

Product Information

Glucose-6-Phosphate Colorimetric Assay Kit

I. Kit Contents:

Components	K2208-100 100 assays	Cap Color	Part Number
G6P Assay Buffer	25 ml	WM	K2208-C-1
G6P Enzyme Mix	lyophilized	Green	K2208-C-2
G6P Substrate Mix	lyophilized	Red	K2208-C-3
G6P Standard (10 μ mol)	lyophilized	Yellow	K2208-C-4

II. Introduction:

Glucose-6-phosphate (G6P) is an important intermediate for glucose to get into cells. G6P is involved in pentose phosphate pathway, glycolytic pathway or be stored as glycogen or starch. G6P is utilized by glucose 6-phosphate dehydrogenase to produce NADPH, which is very important in red blood cells. The deficiency of G6PDH will lead to hemolytic anemia.

The Glucose-6-Phosphate Colorimetric Assay Kit provides a sensitive, simple, fast and convenient way for detection of G6P levels in various biological samples based on colorimetric method. In the assay, G6P is oxidized to generate a product which is used to transform a nearly colorless probe to an intensely colored product with an absorbance at 450 nm. The kit can detect 1 to 30 nmole G6P with detection sensitivity \sim 10 μ M.

III. Storage and Handling:

Store kit at -20°C , protect from light. Warm G6P Assay Buffer to room temperature before use. Briefly centrifuge all small vials prior to opening. Keep enzyme mix on ice while in use.

VI. Reagent Preparation and Storage Conditions:

G6P Enzyme Mix: Dissolve with 220 μ l dH₂O. Pipette up and down to dissolve. Aliquot into portions and store at -20°C . Avoid repeated freeze/thaw cycles. Use within two months.

G6P Substrate Mix: Dissolve with 220 μ l of G6P Assay Buffer. Pipette up and down to dissolve. Stable for 2 months at 4°C .

G6P Standard: Dissolve in 100 μ l dH₂O to generate 100 mM (100 nmol/ μ l) G6P Standard solution. Keep cold while in use. Store at -20°C .

V. Assay Protocol:

1. Sample Preparation: Liquid or solution samples can be assayed directly. For tissue or cell samples: 10 - 100 mg tissue or 5 million cells should be rapidly homogenized with 2 - 3 volume of ice cold PBS or other buffer (pH 6.5 - 8). Centrifuge at top speed for 10 min to remove insoluble materials. Add 1 - 50 μ l samples into duplicate wells of a 96-well plate and bring volume to 50 μ l with Assay Buffer. For unknown samples, we suggest testing several doses of your samples to ensure readings are within the standard curve range.

Notes:

A. Enzymes in sample may convert or consume G6P. We suggest to deproteinize samples using a perchloric acid/KOH protocol or 10 kDa molecular weight cut off spin filter to remove enzymes. Samples may be homogenized in perchloric acid, then neutralize with 10N KOH to minimize G6P conversion. For tissues or cells containing low level of free G6P (5 - 60 μ M), try to minimize sample dilutions.

B. NADH or NADPH in samples will generate background readings. If NADH or NADPH is in your sample, you may do a background control (omit

G6P Enzyme Mix from the reaction mix) to read the background, then subtracted the background from G6P readings.

2. Standard Curve Preparations:

Dilute the G6P Standard to 1 nmol/μl by adding 10 μl of the 100 nmol/μl Standard to 990 μl of dH₂O, mix well. Add 0, 2, 4, 6, 8, 10 μl into a series of standards wells on a 96 well plate. Adjust volume to 50 μl/well with Assay Buffer to generate 0, 2, 4, 6, 8, 10 nmol/well of G6P Standard.

3. Develop: Mix enough reaction mix for the number of samples and standards to be performed: For each well, prepare a total 50 μl Reaction Mix containing:

	Reaction Mix	Background
G6P Assay Buffer	46 μl	48 μl
G6P Enzyme Mix	2 μl	---
G6P Substrate Mix	2 μl	2 μl

Add 50 μl of the Reaction Mix to each well containing the G6P Standard and samples. Add 50 μl of the background mix into background control wells.

4. Incubate for 30 min at room temperature, protect from light.

5. Measure OD at 450 nm.

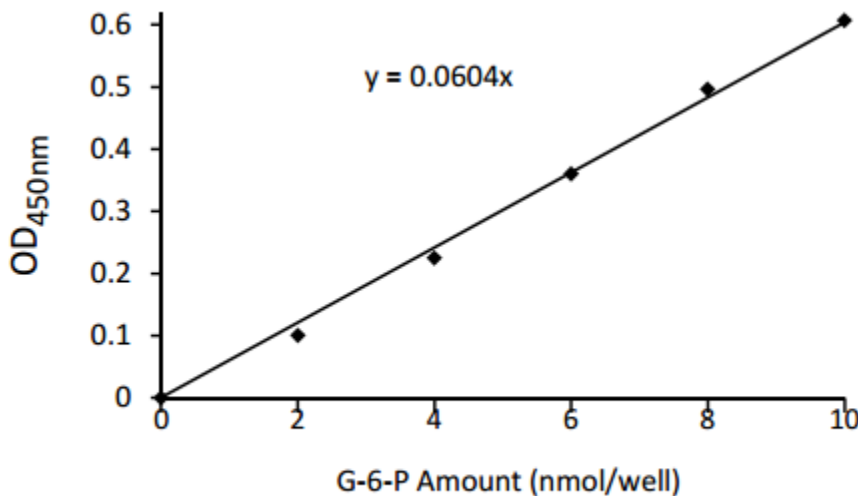
6. Calculation: Correct background by subtracting the value of the 0 G6P blank from all sample readings. If background control reading is significant, subtract the background reading from sample reading. Plot the standard curve. Apply the corrected sample readings to the standard curve to get G6P amount in the sample wells. The G6P concentrations in the test samples:

$$C = Ay/Sv \text{ (nmol/}\mu\text{l; or } \mu\text{mol/ml; or mM)}$$

Where: Ay is the amount of G6P (nmol) in your sample from the standard curve.

Sv is the sample volume (μl) added to the sample well.

Glucose-6-phosphate molecular weight: 260.14.



Glucose-6-phosphate standard curve generated using this kit protocol

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 	<ul style="list-style-type: none"> • Assay buffer must be at room temperature • Refer and follow the data sheet precisely

	<ul style="list-style-type: none"> • Plate read at incorrect wavelength • Use of a different 96-well plate 	<ul style="list-style-type: none"> • 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 • 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"> • 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 • 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
<p>Note: The most probable list of causes is under each problem section. Causes/ Solutions may overlap with other problems.</p>		

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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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