



- a) For Sample(s) with high background such as liver tissue lysate, dilute the lysate with HAO assay buffer 5-10 times and filter through 10 kDa Spin Columns (BV# 1997). Small molecules will be removed in the ultrafiltrate and the ultraconcentrate is used for the HAO activity assay.
- b) We recommend using the Samples for activity analysis immediately. Otherwise, store the Sample(s) at -80°C for 3-4 days.
- c) For Unknown Samples, we suggest testing several concentrations to ensure that the readings are within the Standard Curve range.
- d) Do not dilute Positive Control in HAO Assay Buffer. Do not dilute the entire vial at one time. Dilute enough for the number of Positive Control being run at a time.
- 2. H<sub>2</sub>O<sub>2</sub> Standard Curve Generation:** Dilute 10 µl 0.88M H<sub>2</sub>O<sub>2</sub> Standard into 870 µl dH<sub>2</sub>O to generate 10 mM H<sub>2</sub>O<sub>2</sub> Standard. Dilute 10 µl of the 10 mM H<sub>2</sub>O<sub>2</sub> Standard into 990 µl dH<sub>2</sub>O to generate a 0.1 mM H<sub>2</sub>O<sub>2</sub> Standard. Further dilute 100 µl of the 0.1 mM H<sub>2</sub>O<sub>2</sub> Standard into 900 µl dH<sub>2</sub>O to generate 10 µM H<sub>2</sub>O<sub>2</sub> Standard.

Add 0, 2, 4, 6, 8, 10 µl of the 10 µM H<sub>2</sub>O<sub>2</sub> Standard into a 96-well white plate in duplicates to generate 0, 20, 40, 60, 80, 100 pmol/well H<sub>2</sub>O<sub>2</sub> Standard. Adjust the volume of each well to 50 µl with HAO Assay Buffer.

- 3. Reaction Mix:** Mix enough reagents for the number of assays to be performed. Make sufficient amount of each type of the mix to add 50 µl to all assay wells of that type.

	H <sub>2</sub> O <sub>2</sub> Standard Curve/SBC Mix	Reaction Mix
HAO Assay Buffer	46 µl	44 µl
HAO Substrate	-	2 µl
HAO Enzyme	2 µl	2 µl
HAO Probe	2 µl	2 µl

Mix well. Add 50 µl of the H<sub>2</sub>O<sub>2</sub> Standard Curve/SBC Mix to "H<sub>2</sub>O<sub>2</sub> Standard Curve" and "SBC" wells and 50 µl of the Reaction Mix to SC, Sample and Positive Control wells.

**Notes:**

- a). Have the plate reader ready at Ex/Em 535/587 nm in a kinetic mode at RT set to record fluorescence every 30 sec.
- b). Prepare reaction mix immediately before adding to wells.
- 4. Measurement:** Immediately start recording fluorescence in kinetic mode at 30 sec intervals for 35 -400 min at RT. Standard Curve may be read in either kinetic or end point mode.
- 5. Calculation:** Subtract the 0 Standard readings from all Standard readings and SBC readings from Sample readings respectively. If the Substrate Control (SC) reading is higher than the SBC reading, subtract the Substrate Control readings from the Sample readings instead. Plot the H<sub>2</sub>O<sub>2</sub> Standard Curve. Choose any two time points within the linear portion of the curve (t<sub>1</sub> & t<sub>2</sub>) for each Sample. Apply the corrected Sample readings to the H<sub>2</sub>O<sub>2</sub> Standard Curve to get  $\Delta M$  pmol of H<sub>2</sub>O<sub>2</sub> formed during the reaction time ( $\Delta t = t_2 - t_1$ ).

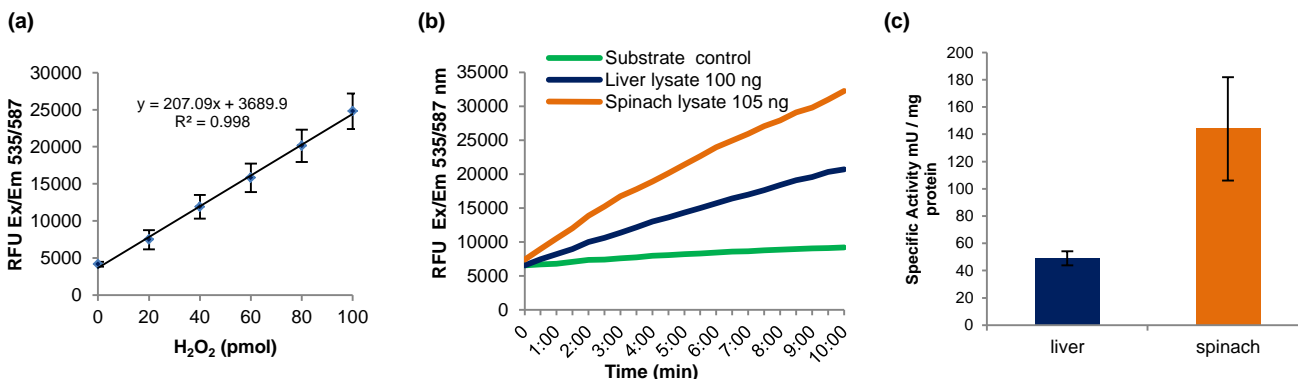
$$\text{Sample HAO Specific Activity} = \Delta M / (\Delta t \times P) \text{ (pmol / (min} \times \mu\text{g))} = \mu\text{Units / } \mu\text{g or mUnits / mg}$$

Where:  $\Delta M$  = linear change in H<sub>2</sub>O<sub>2</sub> concentration during  $\Delta t$  (pmol)

$\Delta t = t_2 - t_1$  (min)

P = Sample protein amount added to well (µg)

**Unit Definition:** One unit of HAO is the amount of enzyme that produces 1 µmol of H<sub>2</sub>O<sub>2</sub> per minute at pH 7.5 at RT.



**Figures. a).** H<sub>2</sub>O<sub>2</sub> Standard Curve. **b).** Enzyme kinetics of Rat Liver lysate (100 ng protein) and Spinach Leaf lysate (105 ng protein). **c).** HAO specific activity in Rat liver and Spinach lysates. Experiments were conducted according to kit protocol.

**VIII. Related Products:**

- Oxalate Oxidase Activity Assay Kit (Fluorometric) (K509)
- Oxalate (Oxalic Acid) Colorimetric Assay Kit (K663)
- Oxalate Decarboxylase Activity Colorimetric Assay Kit (K664)

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