ab185435

Tyrosine Assay Kit
(Colorimetric)

Instructions for Use

For the rapid, sensitive and accurate measurement of Tyrosine in biological samples.

View kit datasheet: www.abcam.com/ab185435
(use www.abcam.cn/ab185435 for China, or www.abcam.co.jp/ab185435 for Japan)

This product is for research use only and is not intended for diagnostic use.

Version: 1 Last Updated: 1 February 2019
# Table of Contents

1. Overview ........................................... 3
2. Protocol Summary .................................. 4
3. Kits Components .................................. 5
4. Storage and Stability ............................. 6
5. Materials Required, Not Supplied ................. 6
6. Reagents Preparation ............................... 7
7. Assay Protocol ..................................... 8
8. Data Analysis ..................................... 11
9. Troubleshooting ................................... 14
1. Overview

Tyrosine (Tyr) is one of the four standard amino acids containing an aromatic group as a side chain. Its hydrophobicity is one of the main characteristics of this uncharged polar amino acid. In addition to being an essential amino acid Tyr is important in number of biological processes such as the synthesis of neurotransmitters, thyroid hormones, melanine, fumarate and acetoacetate. The pathology of abnormal concentrations of Tyr is well known in diseases including phenylketonuria, hypothyroidism, tyrosinemia, albinism, and alkaptonuria.

Abcam's Tyrosine Assay Kit (Colorimetric) (ab185435) is a simple, yet sensitive assay that is able to detect normal and abnormal concentrations of Tyrosine in biological fluids. The assay is based on the enzymatic oxidation of Tyrosine producing a stable signal (OD 492 nm), which is directly proportional to the amount of Tyrosine. Sample preparation is minimal and does not require strenuous or complicated procedures. The assay can detect as low as 50 µM of Tyrosine in a variety of biological samples.

Figure 1: Assay Procedure

```
Tyrosine  Enzyme Mix  Intermediate  Absorbance (OD 492 nm)
```
2. Protocol Summary

- Reagent Preparation
  - Standard Curve Preparation
    - Control and Samples Preparation
      - Reaction
        - Measure Colored substrate and Calculation
3. **Kits Components**

<table>
<thead>
<tr>
<th>Item</th>
<th>Quantity</th>
</tr>
</thead>
<tbody>
<tr>
<td>Tyr Assay Buffer</td>
<td>25 mL</td>
</tr>
<tr>
<td>Tyr Enzyme Mix</td>
<td>1 vial</td>
</tr>
<tr>
<td>Tyr Standard</td>
<td>1 vial</td>
</tr>
</tbody>
</table>
4. Storage and Stability

Upon arrival, store the kit at -20°C and protect from light. Please read the entire protocol before performing the assay. Avoid repeated freeze/thaw cycles.

Briefly centrifuge all small vials prior to opening.

5. Materials Required, Not Supplied

- Distilled water (dH₂O)
- 96-well clear plate with flat bottom
- Multi-well spectrophotometer (ELISA reader)
- Centrifuge with cooling option
- 10kDa spin column (ab93349)
6. Reagents Preparation

1. Tyrosine Assay Buffer:
   Ready to use as supplied. Aliquot and store at +4°C or -20°C.
   Warm to room temperature before use.

2. Tyrosine Enzyme Mix:
   Reconstitute with 220 µL Tyrosine Assay Buffer. Pipette up
   and down to dissolve completely. Aliquot and store at -20°C.
   Use within two months.

3. Tyrosine Standard
   Reconstitute with 100 µL dH₂O to generate 100 mM solution.
   Store at 4°C. Stable for two months.
7. Assay Protocol

1. Sample Preparation
   a. Samples should be deproteinized using 10 kDa Spin Column (ab93349).
   b. Briefly, add sample to the spin column, centrifuge at 10,000 x g for 10 minutes at 4°C. Collect the filtrate.
   c. Add 80-135 µL of filtrate into desired well(s) in 96-well plate.
   d. Adjust the volume to 150 µL/well with dH₂O.

NOTE:

- Tyrosine concentrations can vary over a wide range. Normal ranges in Humans are 55-147 µM for serum, 10-290 µM for urine and 34-112 µM in plasma. Abnormal tyrosine levels can exceed 1.5 mM in tyrosinemia samples. For unknown samples, we recommend doing pilot experiment and testing several doses to ensure the readings are within the Standard Curve range.
- Endogenous compounds may interfere with the assay. To ensure accurate determination of Tyr in the test samples or for samples having low concentration of Tyr, we recommend spiking samples with a known amount of Tyr Standard (30 nmol).
• For samples having background, prepare parallel sample well(s) as background control(s).

2. Standard Curve Preparation:
   a) Dilute the Tyrosine Standard to 2.5 mM by adding 25 µL of 100 mM Tyrosine Standard to 975 µL of dH₂O.
   b) Add 0, 2, 6, 12, 18, 24 and 30 µL of Tyrosine Standard into series of wells in a 96-well plate to generate 0, 5, 15, 30, 45, 60, and 75 nmol/well of Tyrosine Standard.
   c) Adjust the volume to 150 µL/well with dH₂O. The diluted Standard can be stored at 4°C for subsequent assays.
3. Reaction Mix:

Prepare enough Reaction Mix for the number of assays to be performed. For each well, prepare 50 µL Mix containing:

<table>
<thead>
<tr>
<th></th>
<th>Reaction Mix</th>
<th>*Background Control Mix</th>
</tr>
</thead>
<tbody>
<tr>
<td>Tyrosine Assay Buffer</td>
<td>48 µL</td>
<td>50 µL</td>
</tr>
<tr>
<td>Tyrosine Enzyme Mix</td>
<td>2 µL</td>
<td>--</td>
</tr>
</tbody>
</table>

Add 50 µL of the reaction mix to each well containing the Standards and samples.

*For samples having high background, and 50 µl of Background Control mix to sample background control well(s). Mix well.

4. Measurement

a) Incubate the plate at room temperature for 60 minutes protected from light.

b) Measure absorbance (OD 492 nm) in a microplate reader.
8. Data Analysis

Calculations:

a) Subtract 0 Standard reading from all readings. If sample background control reading is significant then subtract the sample background control reading from sample reading. Plot the Tyr Standard Curve. For unspiked samples, apply the corrected OD to the Tyr Standard Curve to get B nmol of Tyr in the sample well.

Sample Tyr concentration (C) = B/V X D nmol/ml or µM

Where:

B is the amount of Tyr in the sample well from the standard curve (nmol)
V is the sample volume added into the reaction well (ml)
D is the sample dilution factor

Note: For spiked samples, correct for any sample interference by subtracting the sample reading from spiked sample reading.

For spiked samples, Tyr amount in sample well (B):

\[
\left( \frac{OD_{\text{sample (corrected)}}}{OD_{\text{sample + Tyr Std (corrected)}} - OD_{\text{sample (corrected)}}} \right) \times Tyr \ Spike (\text{nmol})
\]

Tyrosine molecular weight: 181.2 g/mol.
Figure 2. Tyrosine Standard Curve.
Figure 3. Measurement of Tyrosine concentration in Human urine and serum (135 µL, each). Both samples were deproteinized using 10 kDa Spin Column (ab93349) and spiked with known amount of Tyrosine (30 nmol). Assays were performed following the kit protocol.
## 9. Troubleshooting

<table>
<thead>
<tr>
<th>Problem</th>
<th>Reason</th>
<th>Solution</th>
</tr>
</thead>
<tbody>
<tr>
<td>Assay not working</td>
<td>Assay buffer at wrong temperature</td>
<td>Assay buffer must not be chilled - needs to be at RT</td>
</tr>
<tr>
<td>Protocol step missed</td>
<td></td>
<td>Re-read and follow the protocol exactly</td>
</tr>
<tr>
<td>Plate read at incorrect wavelength</td>
<td></td>
<td>Ensure you are using appropriate reader and filter settings (refer to datasheet)</td>
</tr>
<tr>
<td>Unsuitable microtiter plate for assay</td>
<td></td>
<td>Fluorescence: Black plates (clear bottoms); Luminescence: White plates; Colorimetry: Clear plates. If critical, datasheet will indicate whether to use flat- or U-shaped wells</td>
</tr>
<tr>
<td>Unexpected results</td>
<td>Measured at wrong wavelength</td>
<td>Use appropriate reader and filter settings described in datasheet</td>
</tr>
<tr>
<td>Samples contain impeding substances</td>
<td></td>
<td>Troubleshoot and also consider deproteinizing samples</td>
</tr>
<tr>
<td>Unsuitable sample type</td>
<td></td>
<td>Use recommended samples types as listed on the datasheet</td>
</tr>
<tr>
<td>Sample readings are outside linear range</td>
<td></td>
<td>Concentrate/ dilute samples to be in linear range</td>
</tr>
<tr>
<td>Problem</td>
<td>Reason</td>
<td>Solution</td>
</tr>
<tr>
<td>--------------------------------------------</td>
<td>-----------------------------------------------</td>
<td>--------------------------------------------------------------------------</td>
</tr>
<tr>
<td>Samples with inconsistent readings</td>
<td>Unsuitable sample type</td>
<td>Refer to datasheet for details about incompatible samples</td>
</tr>
<tr>
<td></td>
<td>Samples prepared in the wrong buffer</td>
<td>Use the assay buffer provided (or refer to datasheet for instructions)</td>
</tr>
<tr>
<td></td>
<td>Samples not deproteinized (if indicated on datasheet)</td>
<td>Use the <strong>10kDa spin column (ab93349)</strong></td>
</tr>
<tr>
<td></td>
<td>Cell/tissue samples not sufficiently homogenized</td>
<td>Increase sonication time/number of strokes with the Dounce homogenizer</td>
</tr>
<tr>
<td></td>
<td>Too many freeze-thaw cycles</td>
<td>Aliquot samples to reduce the number of freeze-thaw cycles</td>
</tr>
<tr>
<td></td>
<td>Samples contain impeding substances</td>
<td>Troubleshoot and also consider deproteinizing samples</td>
</tr>
<tr>
<td></td>
<td>Samples are too old or incorrectly stored</td>
<td>Use freshly made samples and store at recommended temperature until use</td>
</tr>
<tr>
<td>Lower/Higher readings in samples and standards</td>
<td>Not fully thawed kit components</td>
<td>Wait for components to thaw completely and gently mix prior use</td>
</tr>
<tr>
<td></td>
<td>Out-of-date kit or incorrectly stored reagents</td>
<td>Always check expiry date and store kit components as recommended on the datasheet</td>
</tr>
<tr>
<td></td>
<td>Reagents sitting for extended periods on ice</td>
<td>Try to prepare a fresh reaction mix prior to each use</td>
</tr>
<tr>
<td></td>
<td>Incorrect incubation time/temperature</td>
<td>Refer to datasheet for recommended incubation time and/or temperature</td>
</tr>
<tr>
<td>Problem</td>
<td>Reason</td>
<td>Solution</td>
</tr>
<tr>
<td>-----------------------------</td>
<td>-----------------------------------------</td>
<td>--------------------------------------------------------------------------</td>
</tr>
<tr>
<td>Incorrect amounts used</td>
<td>Check pipette is calibrated correctly (always use smallest volume pipette that can pipette entire volume)</td>
<td></td>
</tr>
<tr>
<td>Standard curve is not linear</td>
<td>Not fully thawed kit components</td>
<td>Wait for components to thaw completely and gently mix prior use</td>
</tr>
<tr>
<td></td>
<td>Pipetting errors when setting up the standard curve</td>
<td>Try not to pipette too small volumes</td>
</tr>
<tr>
<td></td>
<td>Incorrect pipetting when preparing the reaction mix</td>
<td>Always prepare a master mix</td>
</tr>
<tr>
<td></td>
<td>Air bubbles in wells</td>
<td>Air bubbles will interfere with readings; try to avoid producing air bubbles and always remove bubbles prior to reading plates</td>
</tr>
<tr>
<td></td>
<td>Concentration of standard stock incorrect</td>
<td>Recheck datasheet for recommended concentrations of standard stocks</td>
</tr>
<tr>
<td></td>
<td>Errors in standard curve calculations</td>
<td>Refer to datasheet and re-check the calculations</td>
</tr>
<tr>
<td></td>
<td>Use of other reagents than those provided with the kit</td>
<td>Use fresh components from the same kit</td>
</tr>
</tbody>
</table>
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