

“OMICS” Assessment of the Placenta in Real Time

**Human Placenta Project
May 26-27, 2014**

Diana W. Bianchi M.D.

“The Assignment”

- Develop noninvasive markers of placental development and function
- Markers should permit longitudinal assessment
- Use “omic” tools to improve molecular definitions of placental biology and disease



Outline of Remarks

- **“Non-invasive”**
 - *In silico*, blood, saliva, urine, cervix
- **Intact cells**
 - Syncytiotrophoblast, cytotrophoblast
- **The genome and epigenome**
- **The transcriptome**
- **Importance of annotation**
- **No time for metabolome, proteome!**



In 6 minutes!

Cells: Trophoblasts as a Biomarker of Eclampsia



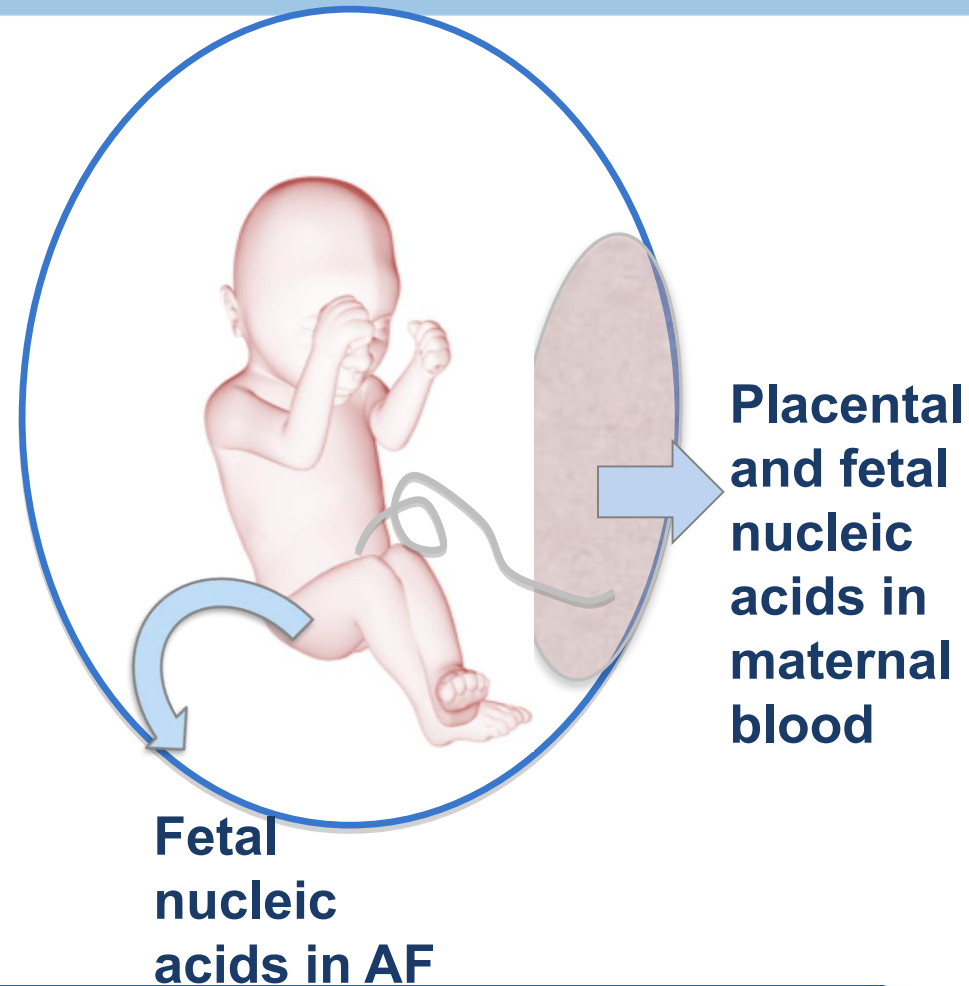
Georg Schmorl
1861-1932

- Multi-nucleated syncytial giant cells in lung capillaries of pregnant women who died of eclampsia
- Women with pre-eclampsia have 5-fold ↑ in number of circulating fetal cells
- Experimental model: injected placental supernatant from pregnant rabbits and caused seizures
- Re-emergence of interest in isolation of intact cells from maternal blood
(Lapaire O et al. *Placenta* 2007; 28: 1-5)

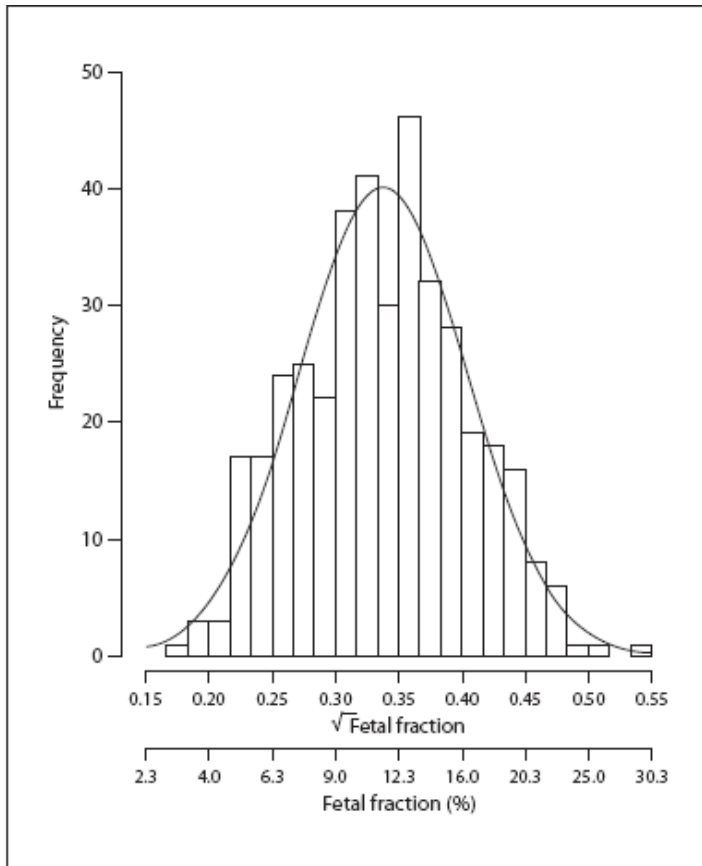
Cell Free Nucleic Acids as a Biomarker

Cell-free fetal (cff) DNA and RNA are present both in AF and maternal blood

- Compared with circulating cff nucleic acids, those in AF are:
 - in direct physical contact with the fetus
 - 100-200 fold more concentrated
 - uncontaminated by maternal nucleic acids



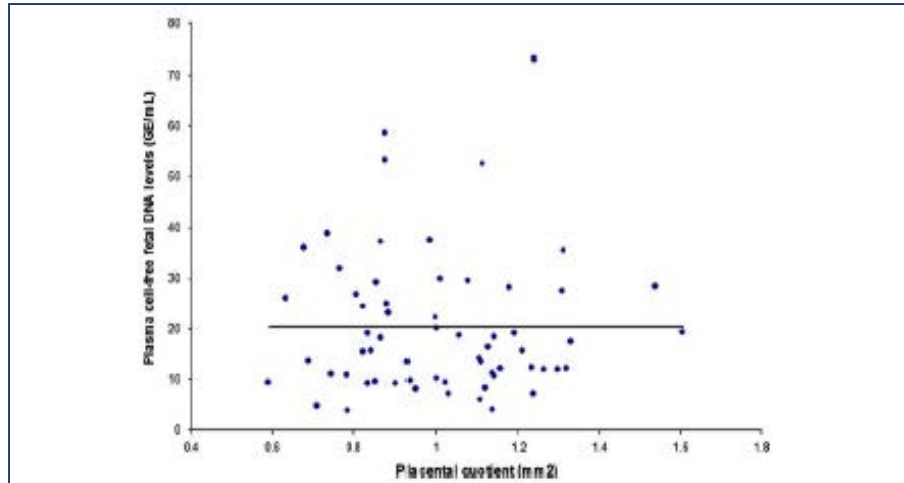
Fetal Fraction is a Placental Biomarker



- Fetal fraction has a normal distribution that peaks between 10 and 20% at 10-21 weeks of gestation
- It is primarily affected by four factors
 - Maternal BMI
 - Gestational age
 - Type of aneuploidy
 - Singleton vs. multiple

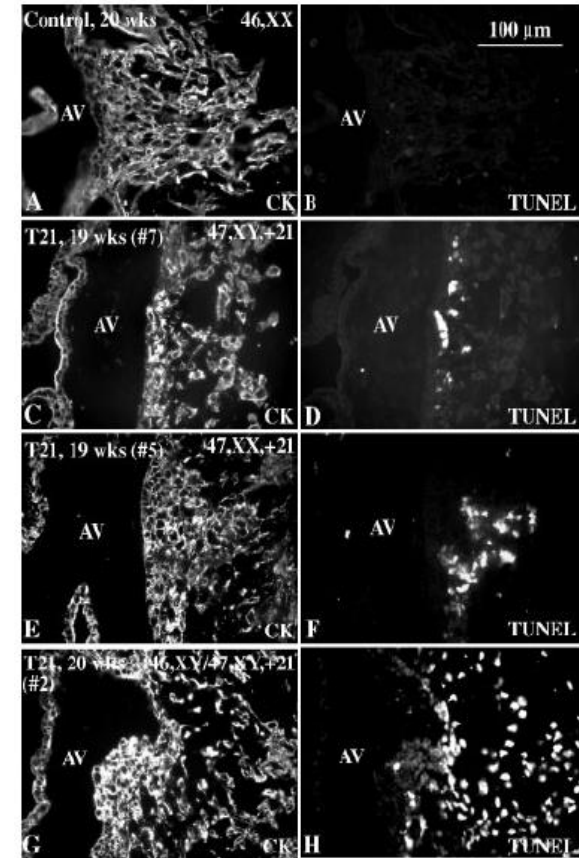
Ashoor et al. Fetal Diagn Ther 2012

Cell-Free Fetal (Placental) DNA in Maternal Blood



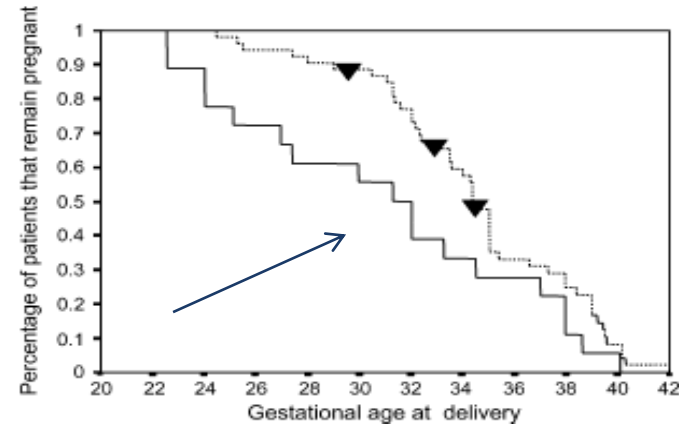
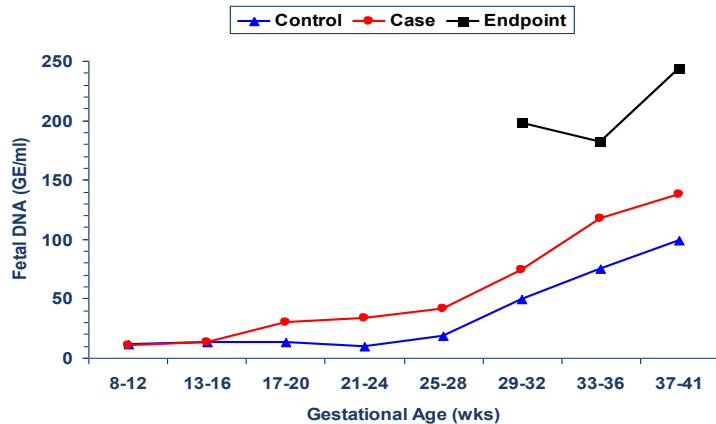
T. Wataganara, *Am J Obstet Gynecol* 2005

- Does not reflect placental volume
- Does reflect oxidative stress and apoptosis



A. Wright, *Am J Med Genet* 2004

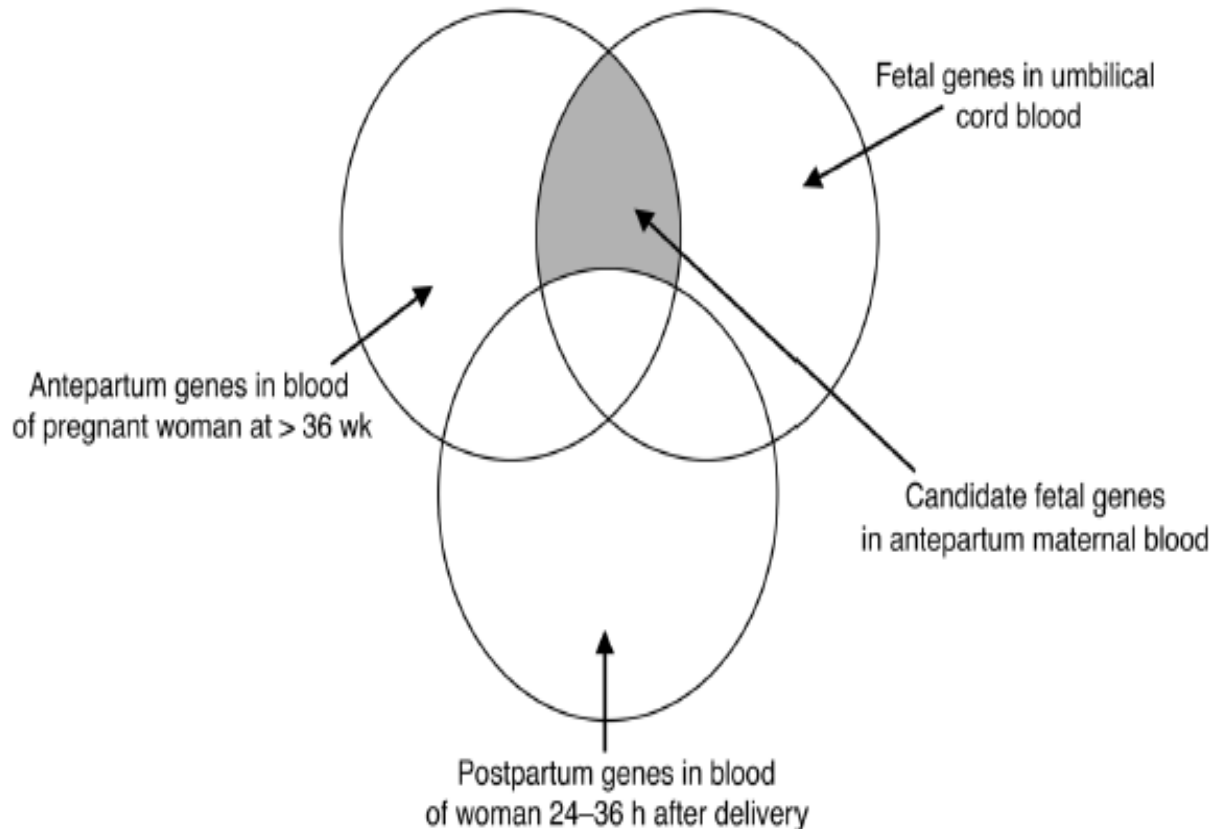
Cell-free DNA as a placental biomarker



- Increased DNA levels in maternal plasma in pre-eclampsia, unstoppable preterm labor, IUGR
- Increased tag counts can suggest the presence of confined placental mosaicism for aneuploidy
- Differential methylation occurs in placenta (epigenetics)
RASSF1A is hyper-methylated in placenta, hypo-methylated in blood

Approach to detection of feto-placental transcripts in maternal blood using cell-free RNA

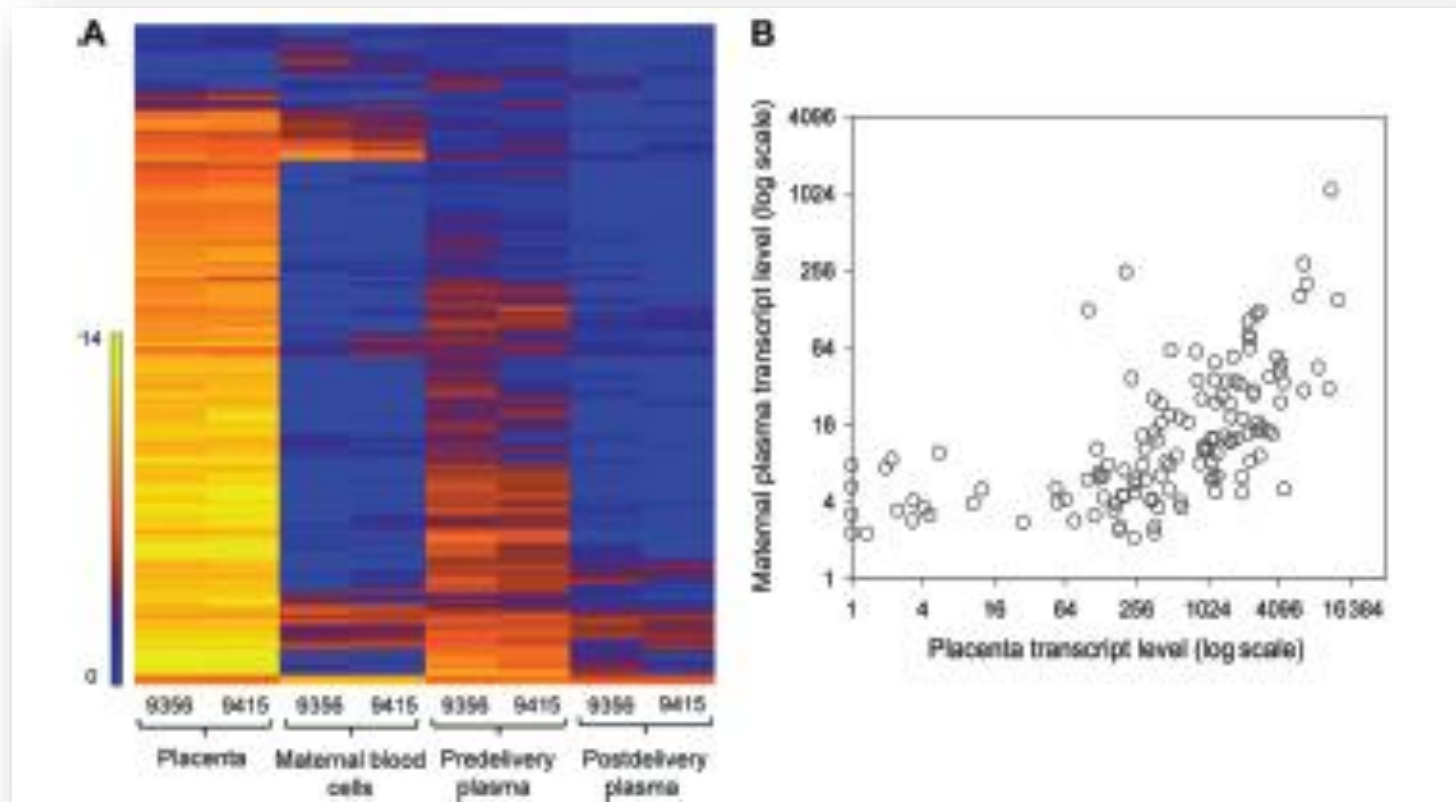
Maron et al. *J Clin Invest* 2007; 117: 3007-3019



- Studied whole blood and placental gene transcripts at term using microarrays
- Sensory perception genes up-regulated
- Fetal nervous system genes
- Immune-related genes
- Placenta-specific genes more easily detected in plasma

Placental-specific genes are more easily detected in maternal plasma compared to maternal blood

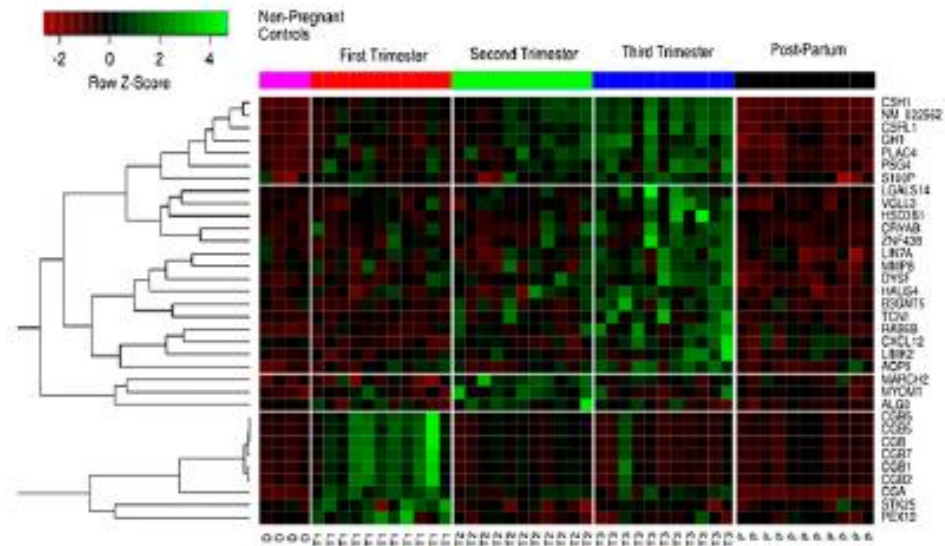
Tsui et al. Clin Chem 2014; April 16



Placental Transcripts Detected in Maternal Plasma

From Koh et al. *Proc Natl Acad Sci USA* 2014;doi/10.1073/pnas.1405528111

- *PLAC1*
- *PLAC4*
- *hCG*
- *PSG4*
- *GH1*
- *CSHL1*
- *PAPP-A*

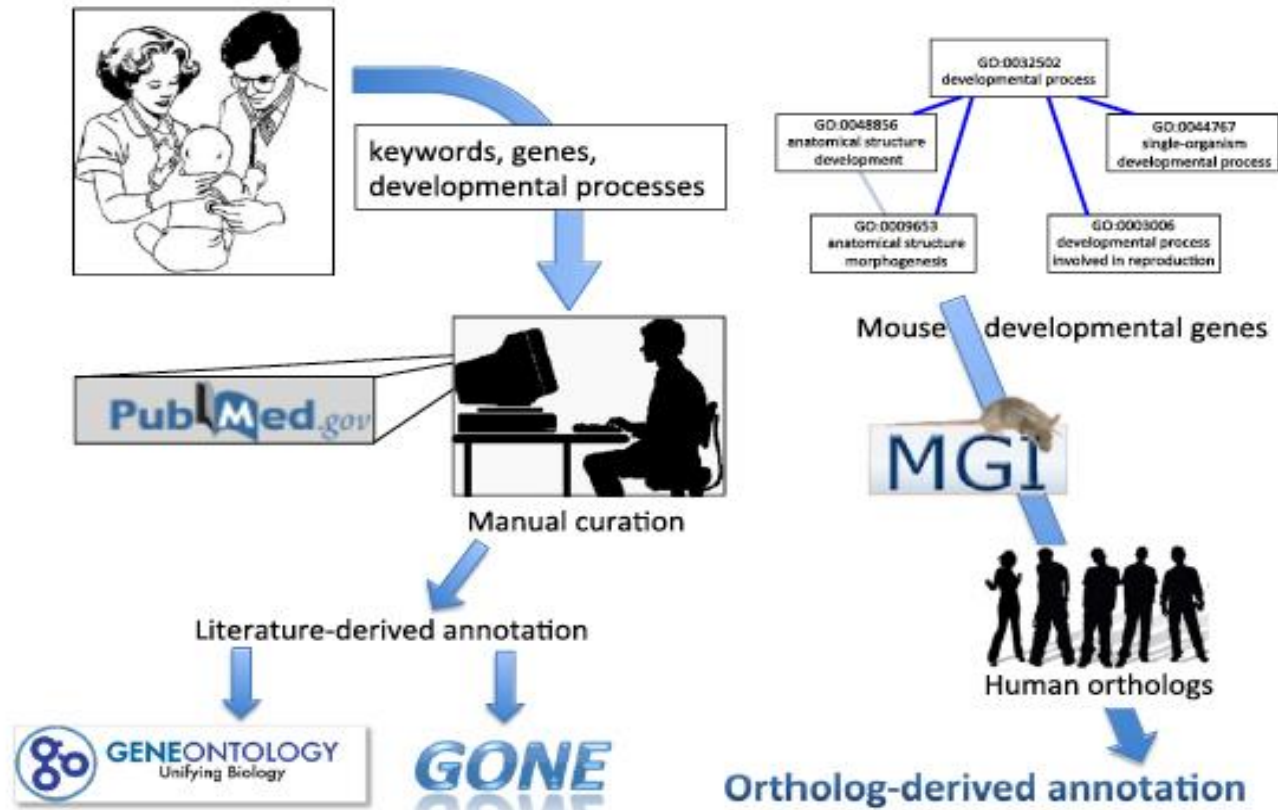


-MicroRNAs: very stable in plasma
-Many placenta-specific miRNAs
-Placental-miRNA expression associated with infant behavior

- How do you know if a transcript is unique to the placenta?
- Publicly-available atlas of human protein-encoding transcriptome (Su *PNAS* 2004)
- Relative tissue expression patterns of individual gene probes across 78 normal human tissues including fetal brain, liver, thyroid, lung (15-33 wks) and **placenta** (n=4)
- Definition of organ-specific gene expression
 - expression value > 30 MoM in a single tissue type
 - no unrelated tissue expression $> 1/3$ of max. expression value

Developmentally-appropriate annotation is critical!

<http://dflat.cs.tufts.edu>



From: HC Wick et al. Bioinformatics 2014; 15:45