ab102523

Amylase Assay kit (Colorimetric)

Instructions for Use

For the rapid, sensitive and accurate measurement of Amylase activity in various samples.

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

Amylase Assay kit (Colorimetric) (ab102523) detects activity of α-amylase through a two-step reaction. α-Amylase will cleave the substrate ethylidene-\(\rho\)NP-G7 to produce smaller fragments that are eventually modified by α-glucosidase, causing the release of a chromophore that can then be measured at OD = 405 nm. The assay can detect α-amylase content as low as 0.2 mU.

Amylases are enzymes that break starch down to sugar molecules. α-amylase is the major form of amylase found in humans and other mammals as well as an enzyme present in seeds, or in fungi (baker’s yeast for instance). α-amylase is a calcium metalloenzyme, completely unable to function in the absence of calcium.

In human physiology, both the salivary and pancreatic amylases are major digestive enzymes. Increased enzyme levels in humans are associated with salivary trauma; mumps due to inflammation of the salivary glands, pancreatitis and renal failure. A simple, direct and automation-ready procedure for measuring α-amylase activity is, therefore, very desirable.
2. **ASSAY SUMMARY**

Sample preparation

↓

Standard curve preparation

↓

Add reaction mix

↓

Measure optical density (OD405 nm) in a kinetic mode at 25°C for 30 – 60 min

*For kinetic mode detection, incubation time given in this summary is for guidance only.*
3. **PRECAUTIONS**

Please read these instructions carefully prior to beginning the assay.

All kit components have been formulated and quality control tested to function successfully as a kit. Modifications to the kit components or procedures may result in loss of performance.

4. **STORAGE AND STABILITY**

Store kit at -20ºC in the dark immediately upon receipt. Kit has a storage time of 1 year from receipt, providing components have not been reconstituted.

Refer to list of materials supplied for storage conditions of individual components. Observe the storage conditions for individual prepared components in section 5.

Aliquot components in working volumes before storing at the recommended temperature. **Reconstituted components are stable for 2 months.**
5. **MATERIALS SUPPLIED**

<table>
<thead>
<tr>
<th>Item</th>
<th>Amount</th>
<th>Storage Condition (Before Preparation)</th>
<th>Storage Condition (After Preparation)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Amylase Assay Buffer</td>
<td>55 mL</td>
<td>-20°C</td>
<td>-20°C</td>
</tr>
<tr>
<td>Amylase Substrate Mix</td>
<td>5 mL</td>
<td>-20°C</td>
<td>-20°C</td>
</tr>
<tr>
<td>Amylase Positive Control</td>
<td>1 vial</td>
<td>-20°C</td>
<td>-20°C</td>
</tr>
<tr>
<td>(Lyophilized)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Nitrophenol Standard (2mM)</td>
<td>150 µL</td>
<td>-20°C</td>
<td>-20°C</td>
</tr>
</tbody>
</table>

6. **MATERIALS REQUIRED, NOT SUPPLIED**

These materials are not included in the kit, but will be required to successfully utilize this assay:

- MilliQ water or other type of double distilled water (ddH₂O)
- PBS
- Microcentrifuge
- Pipettes and pipette tips
- Colorimetric microplate reader – equipped with filter for OD 405 nm
- 96 well plate: clear plate for colorimetric assay
- Orbital shaker
- Vortex
- Dounce homogenizer or pestle (if using tissue)
7. **LIMITATIONS**

- Assay kit intended for research use only. Not for use in diagnostic procedures.
- Do not use kit or components if it has exceeded the expiration date on the kit labels.
- Do not mix or substitute reagents or materials from other kit lots or vendors. Kits are QC tested as a set of components and performance cannot be guaranteed if utilized separately or substituted.
8. **TECHNICAL HINTS**

- This kit is sold based on number of tests. A ‘test’ simply refers to a single assay well. The number of wells that contain sample, control or standard will vary by product. Review the protocol completely to confirm this kit meets your requirements. Please contact our Technical Support staff with any questions.

- Keep enzymes and heat labile components and samples on ice during the assay.

- Make sure all buffers and developing solutions are at room temperature before starting the experiment.

- Avoid cross contamination of samples or reagents by changing tips between sample, standard and reagent additions.

- Avoid foaming or bubbles when mixing or reconstituting components.

- Samples generating values higher than the highest standard should be further diluted in the appropriate sample dilution buffers.

- Ensure plates are properly sealed or covered during incubation steps.

- Ensure complete removal of all solutions and buffers from tubes or plates during wash steps.

- Make sure you have the appropriate type of plate for the detection method of choice.

- Make sure the heat block/water bath and microplate reader are switched on before starting the experiment.
9. **REAGENT PREPARATION**

- Briefly centrifuge small vials at low speed prior to opening.

9.1 **Amylase Assay Buffer:**

   Ready to use as supplied. Equilibrate to room temperature before use. Store at -20°C.

9.2 **Amylase Substrate Mix:**

   Ready to use as supplied. Keep on ice during the assay. Aliquot substrate so that you have enough to perform the desired number of tests. Store at -20°C.

9.3 **Amylase Positive Control:**

   Dissolve Amylase Positive Control into 50 µL Assay Buffer. Aliquot positive control so that you have enough to perform the desired number of tests. Keep positive control on ice during the assay. Store at -20°C.

9.4 **Nitrophenol Standard (2 mM):**

   Ready to use as supplied. Aliquot standard so that you have enough to perform the desired number of tests. Store at -20°C.
10. STANDARD PREPARATION

- Always prepare a fresh set of standards for every use.
- Diluted standard solution is unstable and must be used within 4 hours.

10.1 Using 2mM Nitrophenol standard, prepare standard curve dilution as described in the table in a microplate or microcentrifuge tubes:

<table>
<thead>
<tr>
<th>Standard #</th>
<th>Volume of Standard (µL)</th>
<th>Assay Buffer (µL)</th>
<th>Final volume standard in well (µL)</th>
<th>End [amylase] in well</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>0</td>
<td>150</td>
<td>50</td>
<td>0 nmol/well</td>
</tr>
<tr>
<td>2</td>
<td>6</td>
<td>144</td>
<td>50</td>
<td>4 nmol/well</td>
</tr>
<tr>
<td>3</td>
<td>12</td>
<td>138</td>
<td>50</td>
<td>8 nmol/well</td>
</tr>
<tr>
<td>4</td>
<td>18</td>
<td>132</td>
<td>50</td>
<td>12 nmol/well</td>
</tr>
<tr>
<td>5</td>
<td>24</td>
<td>126</td>
<td>50</td>
<td>16 nmol/well</td>
</tr>
<tr>
<td>6</td>
<td>30</td>
<td>120</td>
<td>50</td>
<td>20 nmol/well</td>
</tr>
</tbody>
</table>

Each dilution has enough amount of standard to set up duplicate reading (2 x 50 µL).
11. **SAMPLE PREPARATION**

**General Sample information:**

- We recommend performing several dilutions of your sample to ensure the readings are within the standard value range.
- We recommend that you use fresh samples. If you cannot perform the assay at the same time, we suggest that you complete the Sample Preparation step before storing the samples. Alternatively, if that is not possible, we suggest that you snap freeze cells or tissue in liquid nitrogen upon extraction and store the samples immediately at -80°C. When you are ready to test your samples, thaw them on ice. Be aware however that this might affect the stability of your samples and the readings can be lower than expected.

11.1 **Cell (adherent or suspension) samples:**

11.1.1 Harvest the amount of cells necessary for each assay (initial recommendation = 4 x 10^6 cells).

11.1.2 Wash cells with cold PBS.

11.1.3 Resuspend cells in 500 µL Assay Buffer.

11.1.4 Homogenize cells quickly by pipetting up and down a few times.

11.1.5 Centrifuge sample for 2 – 5 minutes at 4°C at top speed using a cold microcentrifuge to remove any insoluble material.

11.1.6 Collect supernatant and transfer to a clean tube.

11.1.7 Keep on ice.

11.2 **Tissue samples:**

11.2.1 Harvest the amount of cells necessary for each assay (initial recommendation = 100 mg tissue).

11.2.2 Wash tissue in cold PBS.

11.2.3 Resuspend tissue in 500 µL Assay Buffer.
11.2.4 Homogenize tissue with a Dounce homogenizer sitting on ice, with 10 – 15 passes.

11.2.5 Centrifuge samples for 2 – 5 minutes at 4°C at top speed, using a cold microcentrifuge to remove any insoluble material.

11.2.6 Collect supernatant and transfer to a clean tube.

11.2.7 Keep on ice.

11.3 **Serum and Urine:**

Serum and urine samples can be tested directly by adding sample to the microplate wells.

However, to find the optimal values and ensure your readings will fall within the standard values, we recommend performing several dilutions of the sample (1/2 – 1/5 – 1/10).

**NOTE:** We suggest using different volumes of sample to ensure readings are within the Standard Curve range.
12. ASSAY PROCEDURE and DETECTION

- Equilibrate all materials and prepared reagents to room temperature prior to use.
- It is recommended to assay all standards, controls and samples in duplicate.

12.1 Set up Reaction wells:
- Standard wells = 50 µL Standard dilutions.
- Positive control wells – Add 5 µL of Amylase Positive Control and adjust to 50 µL/well with ddH$_2$O.
- Sample wells = 2 – 50 µL samples (adjust volume to 50 µL/well with ddH$_2$O).

12.2 Reaction Mix:

Prepare 100 µL of Reaction Mix for each reaction:

<table>
<thead>
<tr>
<th>Component</th>
<th>Reaction Mix Samples (µL)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Assay Buffer</td>
<td>50</td>
</tr>
<tr>
<td>Substrate Mix</td>
<td>50</td>
</tr>
</tbody>
</table>

Mix enough reagents for the number of assays (samples, standards, positive control and background control) to be performed. Prepare a Master Mix of the Reaction Mix to ensure consistency. We recommend the following calculation:

$X \mu$L component x (Number samples + standards +1)

12.3 Add 100 µL of Reaction Mix into each well. Mix thoroughly.

12.4 Measure absorbance immediately at OD=405 nm in a kinetic mode, every 2 – 3 minutes, for 30 – 60 min at 25°C protected from light.

**NOTE:** Sample incubation time can vary depending on α-amylase activity in samples. We recommend observing the reaction kinetics then choosing two time points (T1 and T2) in the linear range to calculate the amylase activity. The Standard Curve will not change as incubation time increases.
13. **CALCULATIONS**

- Samples producing signals greater than that of the highest standard should be further diluted in appropriate buffer and reanalyzed, then multiplying the concentration found by the appropriate dilution factor.
- For statistical reasons, we recommend each sample should be assayed with a minimum of two replicates (duplicates).

13.1 Average the duplicate reading for each standard and sample.

13.2 Subtract the mean absorbance value of the blank (Standard #1) from all standard and sample readings. This is the corrected absorbance.

13.3 Plot the corrected absorbance values for each standard as a function of the final concentration of Nitrophenol.

13.4 Draw the best smooth curve through these points to construct the standard curve. Most plate reader software or Excel can plot these values and curve fit. Calculate the trendline equation based on your standard curve data (use the equation that provides the most accurate fit).

13.5 Activity of amylase is calculated as:

\[ \Delta A_{405nm} = A_2 - A_1 \]

Where:
- A1 is the sample reading at time T1
- A2 is the sample reading at time T2

13.6 Use the \( \Delta A_{405nm} \) to obtain B nmol of Nitrophenol generated by amylase during the reaction time (\( \Delta T = T_2 - T_1 \)).

13.7 Activity of LDH in the test samples is calculated as:

\[ \text{Amylase Activity} = \left( \frac{B}{\Delta T \times V} \right) \times D \]

Amylase = nmol/min/mL = mU/mL
Where:
B = Nitrophenol amount from the standard curve (in nmol).
ΔT = reaction time (T2 – T1) (min).
V = Pretreated sample volume added to the reaction well (in mL).
D = sample dilution factor.

**Unit Definition:**

1 Unit Amylase = amount of amylase that cleaves ethylidene-\(p\)NP-G7 to generate 1.0 \(\mu\)mol of nitrophenol per min at pH 7.20 at 25 °C.
14. **TYPICAL DATA**

**TYPICAL STANDARD CURVE** – Data provided for demonstration purposes only. A new standard curve must be generated for each assay performed.

![Figure 1: Typical Nitrophenol standard calibration curve using colorimetric reading.](image-url)
Figure 2: Amylase Activity measured in biological fluids showing quantity (mU) per mL of tested sample after 12 min of incubation.
**Figure 3:** Amylase Activity measured in tissue lysates showing quantity (mU) per mg of protein after 12 min of incubation.

**Figure 4:** Amylase Activity measured in cell lysates showing quantity (mU) per 1 mln cells after 12 min of incubation.
15. QUICK ASSAY PROCEDURE

NOTE: This procedure is provided as a quick reference for experienced users. Follow the detailed procedure when performing the assay for the first time.

- Solubilize Amylase Positive Control, thaw the Assay Buffer, Substrate Mix and Nitrophenol standard (aliquot if necessary); get equipment ready.
- Prepare appropriate standard curve.
- Prepare samples in duplicate (find optimal dilutions to fit standard curve readings).
- Set up plate for standard (50 µL) and samples (50 µL).
- Prepare Amylase Reaction Mix (number samples + standards + 1).

<table>
<thead>
<tr>
<th>Component</th>
<th>Reaction Mix (µL)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Assay Buffer</td>
<td>50</td>
</tr>
<tr>
<td>Substrate Mix</td>
<td>50</td>
</tr>
</tbody>
</table>

- Add 100 µL of Reaction Mix to standard and sample wells.
- Measure absorbance at OD=405 nm on a microplate reader in a kinetic mode at 25°C for 30 – 60 min protected from light.
16. **TROUBLESHOOTING**

<table>
<thead>
<tr>
<th>Problem</th>
<th>Cause</th>
<th>Solution</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Assay not working</strong></td>
<td>Use of ice-cold buffer</td>
<td>Buffers must be at room temperature</td>
</tr>
<tr>
<td></td>
<td>Plate read at incorrect wavelength</td>
<td>Check the wavelength and filter settings of instrument</td>
</tr>
<tr>
<td></td>
<td>Use of a different 96-well plate</td>
<td>Colorimetric: Clear plates</td>
</tr>
<tr>
<td></td>
<td></td>
<td>Fluorometric: black wells/clear bottom plate</td>
</tr>
<tr>
<td><strong>Sample with erratic readings</strong></td>
<td>Samples not deproteinized (if indicated on protocol)</td>
<td>Use PCA precipitation protocol for deproteinization</td>
</tr>
<tr>
<td></td>
<td>Cells/tissue samples not homogenized completely</td>
<td>Use Dounce homogenizer, increase number of strokes</td>
</tr>
<tr>
<td></td>
<td>Samples used after multiple free/thaw cycles</td>
<td>Aliquot and freeze samples if needed to use multiple times</td>
</tr>
<tr>
<td></td>
<td>Use of old or inappropriately stored samples</td>
<td>Use fresh samples or store at -80°C (after snap freeze in liquid nitrogen) till use</td>
</tr>
<tr>
<td></td>
<td>Presence of interfering substance in the sample</td>
<td>Check protocol for interfering substances; deproteinize samples</td>
</tr>
<tr>
<td><strong>Lower/Higher readings in samples and Standards</strong></td>
<td>Improperly thawed components</td>
<td>Thaw all components completely and mix gently before use</td>
</tr>
<tr>
<td></td>
<td>Allowing reagents to sit for extended times on ice</td>
<td>Always thaw and prepare fresh reaction mix before use</td>
</tr>
<tr>
<td></td>
<td>Incorrect incubation times or temperatures</td>
<td>Verify correct incubation times and temperatures in protocol</td>
</tr>
<tr>
<td>Problem</td>
<td>Cause</td>
<td>Solution</td>
</tr>
<tr>
<td>---------</td>
<td>-------</td>
<td>----------</td>
</tr>
<tr>
<td>Pipetting errors in standard or reaction mix</td>
<td>Avoid pipetting small volumes (&lt; 5 µL) and prepare a master mix whenever possible</td>
<td></td>
</tr>
<tr>
<td>Air bubbles formed in well</td>
<td>Pipette gently against the wall of the tubes</td>
<td></td>
</tr>
<tr>
<td>Standard stock is at incorrect concentration</td>
<td>Always refer to dilutions described in the protocol</td>
<td></td>
</tr>
<tr>
<td>Standard readings do not follow a linear pattern</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Measured at incorrect wavelength</td>
<td>Check equipment and filter setting</td>
<td></td>
</tr>
<tr>
<td>Samples contain interfering substances</td>
<td>Troubleshoot if it interferes with the kit</td>
<td></td>
</tr>
<tr>
<td>Sample readings above/ below the linear range</td>
<td>Concentrate/ Dilute sample so it is within the linear range</td>
<td></td>
</tr>
<tr>
<td>Unanticipated results</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
17. **FAQ**

**Will this kit work for beta or gamma amylases as well?**
This kit is specific for alpha-amylase activity.

**Is there an upper limit for this kit regarding alpha amylase concentration?**
To be able to calculate Amylase activity in samples, it is essential to get values within the linear range of the standard curve. The highest datapoint for the standard is 20 nmol of Nitrophenol per well. As mentioned on our datasheet, 1 unit of amylase generates 1.0 µmol of nitrophenol per min at pH 7.20 at 25 °C. Hence the maximum amount of enzyme activity within the linear range of the standard curve would be 20 mU per well. The sample amount needed to get values within the linear range of the std. curve is sample dependent and has to be optimized in a pilot assay by the user.
18. **INTERFERENCES**
19. **NOTES**
UK, EU and ROW
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