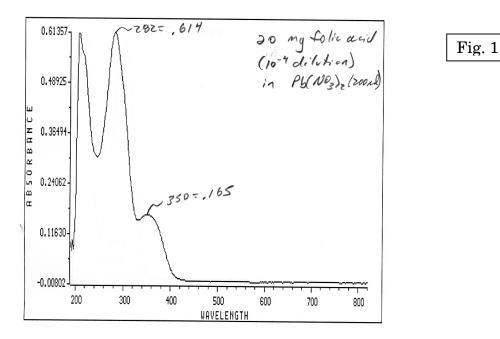
## Synthesis of (6-R,S)5-CH<sub>3</sub>-H<sub>4</sub>PteGlu<sub>n</sub> from PteGlu<sub>n</sub>

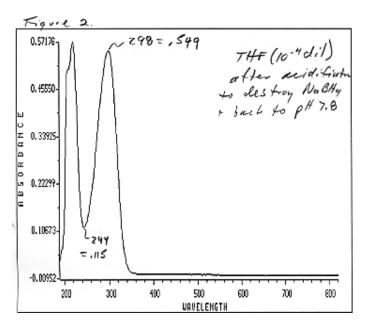
- **References**: Yeo, E. J., and Wagner, C. (1992) Purification and properties of pancreatic glycine N-methyltransferase. *J Biol Chem* 267, 24669-24674; Nando Todari, A., and Marazza, N. F. (2000) Process for the reduction of pterins. (US Patent 6162914) Cerbios-Pharma S.A., Switzerland
- **Outline:** This method uses NaBH<sub>4</sub> to reduce folic acid (PteGlu<sub>n</sub>) to H<sub>4</sub>PteGlu<sub>n</sub> with the modification of the water-soluble  $Pb(NO_3)_2$  to catalyze the reduction. The inclusion of  $Pb(NO_3)_2$  allows less NaBH<sub>4</sub> to be used. The H<sub>4</sub>PteGlu<sub>n</sub> is then condensed with HCHO to produce 5,10-methylene-THF (CH<sub>2</sub>-H<sub>4</sub>PteGlu<sub>n</sub>), which is then reduced with NaBH<sub>4</sub> to 5-CH<sub>3</sub>-H<sub>4</sub>PteGlu<sub>n</sub>.

## Method:

- 1. Weigh 40  $\mu$ mol PteGlu<sub>n</sub> (20 40 mg depending on # of glutamates) into a 2 ml graduated free-standing screwcap tube with round bottom. Add 200  $\mu$ l Pb(NO<sub>3</sub>)<sub>2</sub> (0.265 mM) prepared as follows:
  - a. dissolve 20-30 mg Pb(NO<sub>3</sub>)<sub>2</sub> (Mallinkrodt) in ddH<sub>2</sub>O to 8.8 mg/ml = 26.5 mM (use 1-2  $\mu$ l conc HCl to dissolve completely)
  - b. dilute 1:100 with 10 mM Tris-Cl, pH 7.5 to make 0.265 mM working stock
- Adjust pH to 7.5 with 5 N NaOH (15-35  $\mu$ l); spot small amounts on pH paper to check) and vortex to dissolve PteGlu<sub>n</sub> (should get clear, dark yellow solution)(higher polyglu require considerable vortexing and more NaOH to dissolve completely). Check spectrum of 10<sup>-4</sup> dilution in 10 mM Tris-Cl, pH 7.5 (see figure 1).

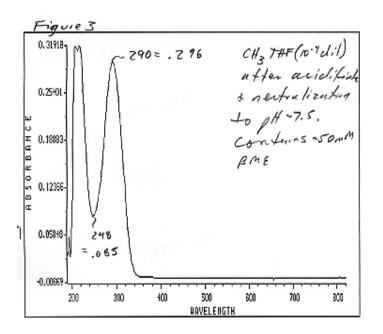


- 2. Add stir bar and move to cold room. Use tube rack to hold tube while stirring at top speed. Add 5 mg NaBH<sub>4</sub> (Sigma) (132  $\mu$ mol) in 25  $\mu$ l ddH<sub>2</sub>O over 20 min (5 X 5  $\mu$ l aliquots). Maintain pH at 8.5-8.8 with 20% (w/v) citric acid (2-5  $\mu$ l per addition of NaBH<sub>4</sub>). Once all NaBH<sub>4</sub> is added, move to room temp, flush tube with argon (or N<sub>2</sub>), and stir at top speed in dark for 2 hrs.
- 3. Cool to 4°C, lower pH to 5.0 with 5 N HOAc (~20  $\mu$ l) to destroy excess NaBH<sub>4</sub> (some bubbling will occur). Bring pH back to 7.8 with 5 N NaOH. Product should be H<sub>4</sub>PteGlu<sub>n</sub>. Check spectrum of 10<sup>-4</sup> dilution (see figure 2).

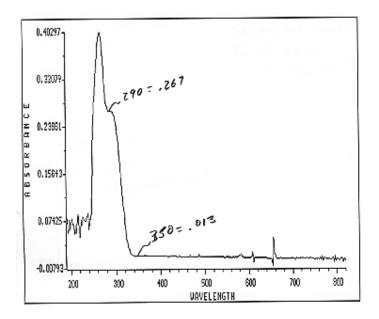


- 4. Add 6.4  $\mu$ l (80  $\mu$ mol) HCHO (37% soln), flush tube with argon, and incubate at 45°C, 15 min in dark to convert H<sub>4</sub>PteGlu<sub>n</sub> to CH<sub>2</sub>-H<sub>4</sub>PteGlu<sub>n</sub>.
- 5. Add 10 mg NaBH<sub>4</sub> (264  $\mu$ mol) in 50  $\mu$ l ddH<sub>2</sub>O over 20 min (5 X 10  $\mu$ l aliquots). Maintain pH at 8.5-8.8 with 20% (w/v) citric acid as before. Once all NaBH<sub>4</sub> is added, flush tube with argon and incubate at 45°C, 60 min in dark to reduce to 5-CH<sub>3</sub>-H<sub>4</sub>PteGlu<sub>n</sub>.

6. Remove stir bar, cool to 4°C, add 2-mercaptoethanol to 50 mM (~1.4  $\mu l$  of 14.3 M stock). Adjust pH to 5.0 with 5 N HOAc to destroy excess NaBH<sub>4</sub> (vortex till no more bubbling). Bring back to ~7.5 with 5N NaOH. Check spectrum of 10<sup>-4</sup> dilution (see figure 3). Store under argon (or N<sub>2</sub>) in dark at –20°C.



Notes: Starting material (PteGlu<sub>n</sub>) has maxima at 282 (E = 23.4 mM<sup>-1</sup>) and 350 nm (Fig. 1).  $H_4PteGlu_n$  has a maximum at 298 nm (E = 30 mM<sup>-1</sup>), and a min at 244 nm (Fig. 2). The final product (5-CH<sub>3</sub>-H<sub>4</sub>PteGlu<sub>n</sub>) has a max at 292 nm (E = 31.7 mM<sup>-1</sup>) and a min at 248 nm (Fig. 3). With PteGlu<sub>1</sub>, this procedure gave ~93% yield, and the final product had less than 3%  $H_4PteGlu_n$  based on its reaction with 5N HCl/60% formate:



Diluted 2  $\mu$ l of product into 200  $\mu$ l Tris, and then 10  $\mu$ l of this into 400  $\mu$ l Tris (7.5). Add 100  $\mu$ l 5N HCl/60% formate, heat at 80°C, 10 min, cool to RT and read (blank against buffer + HCl/formate). A peak at 350 nm is diagnostic of THF; lack of peak indicates CH<sub>3</sub>-THF.