



- b. For uncharacterized enzymes, we suggest testing several doses to ensure the reading is within the Standard Curve range.
2. **SAH Standard Curve:** Prepare 200 μM SAH Standard stock in MT Assay Buffer by diluting 4 μl of 50 mM SAH Standard in 996 μl MT Assay Buffer. Add 0 (Background Control), 5, 15, 25, 35, 45 μl of 200 μM SAH standard into a series of wells on a 96-well plate to generate 0, 1, 3, 5, 7, 9 nmol/well of SAH Standard. Adjust the volume to 50 μl with MT Assay Buffer.
3. **Reaction Mix:** Mix enough reagents for the number of samples and standards to be performed:

Sample Reaction Mix (1 assay)	
MT Assay Buffer	37 μl
Enzyme Mix I	2 μl
Enzyme Mix II	2 μl
Enzyme Mix III	6 μl
SAM Cofactor (50 mM)	1 μl
OxiRed™ Probe	2 μl

Mix and add 50 μl of the Sample Reaction Mix to each well containing the Positive Control, Test Samples, Standards and Background Control.

4. **Measurement:** For positive control, test samples, and background control measure absorbance at 570 nm in kinetic mode every 30 seconds for at least 45 minutes at 37 °C. To generate the SAH Standard Curve, incubate SAH standard reactions for 45 minutes at 37 °C and measure absorbance at 570 nm in end-point mode or simply take absorbance reading at the 45 minute mark from the kinetic reading.

Note: Your sample methyltransferase may have a different optimal temperature. You may change reaction temperature to suit your needs. Similarly, your sample methyltransferase may have a different K_M for the SAM Cofactor. In this case, the user may decide on the optimal SAM Cofactor concentration to incorporate into the master mix. For Methyltransferase enzymes showing very low activities, it may be advantageous to use a more sensitive Methyltransferase Activity Assay Kit (Fluorometric) (BV Cat No. K521-100).

5. **Calculations:** *Standard Curve:* Subtract 0 nmol SAH Standard reading from all SAH standards to obtain normalized standard curve. Plot the SAH standard curve. Apply a linear fit to the SAH standard values and determine the standard curve equation. *Samples/Positive Control:* subtract each point on the "no methyltransferase" background control curve from each corresponding point generated in Sample and Positive Control readings. Apply OD values at each time point to the SAH standard curve equation to determine nmol of SAH generated at each time point; multiply these values by 1000 to determine pmol of SAH generated at each time point. Plot pmol SAH on the y-axis vs. time (in minutes) on the x-axis and determine the sample reaction slope (pmol/min) of the linear portion of the curve.

$$\text{Sample Methyltransferase Activity} = [\text{sample reaction slope}/V] \times D \text{ (pmol/min/ml} \equiv \mu\text{U/ml)}$$

$$\text{Sample Specific Activity} = \text{slope}/\mu\text{g (pmol/min}/\mu\text{g} \equiv \mu\text{U}/\mu\text{g})$$

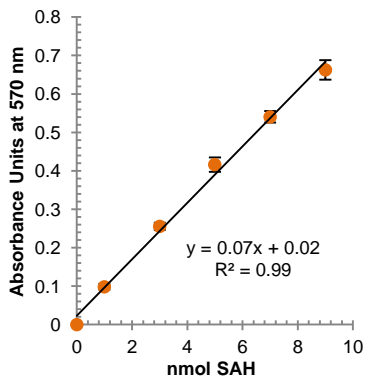
Where: V = sample volume added into the reaction well (ml).

D = Dilution Factor

$$\text{Sample Reaction Slope} = \text{pmol/min (calculated using the SAH standard curve equation)}$$

Unit Definition: One unit of methyltransferase is the amount of enzyme that generates 1.0 μmol of SAH per min. at 37°C.

a)



b)

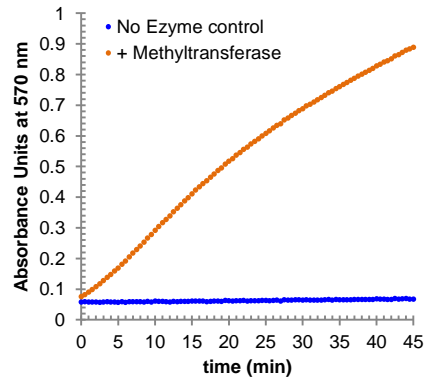


Figure: a) Normalized SAH Standard Curve; **b)** Representative activity curve for human recombinant Nicotinamide N-Methyltransferase (NNMT) (BV Cat No.7261) with Nicotinamide Substrate at 37°C. Assays were performed using kit protocol.

VIII. RELATED PRODUCTS:

Adenosylhomocysteinase (AHCY) Activity Fluorometric Assay Kit (K807)
Methyltransferase Activity Assay Kit (Fluorometric) (K521)
SGI-1027 (2726)
Zebularine (2225)

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