Video Article

Measurement of Heme Synthesis Levels in Mammalian Cells

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Abstract

Heme serves as the prosthetic group for a wide variety of proteins known as hemoproteins, such as hemoglobin, myoglobin and cytochromes. It is involved in various molecular and cellular processes such as gene transcription, translation, cell differentiation and cell proliferation. The biosynthesis levels of heme vary across different tissues and cell types and is altered in diseased conditions such as anemia, neuropathy and cancer. This technique uses [4-¹⁴C] 5-aminolevulinic acid ([¹⁴C] 5-ALA), one of the early precursors in the heme biosynthesis pathway to measure the levels of heme synthesis in mammalian cells. This assay involves incubation of cells with [¹⁴C] 5-ALA followed by extraction of heme and measurement of the radioactivity incorporated into heme. This procedure is accurate and quick. This method measures the relative levels of heme biosynthesis rather than the total heme content. To demonstrate the use of this technique the levels of heme biosynthesis were measured in several mammalian cell lines.

Video Link

The video component of this article can be found at <http://www.jove.com/video/51579/>

Introduction

Heme, a complex of ferrous iron and protoporphyrin IX is a central molecule for transporting and utilizing oxygen in virtually all living organisms¹⁻³. The unique structure of heme enables it to function as a carrier of diatomic gases and electrons, as well as to perform various other functions¹⁻⁵. For example, heme binds to oxygen in hemoglobin and myoglobin for the transfer and storage of oxygen^{6,7}. It also works as an electron carrier in the cytochromes during respiration and acts as an electron donor for redox reactions catalyzed by cytochrome P450 enzymes^{8,9}. One of the most significant features of heme is that, it can play regulatory roles in cellular and molecular processes such as gene transcription, protein synthesis and micro-RNA biogenesis⁴. For example, it affects the transcription of many genes by controlling the activity of mammalian transcriptional repressor Bach1 and the mammalian nuclear receptor Rev-erba¹⁰⁻¹⁵. Heme regulates the activation of heme activator protein (Hap) 1 which plays an important role in the activation of genes involved in respiration and controlling oxidative damage, in response to heme or oxygen¹⁶. Heme also regulates gene transcription in neuronal cells via nerve growth factor (NGF) signaling^{3,17-20}. It also regulates protein synthesis in mammalian erythroid cells by modulating the activity of heme-regulated eIF2 α kinase (HRI)²¹⁻²⁴. Furthermore, heme affects the activity of key signaling proteins such as tyrosine kinase Jak2 and Src, which are essential for proper cell functioning and cell growth^{4,20,25}. It was found that in HeLa cells heme inhibition causes cell cycle arrest and activation of markers associated with senescence and apoptosis²⁶. Both Heme deficiency or increased levels of heme are associated with severe health effects in humans²⁷. Recent molecular and epidemiological studies have shown a positive association of high heme intake and increased risk of diseases, such as type-2 diabetes, coronary heart disease and several cancers including lung cancer, colorectal cancer and pancreatic cancer^{27,28}. Using a matched pair of normal and cancer lung cells authors' lab have found that cancer cells have increased levels of oxygen consumption, heme synthesis and proteins involved in heme uptake and oxygen utilization²⁸. Interestingly, inhibition of heme synthesis decreased oxygen consumption, proliferation, migration and colony formation of cancer cells²⁸. Thus, the fluctuation in the levels of endogenous heme plays an important role in the regulation of molecular and cellular processes^{3,4,28,29}.

In mammals, the biosynthesis of heme occurs in eight steps, involving enzymes located in the mitochondria and the cytosol⁴ (**Figure 1**). Heme biosynthesis begins in the matrix of mitochondria with the condensation of glycine and succinyl-coA to form 5-aminolevulinic acid (5-ALA), catalyzed by ALA synthase (ALAS)^{4,31}. This is the rate limiting step in heme biosynthesis in nonerythroid cells. 5-ALA is then exported out to the cytosol where the next four steps occur to form coproporphyrinogen III (CPgenIII), which is then imported back to the mitochondria, where it is converted into protoporphyrin IX (PPIX). Finally, one molecule of iron is incorporated to the protoporphyrin IX (PPIX) to produce heme, a reaction catalyzed by ferrochelatase (FECH)^{2,4}.

The level of heme biosynthesis depends primarily on the level of ALAS enzyme which is tightly controlled by intracellular iron and heme⁴. The biosynthesis of heme can be affected by genetic defects, availability of certain minerals and vitamins (e.g., riboflavin, zinc), exposure to toxins (e.g., aluminum, lead), anoxia, fever, and levels of certain steroids (e.g., estrogen)³²⁻³⁵. The level of heme synthesis is altered in various diseased conditions. Decreased heme biosynthesis can cause anemia as well as neurological diseases^{3,36}. Alternatively, increased heme biosynthesis plays an important role in the progression of certain cancers^{28,37}. Heme has been shown to be critical for the growth, differentiation and survival of mammalian adipose, erythroid and neuronal cells^{4,38-41}. For example, heme deficiency leads to neurite damage in primary mouse cortical

neurons via the inhibition of glutamate NMDA (*N*-methyl-D-aspartate) receptor¹⁷. Additionally, inhibition of heme synthesis causes programmed cell death in the human epithelial cervix carcinoma HeLa cells^{26,41}. Therefore, measuring the heme biosynthesis levels in various cells under different conditions is important for studying etiology and progression of many diseases.

Here we describe a fast and sensitive method to measure the level of intracellular heme synthesis by using [4-¹⁴C] 5-aminolevulinic acid. This is an alternative method to other methods using ⁵⁵Fe or ⁵⁹Fe. We prefer using ¹⁴C because its radiation is very weak. In contrast, strong protection is required for working with Fe isotopes. Furthermore, this method is intended to measure and compare heme synthesis in different cells in parallel in a quick manner. In order to measure absolute heme levels, one may use the previously established method involving the use of HPLC^{42,43}.

Protocol

CAUTION: While working with radioactivity, take appropriate precautions to avoid contamination of the experimenter and the surroundings. Dispose off all waste following local radiation safety guidelines.

1. Preparation of Cells

- Seed cells in 3.5 cm plates such that the confluency reaches 80%-90% on the day of assay.
 - Note that seeding confluency for cells depends upon the cell type and their growth rate. When treating cells with a reagent for a certain number of days, seed cells so that the confluency is 80%-90% on the day of heme measurement. If the number of cells obtained from 3.5 cm plates is insufficient, use 6 cm plates with same amount of radioactive ALA as in the next step.
Note: Use of individual plates is recommended because it is easier to handle individual plates.
- A day before heme measurement, add 0.1-0.3 µCi [¹⁴C] 5-ALA and cold ALA to reach a final concentration of 20 µM of total ALA to each culture dish in a work area fit for working with radioactive materials and put the plates back into the incubator. Ideally, incubate with radioactive ALA for 16 hr.
Note: Radiolabeled 5-ALA required may vary between cell lines. A radioactive count per minute (cpm) of at least 100 is required to minimize the error associated with counting. It is advisable to determine a minimum amount of [¹⁴C] 5-ALA required for the cell lines under study. For most experiments 0.1-0.3 µCi is sufficient to get a good measurement. Addition of cold ALA is optional.

2. Heme Extraction

- Place the cell culture dishes on ice and take them to the area fit for radioactivity work.
 - Perform the whole extraction and evaporation process in an explosion-proof hood. Because the stored ether can explode, use the ether in small bottles and evaporate any remaining liquids, do not store ether in an open bottle for more than one month. Do not use acetone near open flames and heaters as it is highly flammable. Store acetone containing reagents in glass bottles as it dissolves tissue culture plastics.
- Aspirate off the medium using a pipette. Wash cells one time with 1 ml of 1x cold PBS. Aspirate off PBS.
- Add 1 ml of 1x cold PBS to the plates and collect the cells at the bottom of the plate using a rubber policeman. Transfer the collected cells from each plate to a labeled pre-chilled 2 ml tube. Collect all cells in 2 ml tubes.
Note: Use a fresh rubber policeman for each plate for scraping cells from each plate. For each plate you would require four 2 ml tubes, a 1.5 ml tube and a scintillation vial.
- Spin at 15,000 x g for 1 min at 4 °C.
- Take out PBS using a pipette, give a short spin and remove the residual PBS. Resuspend cells in 100 µl of 1x PBS. Vortex to resuspend cells and keep on ice.
- Take 10 µl from step 5 and transfer to a 1.5 ml microcentrifuge tube and store on ice to measure protein concentration.
Note: Cell lysis buffer should be added to the cells and kept on ice to measure protein concentration later.
- Spin the remaining 90 µl from step 5 at 5,000 x g for 30 sec at 4 °C; remove PBS completely using a 200 µl pipette tip.
- Add 300 µl of heme extraction buffer (see Materials List) to each tube. Then vortex for 15 sec to mix and put on ice for 1 min, repeat this 3 times.
 - Add heme extraction buffer (HEB) and vortex the tube quickly to resuspend the pellet, do only one or two tubes at a time because the pellet should be resuspended quickly without being left in the HEB for too long. Once HEB is added to all the tubes and pellet is resuspended then follow the 15 sec vortex and 1 min on ice cycle as in step 8. Note that pellet may not go completely into the HEB if the pellet is left in the HEB buffer for a longer time and not resuspended quickly.
- Add 1.2 ml of diethyl ether to each tube and vortex for 30 sec. Spin at 15,000 x g for 5 min at 4 °C.
Note: Diethyl ether runs out of pipette tips quickly. To control this, pipette diethyl ether up and down once or twice, doing so helps hold the diethyl ether in the pipette. Repeat this every time a fresh pipette tip is used.
- Transfer top (ether) phase with 1 ml pipette to a new microcentrifuge tube. Add 0.5 ml of 2 N HCl, vortex to mix and spin as in step 9.
Note: It is difficult to see the separation between ether and HCl phases. Therefore, add small amounts of Trypan blue in 2 N HCl solution to be used during the experiment, enough to give it a pale blue color. This will help distinguish the two phases. The lower phase will be blue and the upper ether phase will be colorless. Transfer the upper colorless ether phase to a new tube and discard the lower HCl (blue) phase.
- Repeat step 10 two more times to wash the ether phase with 2 N HCl. Transfer only the top ether phase to a new tube during each repetition and discard the lower HCl phase.
- After the third wash with HCl, transfer the top phase with 1 ml pipette to a scintillation vial. Dry the samples by evaporating the diethyl ether by keeping the samples in the chemical hood O/N.

3. Measurement of Radioactivity

1. Next day, add 5 ml of scintillation fluid to the dried samples from step 2.12.
2. Shake well to mix. Measure the radioactivity by using a scintillation counter.

4. Calculations

1. Lyse the cells from step 2.6 using equal amounts of CellLytic M buffer (see Materials List) and spin at 14,000 x g for 5 min at 4 °C.
2. Take an aliquot of the supernatant and measure the protein concentration using BCA protein assay kit.
Total amount of protein in mg (TP) = (90 µl * conc. of protein in µg/µl) / 1,000
Heme synthesis level (radioactivity/TP) = pmol/mg of protein.

Representative Results

This method was used to compare the levels of heme synthesis in normal (HBEC30KT) vs. cancer (HCC4017) lung cells. **Figure 2** shows a higher level of heme synthesis in cancer cells (HCC4017) than normal lung cells (HBEC30KT). The level of heme synthesis was also measured in normal and cancer cells in the presence of mitochondrial uncoupler carbonyl cyanide 3-chlorophenylhydrazone (CCCP). Cells were treated with 10 µM CCCP for 24 hr before the measurement of heme synthesis levels. As expected, the levels of heme synthesis (**Figure 2**) decreased in the presence of CCCP in both normal and cancer cells. It has been shown previously that heme synthesis can be inhibited by succinyl acetone (SA), a potent and specific inhibitor of 5-aminolevulinic acid dehydratase (ALAD) which is the second enzyme in heme biosynthetic pathway⁴¹. Using this method, the heme synthesis levels were measured in HeLa cells in the presence and absence of 0.5 mM SA. **Figure 3** shows that the levels of heme synthesis decreased by more than 10-folds in the presence of SA. To further demonstrate the use of this technique, the levels of heme synthesis were measured and compared in 4 cancer and in HEK293T cell lines as shown in **Figure 4**.

Figure 1. The heme biosynthesis pathway in humans. ALAS, 5-aminolevulinic acid synthase; 5-ALA, 5-aminolevulinic acid; ALAD, ALA dehydratase; PBG, porphobilinogen; HMBS, hydroxymethylbilane synthase; HMB, hydroxymethylbilane; UROS, uroporphyrinogen III synthase; UPgenIII, uroporphyrinogen III; UROD, uroporphyrinogen decarboxylase; CPgenIII, coproporphyrinogen III; CPOX, coproporphyrinogen III oxidase; PPgenIX, protoporphyrinogen IX; PPOX, protoporphyrinogen IX oxidase; PPIX, protoporphyrin IX; FECH, ferrochelatase. [Please click here to view a larger version of this figure.](#)

Figure 2. The levels of heme biosynthesis in normal (HBEC30KT) and cancer (HCC4017) cell lines, in absence and presence of 10 μ M mitochondrial uncoupler CCCP. The background level was similar to the level of blank sample (scintillation fluid alone) and it was less than 2% of the control. [Please click here to view a larger version of this figure.](#)

Figure 3. Levels of heme biosynthesis in HeLa cells in absence and presence of 0.5 mM of heme synthesis inhibitor succinyl acetone (SA). Cells were treated with SA for 3 days before heme measurement. The background level was similar to the level of blank sample (scintillation fluid alone) and it was less than 2% of the control. [Please click here to view a larger version of this figure.](#)

Figure 4. Comparison of the levels of heme biosynthesis in 4 cancer and HEK293T cell lines. Assay was performed as per the protocol. HCC4017 were grown in ACL4 media, A549 and H1395 in RPMI1640 supplemented with 5% FBS, HEK293T and HeLa in DMEM supplemented with 10% FBS. For statistical analysis, the heme synthesis levels in cancer cells and HEK293T cells were compared to the heme synthesis level in HCC4017 cells, by using Welch 2-sample t-test. **, p value < 0.005. The background level was similar to the level of blank sample (scintillation fluid alone) and it was less than 2% of the control. [Please click here to view a larger version of this figure.](#)

Discussion

Heme plays a key role in the generation of cellular energy via respiration²⁶. Altered heme metabolism is known to be associated with various diseases including cancer^{28,41}. Inhibition of heme synthesis is known to cause cell cycle arrest and apoptosis in Hela cells^{26,41}. It has been shown that high heme synthesis level is associated with the progression of lung cancer cells²⁸. Therefore, it would be of great importance to measure the levels of heme biosynthesis in cells under different conditions. The method discussed here does not quantify the total amount of heme therefore, it does not provide the absolute value of heme present in the cells. There are a variety of colorimetric methods and kits available to measure the total amounts of heme in the cells. For example, heme can be extracted from cells and mitochondria in a mixture of acetone and HCl and mixed with 50% (v/v) acetonitrile. Then the mixture can be applied to a 3.9 x 300 mm C18 Bondclone column followed by elution of heme in a gradient of acetonitrile containing 0.05% (v/v) trifluoroacetic acid and identifying heme compounds at 400 nm^{44,45}.

Previously, to measure the levels of heme biosynthesis in mammalian tissues and cell extracts [¹⁴C]-glycine and [¹⁴C]-5-ALA were used^{46,47}. Although labeled glycine enters the heme biosynthesis pathway at the first step but it is drained off by other metabolic pathways resulting in its limited incorporation into heme. Glycine is not a specific metabolite for heme biosynthesis pathway⁴⁸. On the other hand, labeled 5-ALA is specific to heme biosynthesis pathway. This feature confers a quantitative advantage of using 5-ALA over glycine^{46,48,49}. Therefore, use of radioactive 5-ALA is a more specific and accurate method to measure and compare the levels of heme biosynthesis.

The most critical steps to get consistent results are: 1) make sure the cell confluency does not go higher than 100% and not below 70% across various conditions; 2) the amount of radioactivity added per plate should be identical because this method is very sensitive; 3) heme extraction buffer should not be old, it is recommended to make fresh every time. The volume of radioactivity required per plate is very small and it is difficult to maintain the identical amounts across the samples. Therefore, it is recommended to make a working stock solution in the culture medium, if using the same medium for all the samples, else PBS (phosphate buffered saline) can be used. Working stock solution can be prepared in such a way that the volume dispensed per plate is under 10 µl. Also, to be more accurate, the medium from the replicate samples for one condition can be taken out in a 50 ml tube and the appropriate amounts of radioactivity can be added from the working stock solution, then dispense back equal amounts of medium to the plates. If the number of cells are limited, such as in primary cultures, 12-well or 24-well plates can be used to culture the cells and reduce the amount of radioactivity added per well proportionately.

This protocol is straightforward, simple, and provides accurate measures of the levels of heme synthesis. Sometimes, it may be difficult to resuspend the cell pellet in the heme extraction buffer (step 2.8) completely. To avoid this be quick in step 2.8, add heme extraction buffer to 1-2 samples at a time, vortex and put them on ice. Make sure the heme extraction buffer is not too old. If the standard deviation among the replicate samples is too high, pay attention to the following steps: 1) PBS was removed completely in step 2.7; 2) avoid spilling the sample when in diethyl ether phase, follow the tip in step 2.9; 3) make sure the radioactivity added per plate and the confluency of the replicate samples were the same.

Using radioactive 5-ALA method, the levels of heme synthesis were compared as shown previously²⁸ in normal nonmalignant (HBEC) vs. cancer (HCC) cells, developed from the same patient. The technique described here requires the use of [¹⁴C] 5-ALA, a precursor in heme biosynthesis pathway (**Figure 1**), and measures the radioactivity incorporated into heme. In this technique, porphyrins, including heme, were extracted from the cells in a mixture of acetone, concentrated HCl and water^{50,51}. Heme was further separated from porphyrins by extracting in diethyl ether and washing the ether phase with 2 N HCl^{50,51}. Then, the radioactivity incorporated into heme was determined using a scintillation counter. Results

can be normalized by the total amount of protein or cell number. Heme synthesis in HeLa cells was inhibited by using succinyl acetone (SA) and a decrease in the level of heme synthesis was observed as shown previously²⁶ (Figure 3).

This method is very useful for the comparison of the levels of heme synthesis under different conditions or between cell lines (Figure 4). It requires a small number of cells and provides specific measurement and comparison of the levels of heme synthesis in different cells. It is not intended for the accurate measurement of absolute heme levels in the cells. It is best used to compare the levels of heme synthesis in cells with different properties, such as different cancer cells. Because cobalt, instead of iron, can be incorporated into heme in the last step of heme synthesis and be extracted⁵¹, this method may not yield accurate comparison of heme synthesis, if the growth media contain different levels of cobalt or other metal ions. Nonetheless, the heme synthetic pathway from 5-aminolevulinic acid to porphyrins and heme is very specific, measuring the flux levels of the whole pathway is likely to be more reflective of metabolic activities relevant to heme, regardless of the metal ions. In contrast, the method using ⁵⁹Fe detects only the part of the pathway incorporating Fe to form heme⁵². Although this is more specific, in the presence of significant levels of metal ions, the detected heme synthesis levels may not reflect the metabolic potential of certain cells such as cancer cells. Furthermore, Fe can be sequestered into other cellular locations beside mitochondria to make heme. By and large, we believe that the use of [4-¹⁴C] 5-ALA is a more practical method for measuring heme levels when comparing different cells with varying properties.

Disclosures

We have nothing to disclose.

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