Markers of systemic inflammation in response to osmotic stimulus in healthy volunteers

Clara Odilia Sailer¹², Sophia Julia Wiedemann¹³, Konrad Strauss⁴, Ingeborg Schnyder¹, Wiebke Kristin Fenske⁵, Mirjam Christ-Crain¹²

¹Department of Endocrinology, Diabetology and Metabolism, University Hospital Basel, Basel, Switzerland
²University of Basel, Department of Clinical Research, Basel, Switzerland
³Department of Biomedicine, University of Basel, Basel, Switzerland
⁴Division of Endocrinology and Diabetes, Department of Internal Medicine I, University Hospital, University of Würzburg, Germany
⁵Leipzig University Medical Center, Integrated Center for Research and Treatment Adiposity Diseases, Leipzig, Germany;

Short title: Systemic inflammation and osmotic stimulus

Key Terms: TNF-α, interleukin-8, interleukin-6, copeptin, hyperosmolality

Clinical Trial Information: ClinicalTrials.gov no. NCT02647736 (registered 6 January 2016).

Word count (Introduction through Acknowledgments): 2001

Correspondence: Dr. med Clara Sailer
Department of Endocrinology, Diabetology and Metabolism
University Hospital Basel
Petersgraben 4, 4031 Basel, Switzerland
Phone: +41 61 265 50 78 / Fax: +41 61 265 51 00
E-mail: clara.sailer@usb.ch
Abstract

Hyperosmolality, osmotic stimulus or stress results in vasopressin release. Animal and human in vitro studies have shown that inflammatory parameters such as interleukin-8 (IL-8) and tumour necrosis factor-α (TNF-α) increase in parallel in the central nervous system and bronchial, corneal or intestinal epithelial cell lines in response to osmotic stimulus. Whether osmotic stimulus directly causes a systemic inflammatory response in humans is unknown. We therefore investigated the influence of osmotic stimulus on circulatory markers of systemic inflammation in healthy volunteers. In this prospective cohort study, 44 healthy volunteers underwent a standardized test protocol with an osmotic stimulus leading into the hyperosmotic/hypernatremic range (serum sodium ≥150mmol/l) by hypertonic-saline infusion. Copeptin – a marker indicating vasopressin activity –, serum sodium and osmolality, plasma IL-8 and TNF-α were measured at baseline and directly after osmotic stimulus. Median (range) serum sodium increased from 141 mmol/l (136, 147) to 151 mmol/l (145, 154) (p-value <0.01), serum osmolality increased from 295 mmol/l (281, 306) to 315 mmol/l (304, 325) (p-value <0.01). Median (range) copeptin increased from 4.3 pg/l (1.1, 21.4) to 28.8 pg/l (19.9, 43.4) (p-value <0.01). Median (range) IL-8 levels showed a trend to decrease from 0.79 pg/ml (0.37, 1.6) to 0.7 pg/ml (0.4, 1.9) (p-value 0.09) and TNF-α levels decreased from 0.53 pg/ml (0.11, 1.1) to 0.45 pg/ml (0.12, 0.97) (p-value 0.036). Contrary to data obtained in vitro, circulating proinflammatory cytokines tend to or decrease in human plasma after osmotic stimulus. Osmotic stress does not to stimulate circulating markers of systemic inflammation.
Introduction

Hypernatremia and hyperosmolality are common among hospitalized patients and especially among those in the intensive care unit (up to 9%) and are associated with substantial morbidity and mortality. Causes of hypernatremia and hyperosmolality range from dehydration due to volume loss (e.g., diarrhoea, sweating) or diabetes insipidus, to salt overload. Hypernatremia and hyperosmolality induce shrinking of cells, mainly in the central nervous system, leading to neurological dysfunctions and symptoms such as thirst, disorientation, lethargy, convulsion and ultimately death.

Hypernatremia and a consequent rise in serum osmolality increase the perception of thirst and induce the secretion of the hormone arginine vasopressin (AVP) leading to a reduced renal water excretion. Animal data have shown that upon hypernatremia and hyperosmolality, together with AVP secretion, there is an upregulation of inflammatory markers in the supraoptic and paraventricular nuclei of the hypothalamus. Furthermore, it was shown that hyperosmolality increases the secretion of proinflammatory cytokines from macrophages. Human in vitro studies in bronchial, corneal and intestinal cell lines consistently showed an increased inflammatory response with an increase in interleukin-6 (IL-6), interleukin-8 (IL-8) and tumour necrosis factor alpha (TNF-α) upon hypernatremia and hyperosmolality. Human in vivo studies investigating the effect of hypernatremia on the inflammatory system are scarce. In patients with systemic inflammatory response syndrome (SIRS) due to ketoacidosis and hyperosmolality an increase in inflammatory parameters was observed, but no other data are available.

No study looked at the isolated effect of hypernatremia and hyperosmolality on systemic inflammatory markers in healthy volunteers. The aim of this study was therefore to prospectively investigate circulating markers of systemic inflammation in healthy volunteers undergoing a hypertonic saline infusion in a prospective study.
Materials and Methods

Study Design

This is a secondary analysis of a prospective study including 44 healthy volunteers with available plasma samples\textsuperscript{15}. Inclusion criteria were normonatremia (serum sodium levels of 135-145 mmol/l) and age >18 years. Exclusion criteria involved a history or presence of polyuria/polydipsia syndrome, hyponatremia/hypernatremia, heart failure, liver cirrhosis, nephrotic syndrome or renal failure, anaemia, uncontrolled hypertension, chronic alcohol consumption, body mass index (BMI) <18 kg/m\textsuperscript{2} or >28 kg/m\textsuperscript{2}, drug intake (except oral contraception), and other therapy-requiring diseases. The study was approved by the Ethical Committee Northwest and Central Switzerland, University of Basel, Basel, Switzerland and the Ethical Committee of the University of Würzburg, Würzburg, Germany. Written informed consent has been obtained from each participant after full explanation of the purpose and nature of all procedures used. The study was registered on ClinicalTrials.gov (NCT02647736).

Study procedure

The study procedure has been described in detail elsewhere\textsuperscript{15}. In Brief, smoking and alcoholic beverages were allowed until 24h before, eating until midnight before and drinking until 6 am of the test day. On the test day, participants underwent a standardized medical examination and a baseline blood sample for laboratory evaluation of sodium, osmolality, copeptin, IL-6, IL-8 and TNF-\textgreek{a} was taken following a 30-minute rest in a supine position.

Participants received a 250-ml intravenous bolus of hypertonic saline (3% saline, 513 mOsm/l) within 15 minutes, which was continued thereafter at a rate of 0.15 ml/kg body weight/min until serum sodium reached 150 mmol/l or total infusion time exceeded 180 minutes. At this point, a stimulated blood sample was taken for laboratory evaluation of sodium, osmolality, copeptin, IL-6, IL-8 and TNF-\textgreek{a}.

Laboratory measurements

Laboratory measurements for sodium and osmolality were performed immediately in non-frozen serum samples by automated biochemical analyses. Blood for copeptin, IL-6, IL-8 and TNF-\textgreek{a} analysis...
was sampled in EDTA tubes, immediately centrifuged at 4°C, and plasma was stored at –80°C until analysis. Copeptin was measured in a single blinded batch by a chemiluminescence sandwich immunoassay (B.R.A.H.M.S CT-proAVP LIA; Thermo Scientific Biomarkers, Hennigsdorf, Germany) as previously described\textsuperscript{15}. IL-8 and TNF-\(\alpha\) were measured by MSD Mesoscale (V-PLEX Human Proinflammatory Panel II (4-Plex)) according to the manufacturer’s specifications. IL-6 was measured in a subset of samples by MSD Mesoscale (V-PLEX Human IL-6 Kit) according to the manufacturer’s specifications.

**Statistical analysis**

Discrete variables are expressed as absolute numbers, continuous variables as median and interquartile range (IQR 25\textsuperscript{th}, 75\textsuperscript{th} percentiles) and range (min, max). Comparison between baseline and stimulated laboratory values was computed using a Wilcoxon rank sum test. To adjust for age and sex, a multiple linear regression model and a linear mixed effect model was performed. Graphical analysis was done using boxplots of baseline and stimulated laboratory values, points representing each individual patient.

A two-sided p-value of <0.05 was considered to be statistically significant. Analyses were performed using R statistical software\textsuperscript{16}. 
Results

Baseline characteristics

Forty-four healthy volunteers (23 females, 21 males, aged 30 years (IQR 26, 42)) were included in this analysis. Baseline characteristics are shown in Table 1. At the start of the test, serum sodium and osmolality were in the normal range in all participants (Table 2). In a subset of 13 healthy volunteers, we also measured IL-6 (6 females, 7 males, aged 30 years (IQR 27, 44)) (data not shown).

Sodium, osmolality and copeptin levels upon osmotic stimulus

After osmotic stimulus, serum sodium increased significantly from 141 mmol/l (IQR 140, 142; range 136, 147) to 151 mmol/l (IQR 150, 153; range 145, 154) (p < 0.01) and serum osmolality increased significantly from 295 mmol/kg (IQR 291, 298; range 281, 306) to 315 mmol/l (IQR 312, 317; range 304, 325) (p-value < 0.001). Copeptin increased significantly from 4.3 pg/l (IQR 3.3, 6.7; range 1.1, 21.4) to 28.8 pg/l (IQR 19.9, 43.4; range 3, 125.2) (p-value < 0.001) (Table 2, Figure 1).

Circulatory markers of systemic inflammation upon osmotic stimulus

After osmotic stimulus, IL-8 tended to decrease from 0.79 pg/l (IQR 0.65, 1.11; range 0.37, 1.6) to 0.7 pg/l (IQR 0.57, 0.9; range 0.4, 1.9) (p-value 0.09) and TNF-a decreased significantly from 0.53 pg/l (IQR 0.39, 0.69; range 0.11, 1.1) to 0.45 pg/l (IQR 0.3, 0.53; range 0.12, 0.97) (p-value 0.036) (Table 2, Figure 2, Figure 3). In a subset of patients, IL-6 remained unchanged (0.59 pg/l before osmotic stimulus, 0.67 pg/l after osmotic stimulus, p-value = 0.6) (data not shown).

In a multiple linear regression model adjusting for age and sex, TNF-α remained statistically significant whereas IL-8 and IL-6 remained statistically non-significant. In a mixed effect model with timepoint (before and after stimulus), age and sex as fixed effects and patients as random effect, TNF-α remained statistically significant whereas IL-8 and IL-6 remained statistically non-significant (Table 3).

Correlation of IL-8 and serum sodium was R = -0.04, of IL-8 and serum osmolality was R = -0.12 and of IL-8 and copeptin was R = -0.15. Correlation of TNF-α and serum sodium was R = -0.14, of TNF-α and
serum osmolality was $R = -0.15$ and of TNF-$\alpha$ and copeptin was $R = -0.1$. Correlation of IL-6 and serum sodium was $R = -0.13$, of IL-6 and serum osmolality was $R = 0.1$ and of IL-6 and copeptin was $R = 0.1$. 
Discussion

The main finding of this study is that hypertonic saline infusion leading to hypernatremia and hyperosmolality does not lead to an increase of circulating markers of systemic inflammation in healthy volunteers, but instead leads to a decrease in TNF-α, a possible decrease in IL-8 while IL-6 remains unchanged.

This is in contrast to the literature, where in vitro data suggested an increase in inflammatory parameters upon hyperosmolality\(^5\)\(^–\)\(^7\). Different explanations for these unexpected findings are possible: first, the hyperosmolar range that is feasible to be induced in animals and in human in vitro cell lines is much higher than the one in healthy humans, reaching up to 1200 mmol/l\(^8\),\(^11\),\(^12\). An increase in inflammatory markers was evident at a level of around 400-600 mOsm/kg whereas we increased serum osmolality only up to 315 mmol/l. This higher osmolality in vitro is likely to be more potent in changing gene expression with subsequent production of inflammatory markers. An increase of serum osmolality to around 315 mOsm/kg already induced symptoms such as nausea, vomitus and concentration difficulties\(^15\). A higher serum osmolality could induce serious neurological complications and ultimately death\(^3\) which evidently is not justifiable for healthy humans.

Second, the duration of cell lines and tissue exposed to hyperosmolality was longer in vitro studies than in our population. While the duration of our hyperosmotic stimulus was only for several minutes (as we stopped saline infusion once a sodium \(\geq 150\) mmol/l was reached), cell lines were exposed for several hours to days\(^11\),\(^13\). It can therefore be speculated that chronic hyperosmolality is necessary to induce inflammation\(^17\). It is thus possible that a short stimulus into the hypernatremic range might not be sufficient to observe an increase in circulating markers of inflammation.

Third, animal data showed a specific increase in mRNA expression only within the rat central nervous system (supraoptic and paraventricular nuclei of the hypothalamus)\(^6\). However, samples taken from the rat peripheral circulation after hypertonic saline exposure did not indicate an increase in either IL-6, nor IL-1β\(^18\). We cannot conclude from our experiment, whether a central increase in inflammatory markers was present in our population.
Fourth, the medium to induce a hyperosmotic state was either increasing glucose, glucose polyols (D-Mannitol, Propylene Glycol and Glycerol), or serum sodium (NaCl, sodium chloride)\textsuperscript{11,13}. In our study, we induced hyperosmolality by increasing serum sodium only. In regard of recent advances made regarding the role of glucose induced inflammation\textsuperscript{19}, it could be speculated that the rise in inflammatory parameters observed in these studies might be caused by the increase in serum glucose itself rather than the hyperosmolality.

Some of the measured inflammatory markers, i.e. TNF-\(\alpha\), even decreased upon osmotic stimulus. We can only speculate about possible reasons. It has been shown that inflammatory markers are able to trigger AVP release independent of plasma osmolality\textsuperscript{20,21}. Possibly, the observed decrease in inflammatory parameters in our study may be because of a negative feedback mechanism of AVP on inflammatory markers.

One study in severely ill patients with SIRS due to ketoacidosis and hyperosmolality showed increased inflammatory markers\textsuperscript{14}. In these patients, it is however likely that the underlying disease, such as SIRS, triggered the inflammatory response. Furthermore, as mentioned above, the increase in glucose levels which commonly occurs in critically ill patients is known to induce inflammation\textsuperscript{19}.

According to our results, a short increase into the hyperosmotic range, such as in the saline infusion test to differentiate primary polydipsia from diabetes insipidus\textsuperscript{15}, does not induce an increase in circulating markers of systemic inflammation.

Some limitations of our study need to be mentioned. First, this is a secondary analysis and the results are hypothesis generating and should thus be interpreted with caution. Second, we only increased serum sodium to 150 mmol/l and serum osmolality to 315 mmol/l for a short time period, whereas in vitro studies investigated the inflammatory response with higher sodium and osmolality levels and over a longer period of time. However, this is arguably not possible in humans. We can therefore not exclude a rise in inflammatory parameters at a higher osmotic threshold and/or a later rise in inflammatory response after a longer stimulus.
This is the first study investigating circulating markers of systemic inflammation in humans in response to a strong osmotic stimulus into the hypertonic range. It adds valuable inputs on the understanding of hypernatremia and hyperosmolality and inflammation.

In summary, our data show that in healthy volunteers, hyperosmolality by intravenous saline infusion does not increase circulating markers of systemic inflammation which has been shown in animals and in human bronchial, corneal and intestinal cell lines. Short term exposure into a hyperosmotic state such as practiced in a new approach for diabetes insipidus seems not to induce an inflammatory response. Whether this differs for long term exposure or increased hyperosmolality remains unclear.
Acknowledgments

We thank Cemile Bathelt and Nina Hutter (Clinical Trial Unit, University Hospital Basel) and Mary Beth Gaskill (Clinical Research Unit Laboratory, Northwestern University) for excellent technical assistance.

Financial Support: This work was supported by Thermo Fisher Scientific Biomarkers (Hennigsdorf, Germany) and by the Margot und Erich Goldschmidt & Peter René Jacobson-Stiftung (to COS and to SJW).

Disclosure Summary: W.K.F. and M.C.-C. received speaking honoraria by Thermo Fisher Scientific. The remaining authors have nothing to disclose.
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Figures

Figure 1: Copeptin before and after osmotic stimulus
Figure 2: IL-8 before and after osmotic stimulus
Figure 3: TNF-α before and after osmotic stimulus
### Table 1: Baseline Characteristics

<table>
<thead>
<tr>
<th></th>
<th>Healthy volunteers</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>(n = 44)</td>
</tr>
<tr>
<td>Age (years), median (IQR)</td>
<td>30.0 (26, 42)</td>
</tr>
<tr>
<td>Weight (kg), median (IQR)</td>
<td>70 (60, 80)</td>
</tr>
<tr>
<td>Height (cm), median (IQR)</td>
<td>174 (166, 182)</td>
</tr>
<tr>
<td>BMI (kg/m²), median (IQR)</td>
<td>22.6 (21.2, 24.9)</td>
</tr>
<tr>
<td>Sex (female), n</td>
<td>23</td>
</tr>
<tr>
<td>Systolic blood pressure (mmHg), median (IQR)</td>
<td>122 (116, 128)</td>
</tr>
<tr>
<td>Diastolic blood pressure (mmHg), median (IQR)</td>
<td>77 (70, 82)</td>
</tr>
<tr>
<td>Heat rate (1/min), median (IQR)</td>
<td>63 (58, 73)</td>
</tr>
</tbody>
</table>

IQR = Interquartile range; n = number. Data are presented as absolute number (n) and median (IQR: 25<sup>th</sup>, 75<sup>th</sup>).
Table 2: Laboratory values before and after osmotic stimulus

<table>
<thead>
<tr>
<th>Laboratory values</th>
<th>Healthy volunteers (n = 44)</th>
<th></th>
<th></th>
<th>p value</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Before stimulus</td>
<td>After stimulus</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sodium (mmol/l), median IQR; range</td>
<td>141 (140, 142; 136, 147)</td>
<td>151 (150, 153; 145, 154)</td>
<td></td>
<td>&lt;0.0001</td>
</tr>
<tr>
<td>Osmolality (mmol/kg), median IQR; range</td>
<td>295 (291, 298; 281, 306)</td>
<td>315 (312, 317; 304, 325)</td>
<td></td>
<td>&lt;0.0001</td>
</tr>
<tr>
<td>Copeptin (pg/l), median IQR; range</td>
<td>4.3 (3.3, 6.7; 1.1, 21.4)</td>
<td>28.8 (19.9, 43.4; 19.9, 43.4)</td>
<td></td>
<td>&lt;0.0001</td>
</tr>
<tr>
<td>IL-8 (pg/ml), median IQR; range</td>
<td>0.79 (0.65, 1.11; 0.37, 1.6)</td>
<td>0.7 (0.57, 0.9; 0.4, 1.9)</td>
<td></td>
<td>0.09</td>
</tr>
<tr>
<td>TNF-a (pg/ml), median IQR; range</td>
<td>0.53 (0.39, 0.69; 0.11, 1.1)</td>
<td>0.45 (0.3, 0.53; 0.12, 0.97)</td>
<td></td>
<td>0.036</td>
</tr>
</tbody>
</table>

IQR = Interquartile range; n = number. Data are presented as median (IQR: 25th, 75th; range: min, max).

Two group comparisons were made using the Wilcoxon rank sum test. Bold p-values represent statistically significant results.
Table 3: Circulating markers of systemic inflammation upon osmotic stimulus

<table>
<thead>
<tr>
<th>Inflammatory marker</th>
<th>Multiple linear regression model*</th>
<th>Mixed effect model*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Effect estimate of osmotic stimulus</td>
<td>p-value</td>
</tr>
<tr>
<td>IL-8 (pg/l)</td>
<td>-0.08</td>
<td>0.3</td>
</tr>
<tr>
<td>TNF-α (pg/l)</td>
<td>-0.1</td>
<td><strong>0.03</strong></td>
</tr>
<tr>
<td>IL-6 (pg/l)</td>
<td>0.1</td>
<td>0.7</td>
</tr>
</tbody>
</table>

IL-8 = interleukin-8; TNF-α = tumor necrosis factor-α, IL-6 = interleukin-6.

*Adjusted for age and sex. *With timepoint (before and after stimulus), age and sex as fixed effects and patients as random effect. Bold p-values represent statistically significant results.
Figure 1: Copeptin before and after osmotic stimulus

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Figure 2: IL-8 before and after osmotic stimulus
Figure 3: TNF-α before and after osmotic stimulus

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