

Product Information

L-Amino Acid Quantitation Colorimetric/Fluorometric Kit

I. Kit Contents:

| Components | K2141-100 100 assays | Cap Color | Part Number |
|----------------------------------|-------------------------|-----------|-------------|
| L-Amino Acid Assay Buffer | 25 ml | WM | K2141-C-1 |
| L-Amino Assay Probe | 0.2 ml | Red | K2141-C-2 |
| L-Amino Acid Enzyme Mix | 1 Vial | Green | K2141-C-3 |
| L-Amino Acid Standard (4nmol/μl) | 300μl | Yellow | K2141-C-4 |

Note: Mixture of all amino acids at equal molar ratio. Glycine is not detectable in this assay

II. Introduction:

L-amino acids are found in proteins during translation in the ribosome. D-amino acids are produced by enzyme posttranslational modifications in some proteins. Accurately quantitating L-amino acids in purified samples or body fluids can provide useful information for basic research or diagnostic studies.

The L-Amino Acid Quantitation Colorimetric/Fluorometric Kit provides a sensitive, fast and convenient way for detection of L-amino acid in various biological samples based on colorimetric and fluorometric method. The L-amino acid(s) level can be detected using colorimetric (at $\lambda = 570$ nm) or fluorometric (at Ex/Em = 535/587 nm) method in 96-well plates. The assay is performed without requirement for sample pretreatment or purification.

III. Reagent Preparation and Storage Conditions:

Probe: Ready to use as supplied. Warm to room temperature to thaw the DMSO solution before use. Store at -20°C , protect from light. Use within two months.

Enzyme Mix: Dissolve in 220 μl L-Amino Acid Assay Buffer. Pipette up and down to complete dissolve the content. Store at -20°C . Use within two months.

IV. Assay Protocol:

1. Standard Curve Preparations: For the colorimetric assay, add 0, 2, 4, 6, 8, 10 μl L-Amino Acid Standard into a series of wells of a 96-well plate to generate 0, 8, 16, 24, 32, 40 nmol/well of L-Amino Acid Standard. Adjust volume to 50 μl /well with L-Amino Acid Assay Buffer. For the fluorometric assay, dilute the L-Amino Acid to 0.4 nmol/ μl by adding 10 μl of the L-Amino Acid to 90 μl of L-Amino Acid Assay Buffer, mix well. Add 0, 2, 4, 6, 8, 10 μl into each well individually to generate 0, 0.8, 1.6, 2.4, 3.2, 4.0 nmol/well of the L-Amino Acid Standard. Adjust volume to 50 μl /well with L-Amino Acid Assay Buffer.

2. Sample Preparations: Prepare test samples in a final volume of 50 μl /well with L-Amino Acid Assay Buffer in the 96-well plate. We suggest using several doses of your sample to ensure the readings are within the standard curve range.

3. Reaction Mix Preparation: Mix enough reagents for the number of assays performed: For each well, prepare a total 50 μl Reaction Mix containing:

| | |
|---------------------------|------------------|
| L-Amino Acid Assay Buffer | 46 μl |
| L-Amino Acid Probe | 2 μl |
| L-Amino Acid Enzyme Mix | 2 μl |

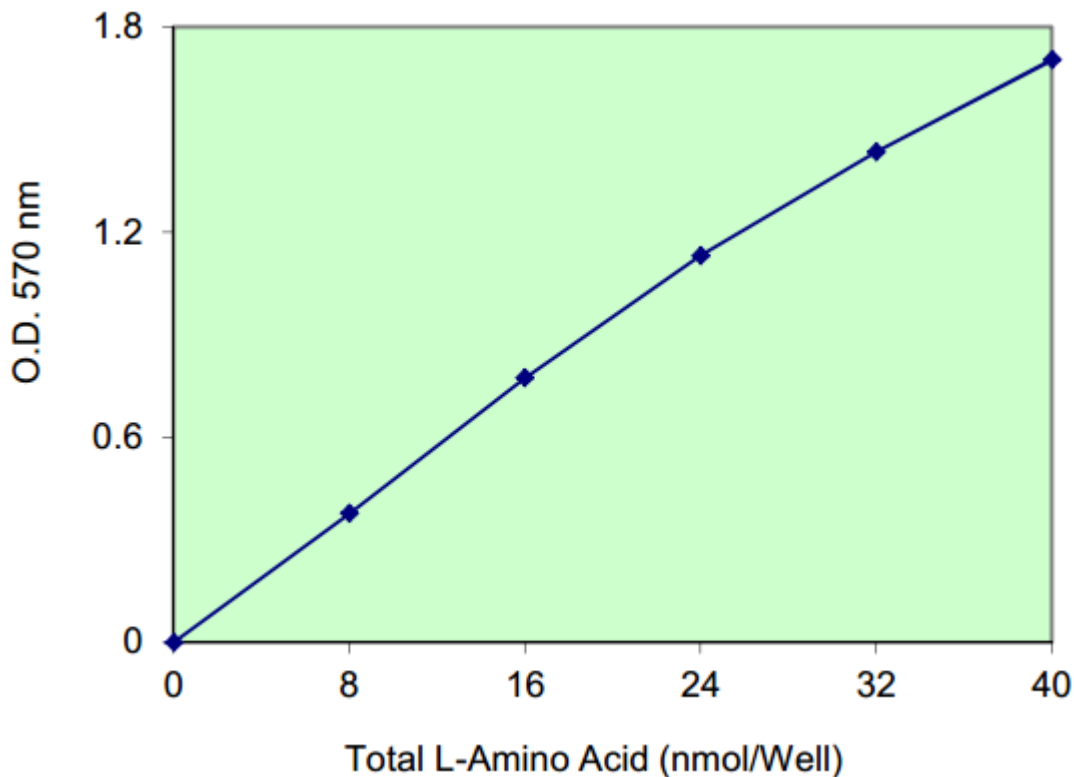
4. Add 50 μl of the Reaction Mix to each well containing the L-Amino Acid standard or test samples.

5. Incubate the reaction for 30 min at 37°C, protect from light.
6. Measure OD at 570nm for the colorimetric assay or fluorescence at Ex/Em = 535/590 nm in a micro-plate reader.
7. Correct background by subtracting the value of the 0 L-amino acid control from all samples (The background reading can be significant and must be subtracted from sample readings). Then apply the sample readings to the L- amino acid standard curve to obtain the total amino acid amount.

$$\text{L-Amino Acid Concentration} = A/S_v \text{ (nmol/}\mu\text{l or mM)}$$

Where A: L-Amino acid amount (nmol) from the standard curve based on Absorbance OD 570 or fluorescence of your samples.

S_v: Sample volume (μl) you added into the sample wells.



L-Amino Acid Standard Curve. Different amounts of L-Amino Acids were measured according to the kit procedure.

General Troubleshooting Guide:

| Problems | Cause | Solution |
|-------------------------------|--|--|
| Assay not working | <ul style="list-style-type: none"> • Use of a different buffer • Omission of a step in the protocol • Plate read at incorrect wavelength • Use of a different 96-well plate | <ul style="list-style-type: none"> • Assay buffer must be at room temperature • Refer and follow the data sheet precisely • Check the wavelength in the data sheet and the filter settings of the instrument • Fluorescence: Black plates ; Luminescence: White plates; Colorimeters: Clear plates |
| Samples with erratic readings | <ul style="list-style-type: none"> • Use of an incompatible sample type • Samples prepared in a different buffer • Samples were not deproteinized (if indicated in datasheet) • Cell/ tissue samples were not completely homogenized | <ul style="list-style-type: none"> • Refer data sheet for details about incompatible samples • Use the assay buffer provided in the kit or refer data sheet for instructions • Use the 10 kDa spin cut-off filter or PCA precipitation as indicated |

| | | |
|--|--|--|
| | <ul style="list-style-type: none"> • Samples used after multiple free-thaw cycles • Presence of interfering substance in the sample • Use of old or inappropriately stored samples | <ul style="list-style-type: none"> • Use Dounce homogenizer (increase the number of strokes); observe for lysis under microscope • Aliquot and freeze samples if needed to use multiple times • Troubleshoot if needed, deproteinize samples • Use fresh samples or store at correct temperatures till use |
| Lower/ Higher readings in Samples and Standards | <ul style="list-style-type: none"> • Improperly thawed components • Use of expired kit or improperly stored reagents • Allowing the reagents to sit for extended times on ice • Incorrect incubation times or temperatures • Incorrect volumes used | <ul style="list-style-type: none"> • Thaw all components completely and mix gently before use • Always check the expiry date and store the components appropriately • Always thaw and prepare fresh reaction mix before use • Refer data sheet & verify correct incubation times and temperatures • Use calibrated pipettes and aliquot correctly |
| Readings do not follow a linear pattern for Standard curve | <ul style="list-style-type: none"> • Use of partially thawed components • Pipetting errors in the standard • Pipetting errors in the reaction mix • Air bubbles formed in well • Standard stock is at an incorrect concentration • Calculation errors • Substituting reagents from older kits/ lots | <ul style="list-style-type: none"> • Thaw and resuspend all components before preparing the reaction mix • Avoid pipetting small volumes • Prepare a master reaction mix whenever possible • Pipette gently against the wall of the tubes • Always refer the dilutions in the data sheet • Recheck calculations after referring the data sheet • Use fresh components from the same kit |
| Unanticipated results | <ul style="list-style-type: none"> • Measured at incorrect wavelength • Samples contain interfering substances • Use of incompatible sample type • Sample readings above/below the linear range | <ul style="list-style-type: none"> • Check the equipment and the filter setting • Troubleshoot if it interferes with the kit • Refer data sheet to check if sample is compatible with the kit or optimization is needed • Concentrate/ Dilute sample so as to be in the linear range |

Note: The most probable list of causes is under each problem section. Causes/ Solutions may overlap with other problems.

For research use only! Not to be used in humans.

Our promise

If the product does not perform as described on this datasheet, we will offer a refund or replacement. For more details, please visit <http://www.apexbt.com/> or contact our technical team.

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