



What analytical tests to order and when



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Outline

• Analytical context – What and how?

- ➢ Prohibited list ≠ target compounds
- Methods and instruments of choice



• Routine test menus – Why not just to analyse everything at once?

- > Complementary information from various sample matrices
- Capabilities and limitations of mass spectrometry in analytical routine



- Extended test menus What to add, when and how?
 - > Evaluation of an altered biomarker vs. identification of an exogenous substance
 - > Systematic review of available data for correct analytical test







Prohibited substances - IC only



Prohibited substances - IC and OOC

Endogenous AAS



Metabolism – Phase-I



Metabolism – Phase-II



Analytical procedures – few tests, wide coverage



Analytical procedures – criteria and limitations

5.2.4.3.1.2 Mass spectrometry (MS) coupled to either gas (GC) or liquid chromatography (LC) is the analytical technique of choice for confirmation of *Prohibited Substances*, *Metabolite(s)* of *Prohibited Substance(s)*, or *Marker(s)* of the *Use* of a *Prohibited Substance* or *Prohibited Method*. GC or High Performance Liquid Chromatography (HPLC) coupled with MS or MS-MS are acceptable for both Initial Testing Procedures and Confirmation Procedures for a specific analyte.



Additional tests e.g. GHRFs, IGF-1 analogs, insulins and GC/C/IRMS

5.2.4.3.1.3 Affinity Binding Assays (e.g. Immunoassays) are also routinely used for detection of macromolecules in urine samples. Affinity Binding Assays applied for the <u>Initial Testing</u> <u>Procedures</u> and <u>Confirmation Procedures</u> shall use affinity reagents (e.g. antibodies) recognizing different epitopes of the macromolecule analyzed, unless a purification or separation method is used prior to application of the Affinity Binding Assay to eliminate the potential of cross-reactivity. The <u>Laboratory</u> shall document, as part of the method validation, the <u>Fitness-forpurpose</u> of any such purification or separation method.

assays) are es in urine tial Testing use affinity opes of the separation







Atypical passport finding – data review

- Is the sample valid?
 - > Blood stability score, microbial contamination
 - Confounding factors (CF) and adverse analytical findings (AAF)
 - Contact the laboratory for further information
- Are the data correct?
 - > Analytical data introduced to correct sample (laboratory)
 - Sample assigned to correct BPID (SCA/TA)
 - Steroid profile confirmation procedure
- What has been already done?
 - Which samples were collected?
 - > Test menus requested per each sample matrix?
 - Request the sample to be transferred to long-term storage

















Atypical passport finding – need of additional tests

- What is the scenario?
 - Sharing of information with SCA/DCO and laboratory; I&I
 - > Data not available for passport evaluation manipulated sample
 - Use of prohibited substance(s) additional tests needed for identification



- > Urine integrity:
 - > pH, specific gravity, steroid profile (included in the routine test menu)
 - creatinine, salt analysis, total protein, proteases (outside of routine testing)
- Sample identity
 - DNA-analysis requires a reference sample (outside of routine testing)

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      Int J Leg Med (2006)
DOI 10.1007/5001141
      Research article
      Drug 1 esting
and Analysis

      ORIGINAL AT
Notice 19 May 2014
      Noted 23 January 2015
      Noted 23 January 2015
      Noted 23 January 2015
      Noted 21 October 2016
      Acapted 21 October 2016
      Noted 21 October 2016
      Noted
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11

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28 31

12 15

Additional tests for steroid module – S1.1b

- Steroid profile confirmation procedure
 - GC-MS-analysis: repeated analysis of steroid profile parameters
 - > Deeper insight also to microbial contamination
 - > GC/C/IRMS-analysis: exogenous/endogenous origin of testosterone metabolites





- For suppressed steroid profiles
- > Direct analysis of glucuronide- and sulpho-conjugates





Additional tests for hematological module – S2.1 / M1

- Erythropoiesis stimulating agents (ESAs)
 - SAR-PAGE-based methods
 - Limited number of specific MS-methods (e.g. Hematide)
 - Selection of matrix; serum often better (e.g. CERA)

- Agents activating hypoxia-inducible factor (HIF)
 - Molidustat, roxadustat (FG-4592)
 - Cobalt
 - Xenon

Detection of blood transfusions

- Phthalates by MS-methods in urine
- Flow cytometric analysis for homologous transfusions
- DNA-analyses for mixed populations





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Conclusions

• Routine analytical testing

- > Preference of mass spectrometric methods to detected relevant target compounds
- > Arranged according to instrumentation and capacity by each laboratory
- Wide coverage, but specific methods applied only upon additional analysis requests

Atypical passport profile – data review

- Verify the correctness of analytical data
- Validate the connection between the results and appropriate BPID
- > Check the availability of various sample matrices collected in the test

Additional tests

- Scenario: sample manipulation or use of prohibited substances?
- > Steroid module: S1.1b substances, Hematological module: S2.1 and M1
- > Discuss with the laboratory, transfer the sample for long term storage



kde.mitre.org



> Dr. Hans Geyer (Cologne)



- > Dr. Norbert Baume & Dr. Sylvain Giraud
- World Association of Anti-Doping Scientists (WAADS)
- LAD team















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