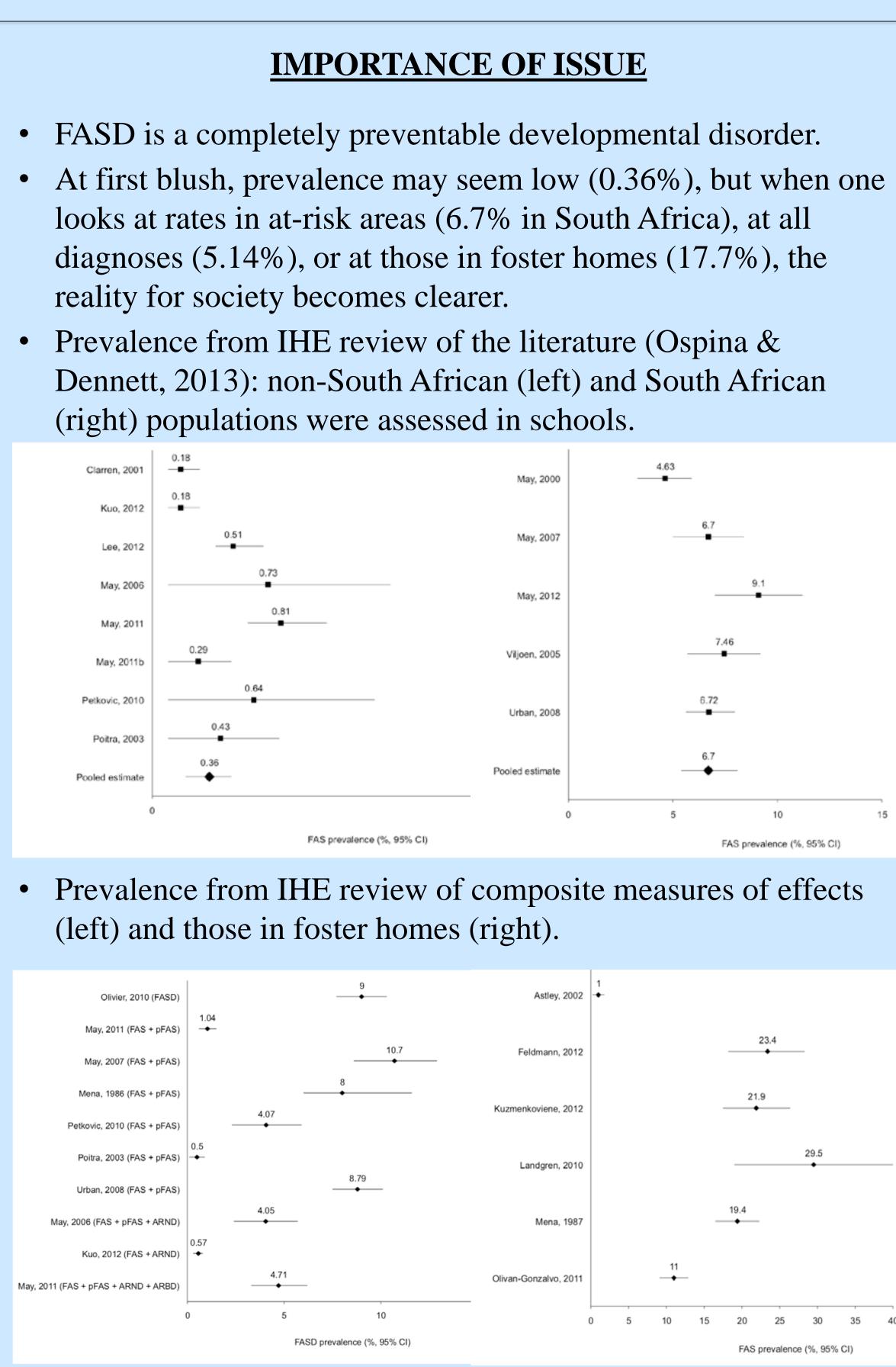


BACKGROUND

- Haycock (2009) posited that epigenetic mechanisms relate to fetal alcohol spectrum disorder (FASD) sequelae given the importance of the epigenome to nervous system development.
- Disruption of methylation and thus, gene expression during the embryonic period may negatively impact the developing fetus.
- Because alcohol disrupts liver 1-carbon metabolism resulting in a deficiency of the choline needed for methyl group transfer (Trimble et al., 1993), it is hypothesized that a similar disruption in methylation is occurring in the fetus in FASD.
- Poor nutrition status and irregularity of meals has been related to higher intake of alcohol (Santolaria et al., 2000). Thus, important nutrients may be low in chronic alcohol consumers.
- We propose that the positive effects of choline supplementation in alcohol-exposed fetuses and individuals are due to epigenetic mechanisms. The epigenome relies on methylation of DNA and on histone modification (Zeisel, 2009); choline is a major source of methyl groups (Zeisel, 2006).
- Here we provide the rationale for our hypothesis that epigenetic modifications enabled by choline supplementation will mitigate the effects of early alcohol exposure.

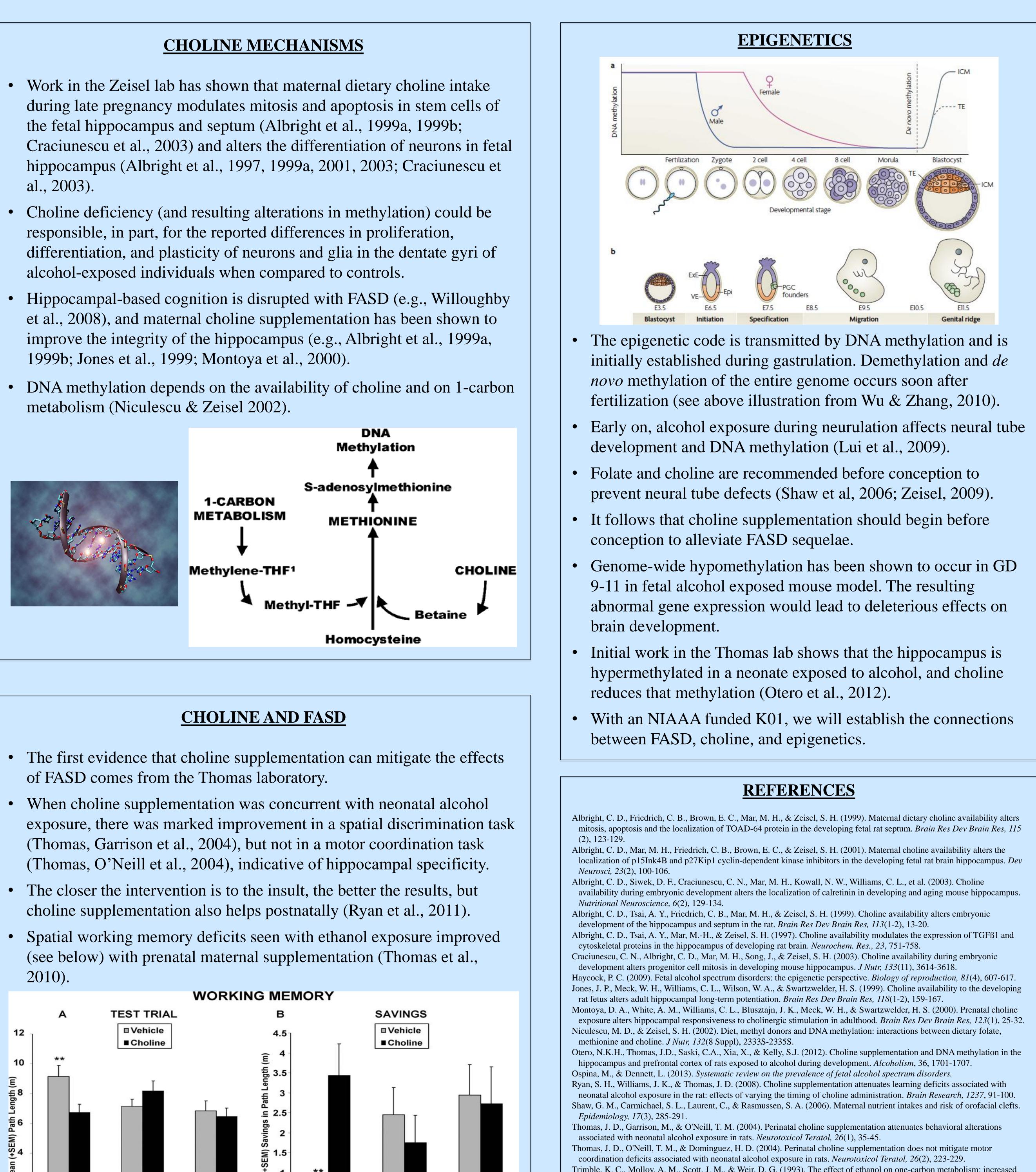


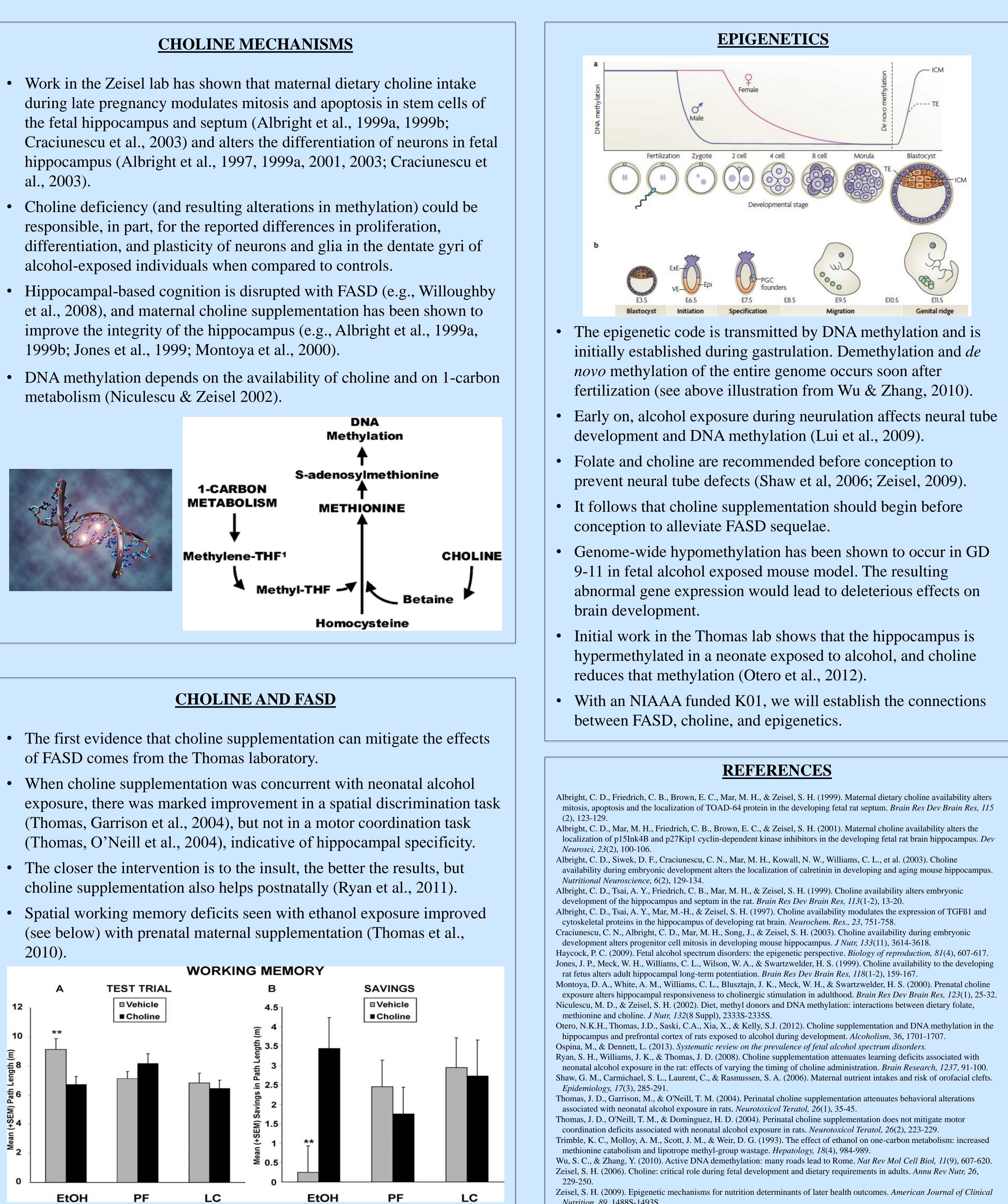
Choline and epigenetic mechanisms: potential for mediation of fetal alcohol spectrum disorders. Carol L. Cheatham,¹ Jennifer D. Thomas,² Philip A. May,¹ & Steven H. Zeisel¹ ¹ University of North Carolina at Chapel Hill Nutrition Research Institute

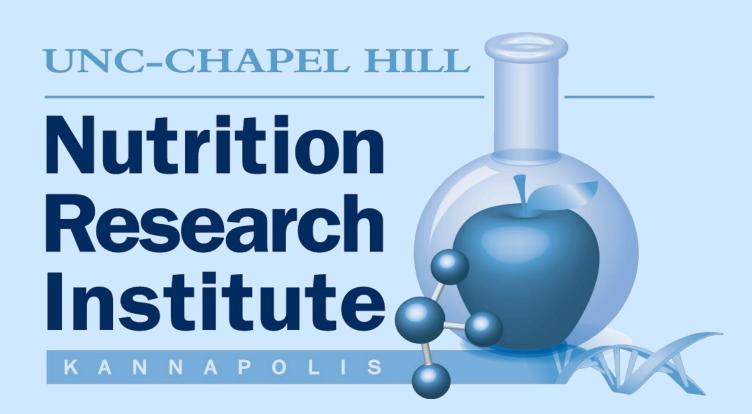
²San Diego State University

23.4 29.5 25 30 35 40 FAS prevalence (%, 95% CI)

- al., 2003).
- 1999b; Jones et al., 1999; Montoya et al., 2000).
- metabolism (Niculescu & Zeisel 2002).







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