# Gene expression in sepsis and sterile sirs in pediatric patients after cardiac surgery

Published: 09-09-2009 Last updated: 17-08-2024

To differentiate between gene expression in sepsis and sterile sirs.

**Ethical review** Approved WMO

**Status** Recruitment stopped

**Health condition type** Congenital cardiac disorders **Study type** Observational non invasive

# **Summary**

## ID

**NL-OMON32059** 

#### Source

**ToetsingOnline** 

#### **Brief title**

Gene expression in sirs and sepsis

#### **Condition**

- Congenital cardiac disorders
- · Helminthic disorders

#### **Synonym**

blood stream infection, sepsis

#### Research involving

Human

# **Sponsors and support**

**Primary sponsor:** Leids Universitair Medisch Centrum

Source(s) of monetary or material Support: Ministerie van OC&W

#### Intervention

**Keyword:** cardiac surgery, gene expression, sepsis, sirs

#### **Outcome measures**

## **Primary outcome**

To differentiate between sepsis and sterile SIRS by means of gene expression.

# **Secondary outcome**

Translate the genes that are expressed in sepsis into a protein which may be used as a diagnostic test in the future.

To reduce the unnecessary use of antibiotics in patients after cardiac surgery

To prohibit the adverse effect of extensive use of antibiotics such as

bacterial resistance or developing superinfection as a result of

microbiological inbalance.

To reduce procedures which are a result of suspecting an infectious cause, such as unnecessarily changing CVC\*s

# **Study description**

# **Background summary**

Because of the high risk of infection and endocarditis in patients after cardiac surgery,

there is a low threshold for starting antibiotics when infection is suspected, or changing the CVC when catheter related infection is suspected. Systemic Inflammatory Response Syndrome (SIRS) appears similar clinically, irrespective if it is caused by an infectious (sepsis) or non-infectious (sterile SIRS) etiology.

In pediatric patients after cardiac surgery there is a low threshold for the use of antibiotics when infection is suspected. However, in a lot of cases no causative microorganism can be demonstrated. Probably, the symptoms are reflecting SIRS, which can for example, be the result of using cardiopulmonary bypass during the procedure.

Often unnecessary antibiotics will be started. Extensive use of antibiotics might cause adverse effects such as bacterial resistance and infections with candida or aspergillus species.

Every seven days, the central venous catheter is electively changed, although in literature there is no evidence found for this routine. In case of suspected (blood stream) infection, often the CVC will be replaced, which can lead to mechanical or other omplications, and also another vessel might be sacrificed due to obstruction.

Despite the similar phenotypic expression, genotypic expression may differ in case of different etiologies causing SIRS. This hypothesis is verified by several studies in critical ill adults with all kinds of underlying diseases. Besides the arguments mentioned above, differentiation between sepsis and sterile SIRS is important considering the fact that the mortality of sepsis is much higher than that of uninfected SIRS.

One study showed a unique pattern of gene expression in the neonatal myocardium of neonates after cardiac surgery for right heart obstructive lesions. However it\*s not clear whether this is age dependent, or reflecting underlying disease.

In several other publications (in adults), gene expression profiles after cardiopulmonary bypass and cardioplegic arrest were described.

## Study objective

To differentiate between gene expression in sepsis and sterile sirs.

## Study design

This is an observational explorative study that will be conducted at the operating room (OR) and pediatric intensive care unit (PICU) at LUMC, Leiden, the Netherlands.

20 patients will be enrolled in the study, undergoing cardiac surgery with bypass for congenital heart defects.

Residual blood from arterial samples will be collected on the OR (moment of study entry), immediate after admittance on the PICU, 12, 24 and 48 hours later. Taking blood samples at these timepoints is routine patient practice. Genetic expression will be analyzed, in order to investigate if there is a differencein expression in patients with SIRS or Sepsis.

The diagnosis of Sepsis and SIRS are based on the clinical criteria as defined by the Members of the International consensus conference on pediatric sepsis.

## Study burden and risks

not applicable

# **Contacts**

#### **Public**

Leids Universitair Medisch Centrum

albinusdreef 2 2300 RC Leiden Nederland **Scientific** 

Leids Universitair Medisch Centrum

albinusdreef 2 2300 RC Leiden Nederland

# **Trial sites**

# **Listed location countries**

**Netherlands** 

# **Eligibility criteria**

## Age

Children (2-11 years)

# **Inclusion criteria**

Pediatric patients who need surgery for congenital heartdefects, and will need cardiopulmonary bypass.

# **Exclusion criteria**

steroid administration, participation in other crt

# Study design

# **Design**

Study type: Observational non invasive

Masking: Open (masking not used)

Control: Uncontrolled

Primary purpose: Diagnostic

#### Recruitment

NL

Recruitment status: Recruitment stopped

Start date (anticipated): 01-03-2010

Enrollment: 20

Type: Actual

# **Ethics review**

Approved WMO

Date: 09-09-2009

Application type: First submission

Review commission: METC Leids Universitair Medisch Centrum (Leiden)

# **Study registrations**

# Followed up by the following (possibly more current) registration

No registrations found.

# Other (possibly less up-to-date) registrations in this register

No registrations found.

# In other registers

Register ID

CCMO NL19974.058.07