

Starch Colorimetric/Fluorometric Assay Kit (#BN00871)

(Catalog #BN00871; 100 assays; Store at -20°C)

I. Introduction:

Starch is a complex carbohydrate consisting of a large number of glucose units. All plants contain starch, present as amylose, (linear α -1, 4 linked polymer) and amylopectin, (highly α -1, 6 branched α -1, 4 polymer). Starch generally contains 0-25% amylose and 75–100% amylopectin. The Assay Genie Starch Assay Kit provides an easy, convenient method to measure starch levels in a variety of samples. In the assay, starch is hydrolyzed to glucose which is oxidized to generate color (λ_{max} = 570 nm) and fluorescence (Ex/Em = 535/587 nm). The assay can detect starch at 0.0004 to 2 mg/ml.

II. Kit Contents:

Components	BN00871	Cap Code	Part Number
Hydrolysis Buffer	25 ml	NM	BN00871-1
Development Buffer	25 ml	WM	BN00871-2
GenieRed Probe	0.4 ml	Red	BN00871-3A
Hydrolysis Enzyme Mix	Lyophilized	Blue	BN00871-5
Development Enzyme Mix	Lyophilized	Green	BN00871-6
Starch Standard (2.0 mg/ml)	100 μ l	Yellow	BN00871-7

III. Storage and Handling:

Store kit at -20°C, protect from light and moisture. Warm Buffers to room temperature before use. Briefly centrifuge all small vials prior to opening. Read entire protocol before the assay.

IV. Reagent Preparation and Storage Conditions:

GenieRed Probe: Ready to use as supplied. Warm up >18°C to melt frozen DMSO before use. Mix well, store at -20°C, protect from light and moisture.

Hydrolysis Enzyme Mix, Development Enzyme Mix: Dissolve Hydrolysis Enzyme Mix with 220 μ l Hydrolysis Buffer and Development Enzyme Mix with 220 μ l Development Buffer. Vortex gently to dissolve. Keep on ice. Store at -20°C. Stable for at least two months.

Starch Standard: Heat the starch standard to 100 °C for 5 minutes in a heat block or boiling water. After boiling, vortex contents for 5 seconds to dissolve any precipitate.

V. Starch Assay Protocol:

1. Standard Curve Preparations:

Colorimetric: Dilute Starch Standard to 0.2 mg/ml by adding 10 μ l of the Standard to 90 μ l of distilled water, mix well. Add 0, 2, 4, 6, 8, 10 μ l into a series of wells. Adjust volume to 50 μ l/well with Hydrolysis Buffer to generate 0, 0.4, 0.8, 1.2, 1.6 and 2.0 μ g/well of the Starch Standard.

Fluorometric: Dilute Starch Standard to 0.02 mg/ml by adding 10 μ l of the Standard to 990 μ l of distilled water, mix well. Add 0, 2, 4, 6, 8, 10 μ l into a series of wells. Adjust volume to 50 μ l/well with Hydrolysis Buffer to generate 0, 0.04, 0.08, 0.12, 0.16 and 0.2 μ g/well of Starch Standard.

2. Sample Preparation:

Depending on your assay purpose (quantitation, mw distribution, compartmentalization, etc.), prepare starch samples according to established protocols¹⁻⁴.

A. Soluble Starch Extraction: Grind up 5-10 mg sample, wash off any free glucose and small oligosaccharides with 1 ml 90% ethanol, warm to 60°C for 5 minutes with occasional vortexing. Centrifuge at 10,000g for 2 minutes. Decant the supernatant. Repeat the wash twice. Soluble starch can be extracted with 1 ml H₂O and heating on a boiling water bath for 5 minutes. Spin at 10,000g for 2 minutes to remove insoluble materials. The supernatant is soluble starch.

B. Resistant Starch Extraction: After extracting soluble starch, extract the water insoluble pellet with 1 ml 10N KOH, heat on boiling water bath for 5 minutes. Neutralize with 1 ml 10M H₃PO₄ slowly. Spin at 10,000g for 2 minutes to remove insoluble materials. The supernatant is resistant starch.

C. Total Starch Extraction: After the 90% ethanol wash (Step A), extract the washed sample directly with 10N KOH/H₃PO₄ as per the procedure for resistant starch (B). The supernatant is

total starch. **For starch sample testing:** Take 20 μ l of the extracted starch, add 180 μ l of Hydrolysis Buffer, mix. Add up to 50 μ l of the diluted sample or buffer (blank) to test wells.

Adjust the volume to 50 μ l with Hydrolysis Buffer. For unknown samples, we suggest testing several doses of the sample to ensure the readings are within the standard curve.

3. Hydrolysis*:

	Colorimetric	Fluorometric
Hydrolysis Enzyme Mix	2 μ l	1 μ l
Mix well; incubate for at least 30 minutes at room temperature to hydrolyze starch.		

***Note:** Glucose generates background. Glucose control is done without the hydrolysis enzyme (add equal volume of Buffer). Glucose background can be subtracted from sample reading.

4. Development:

Mix enough reagents for the number of samples and standards. For each well, prepare a total 50 μ l Reaction Mix.

	Colorimetric	Fluorometric
Development Buffer	46 μ l	48.7 μ l
Development Enzyme Mix	2 μ l	1.0 μ l
GenieRed Probe	2 μ l	0.3 μ l

Add 50 μ l of Development Mix to each well containing Starch Standard or samples.

5. Incubate at room temperature for 30 minutes, protect from light.

6. Measure colorimetrically (OD=570 nm) or fluorometrically (Ex/Em 535/587 nm).

7. **Calculation:** Correct background by subtracting the value of the 0 starch control from all sample readings (Note: The background can be significant and must be subtracted). Plot standard curve μ g/well vs. OD. Apply sample readings to the standard curve to get the amount of starch in the sample wells. The starch concentration in the test samples:

$$C = Ay/Sv \text{ (}\mu\text{g}/\mu\text{l or mg/ml)}$$

Where: Ay is the amount of starch (μ g) in your sample from the standard curve.
Sv is the sample volume (μ l) added to the sample well. Multiply by the dilution factors.
Starch molecular size: ~ 60,000 glucose molecules (MW ~10⁶-10⁷ daltons).

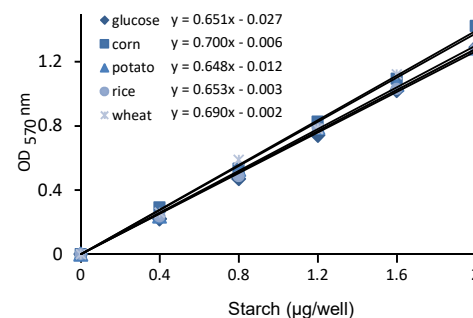


Figure 1. Starch Standard Curve: Different types of pure starch were extracted with 10N KOH/H₃PO₄ as described following the kit protocol.

FOR RESEARCH USE ONLY! Not to be used on humans.

GENERAL TROUBLESHOOTING GUIDE:

Problems	Cause	Solution
Assay not working	<ul style="list-style-type: none"> • Use of ice-cold 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"> • 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 (clear bottoms) ; 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 • 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 datasheet & 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.		