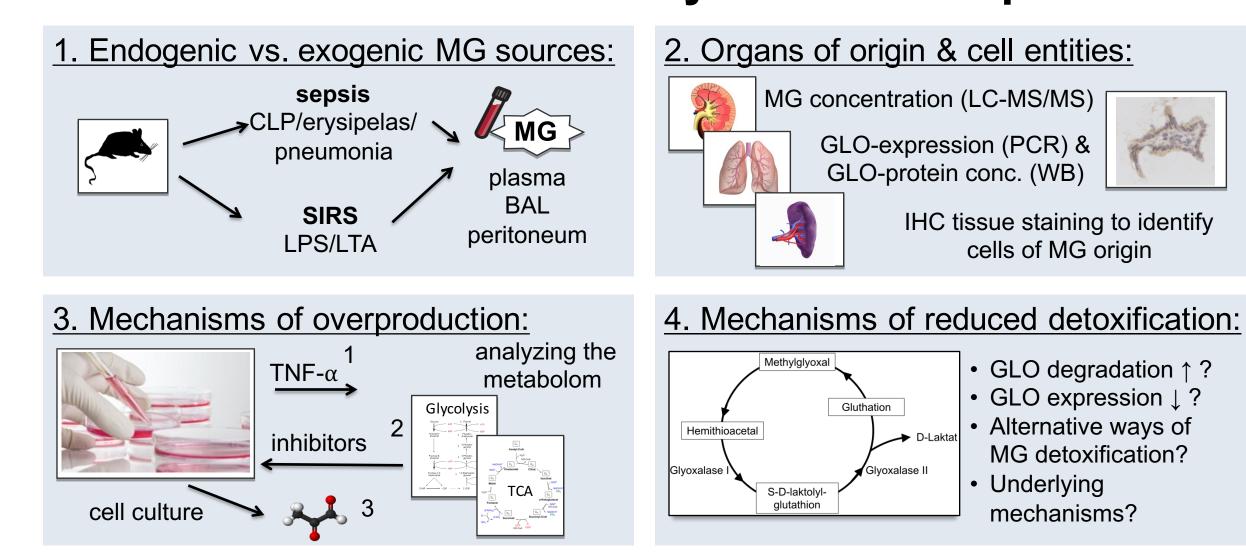
Background

- 1. The plasmatic load of MG and MG-AGE is associated with severe diabetic complications such as nephro-, retino- and neuropathy.
- 2. However, these late sequelae of DM develop within years.

Remaining question: Is MG able to evoke a pivotal influence on the course of the disease in sepsis within hours?

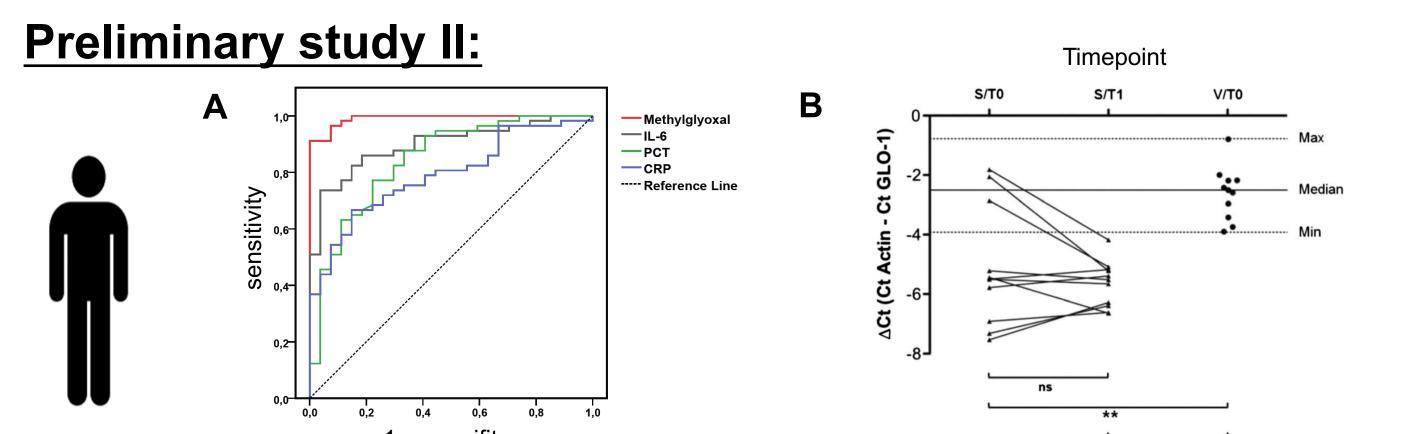
Work Program

Aim A: Main source of MG-derived carbonyl stress in sepsis



Background Data Preliminary study I: B 1000-800-*** onset *** [MM] [Mn] [Vu] 800 600-1000a <u>क</u> 600-400ethylglyo **Methylgly**(400-200-200-healthy sepsis day 1 day 4 day 7 day 14 day 28 post-OP post-OP sepsis

MG plasma levels are significantly increased in patients suffering from sepsis in comparison to healthy volunteers (A) and postoperative controls (B). At the time of sepsis diagnosis, nonsurvivors revealed significantly higher MG plasma levels in comparison to survivors (C).



Question: Which cell entities show such an MG overload, which might serve as an explanation for the MG plasma kinetics observed in septic patients?

Within the 1st step we seek to investigate the relevance of exogenic (=bacterial) MG production in sepsis. Furthermore we will try to answer the question, whether a treatment with antibiotics is able to influence MG-derived carbonyl stress in sepsis. Afterwards we seek to identify the organs as well as the ensuing cell entities, which are responsible for MG-derived carbonyl stress in early sepsis (2nd step). Moreover, we will try to illuminate the underlying mechanisms for MG-accumulation in sepsis (3^{rd} step: production \uparrow vs. 4^{th} step: detoxification \downarrow).

Aim B:

Causal impact of MG-derived carbonyl stress in sepsis

Xcelligence

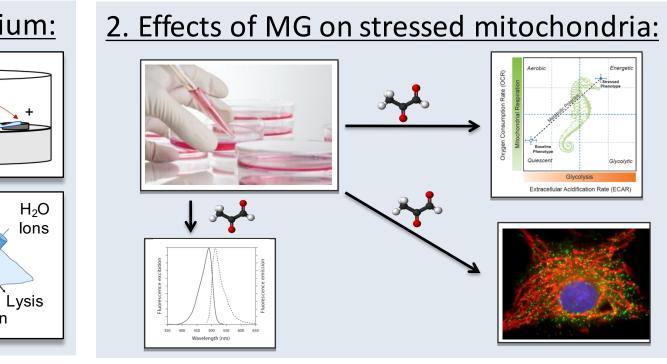
CENTRE A

ιL1β/IL18

<u>1. Effects of MG on activated endothelium:</u> Effects on cell-cell-

- contacts?
- Induction of pyroptosis?
- Changes in morphology?
- Modulation capability via
- scavenger administration? • Are the effects visualisable

via intravital microscopy? **Discharge of Interleukin**

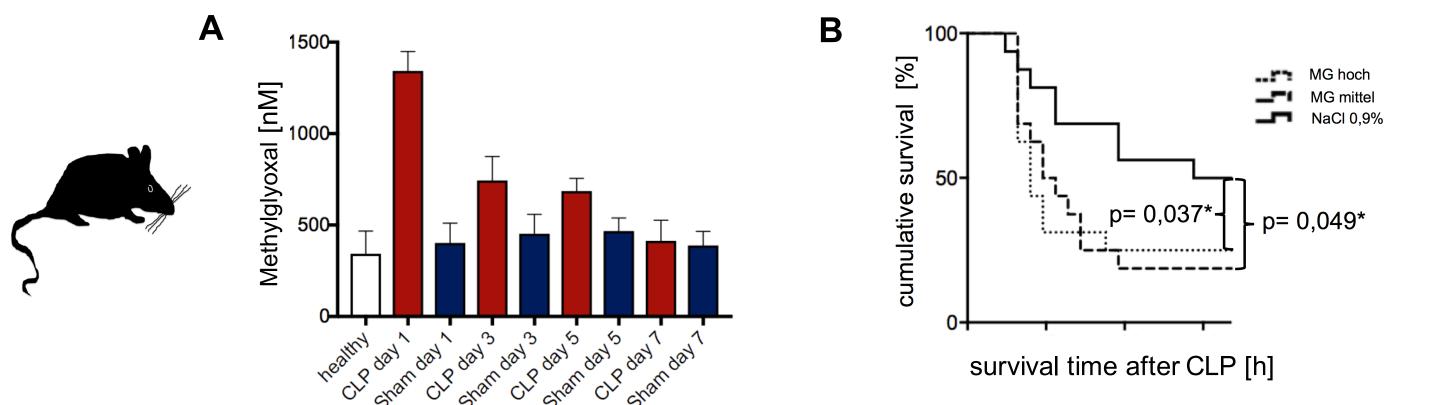


Question: Via which mechanisms is MG able to

1 – specifity

MG plasma levels proved to be superior for the identification of patients with sepsis as compared to the standard parameters PCT, IL-6 und CRP (A). Moreover, GLO-I expression in pBMCs of septic patients was shown to be significantly reduced in comparison to healthy controls (B).

Preliminary study III:



Plasma kinetics of MG in Cecal Ligation and Pucture (CLP)-induced murine sepsis were shown to be comparable to those of septic patients (A). Moreover, an additional MG-administration was associated with a dose-dependent decrease of survival in CLP-mice (B). Accordingly, MGderived carbonyl stress might have a causal impact on the course of the disease in sepsis.

influence the course of the disease in sepsis?

A loss of the endothelial barrier is a decisive factor contributing to serious disturbances of microcirculation, ultimately leading to septic shock. Therefore, we first seek to investigate possible effects of MG-derived carbonyl stress on activated endothelium (1st step). Afterwards, we will try to evaluate the effects of MG on mitochondrial function as well as the global capacity of energy supply (**2**nd step).

Publications

- **Brenner, T.**, Fleming, T., Uhle, F., Silaff, S., Schmitt, F., Salgado, E., Ulrich, A., Zimmermann, S., Bruckner, T., Martin, E., et al. (2014). Methylglyoxal as a new biomarker in patients with septic shock: an observational clinical study. Crit. Care Lond. Engl. 18, 683.
- Hofer, S., Uhle, F., Fleming, T., Hell, C., Schmoch, T., Bruckner, T., Weigand, M.A., and **Brenner, T.** (2016). RAGE-mediated inflammation in patients with septic shock. J. Surg. Res. 202, 315–327.
- Schmoch, T., Uhle, F., Siegler, B.H., Fleming, T., Morgenstern, J., Nawroth, P.P., Weigand, M.A., and **Brenner, T.** (2017). The Glyoxalase System and Methylglyoxal-Derived Carbonyl Stress in Sepsis: Glycotoxic Aspects of Sepsis Pathophysiology. Int. J. Mol. Sci. 18.