

STN	Molekulárne diagnostické vyšetrenia in vitro. Špecifikácie na procesy pred vyšetrovaním metabolitov v moči, sére a plazme.	STN P CEN/TS 16945 85 1026
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Molecular in vitro diagnostic examinations - Specifications for pre-examination processes for metabolomics in urine, venous blood serum and plasma

Táto norma obsahuje anglickú verziu európskej normy.
This standard includes the English version of the European Standard.

Táto norma bola oznámená vo Vestníku ÚNMS SR č. 09/16

Táto predbežná STN je určená na overenie. Pripomienky zasielajte ÚNMS SR najneskôr do mája 2018.

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English Version

Molecular in vitro diagnostic examinations - Specifications for pre-examination processes for metabolomics in urine, venous blood serum and plasma

Tests de diagnostic moléculaire in vitro - Spécifications
relatives aux processus préanalytiques pour l'analyse
du métabolome dans l'urine et le sang veineux (sérum
et plasma)

Molekularanalytische in-vitro-diagnostische Verfahren
- Spezifikationen für präanalytische Prozesse für
Metabolomuntersuchungen in Urin, venöses Blutserum
und -plasma

This Technical Specification (CEN/TS) was approved by CEN on 22 March 2016 for provisional application.

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Contents	Page
European foreword.....	3
Introduction	4
1 Scope.....	5
2 Normative references.....	5
3 Terms and definitions	5
4 General Considerations.....	7
5 Urine	8
5.1 Outside the laboratory	8
5.1.1 Urine collection manual.....	8
5.1.2 Transport requirements.....	9
5.2 Inside the laboratory	9
5.2.1 Specimen reception.....	9
5.2.2 Storage requirements.....	9
5.2.3 Urine sample processing	10
5.2.4 Long-term storage requirements for urine samples.....	10
5.2.5 Urine thawing.....	10
6 Blood.....	10
6.1 Outside the laboratory	10
6.1.1 Primary blood collection manual.....	10
6.1.2 Transport of pre-processed specimens to laboratory.....	12
6.2 Inside the laboratory	12
6.2.1 Specimen reception.....	12
6.2.2 Sample processing	12
6.2.3 Transport of processed samples to a laboratory for metabolomics analysis or transport to a biobank	12
6.2.4 Long-term storage requirements.....	13
6.2.5 Serum and plasma thawing and use	13
Annex A (informative) Long-term stability of urine and serum ¹ H NMR metabolic profiles.....	14
A.1 General.....	14
A.2 Urine ¹ H NMR measurement result	14
A.3 Serum ¹ H NMR measurement result.....	16
A.4 NMR methods for urine and serum	16
Bibliography.....	18

European foreword

This document (CEN/TS 16945:2016) has been prepared by Technical Committee CEN/TC 140 "In vitro diagnostic and medical devices", the secretariat of which is held by DIN.

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Introduction

Molecular *in vitro* diagnostics has enabled a significant progress in medicine. Further progress is expected by new technologies analysing signatures of nucleic acids, proteins, and metabolites in human tissues and body fluids. However, the profiles of these molecules can change drastically during primary sample collection, transport, storage, and processing thus introducing biases and making the outcome from diagnostics or research unreliable or even impossible because the subsequent analytical assay will not determine the situation in the patient but an artificial profile generated during the pre-examination process. Therefore, a standardization of the entire process from sample collection to metabolomics analysis is needed. Studies have been undertaken to determine the important influencing factors. This Technical Specification draws upon such work to codify and standardize the steps for urine, serum and plasma metabolomics analysis in what is referred to as the preanalytical phase.

Metabolomics, the global profiling of metabolites (namely molecules with a molecular weight $MW \leq 2\,000$ Da [3]) in biological samples, is the determination of the dynamic multi-parametric metabolic response of living systems to pathophysiological stimuli and/or genetic modification. Metabolomics studies, which can be semiquantitative or quantitative, help in identifying metabolic profiles that are characteristic for given pathological conditions, for disease prognosis, for the evaluation of the individual response to medical intervention and pharmaceutical treatments. Metabolites are physically and chemically different, and include e.g. sugars, acids, bases, and lipids [3]. This diversity of metabolites and the dynamic range of their concentration in biological samples complicate the separation and detection methods and make it impossible to identify all the metabolites in a single experiment. However, new high-throughput technologies based on NMR (nuclear magnetic resonance) spectroscopy and MS (mass spectrometry) hold great potential due to their ability to look at large parts of the whole metabolome, although with different sensitivity. These two main analytical platforms are now well standardized. Equally well established are the statistical approaches needed to extract information from the huge amount of data resulting from metabolomic analysis.

The metabolic profiles are very sensitive to preanalytical variations that can result from enzymatic activity in the samples and chemical reactions (e.g. oxidation, [4], [5]). This Technical Specification series provides guidelines arising from systematic studies conducted on the most commonly employed biofluids: urine and blood derivatives, serum and plasma.

1 Scope

This Technical Specification covers the preanalytical phase and recommends the handling, documentation and processing of urine, venous blood plasma and serum intended for metabolomics analysis. This Technical Specification is applicable to metabolomics examinations and is of importance to biomedical laboratories, customers of laboratories, *in vitro* diagnostics developers and manufacturers, institutions and companies performing biomedical research, biobanks, and regulatory authorities.

The adoption of the described procedures for the preanalytical phase make it possible to compare and evaluate the results obtained from metabolic profiling analysis.

2 Normative references

The following documents, in whole or in part, are normatively referenced in this document and are indispensable for its application. For dated references, only the edition cited applies. For undated references, the latest edition of the referenced document (including any amendments) applies.

EN ISO 15189:2012, *Medical laboratories - Requirements for quality and competence (ISO 15189:2012, Corrected version 2014-08-15)*

ISO 15190, *Medical laboratories — Requirements for safety*

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