sncRNAs in healthy human pregnancies

Investigating the Small Non-Coding RNA Transcriptome of the Human Placenta

Nikita Telkar^{1,2,3}, Greg L Stewart¹, Michelle E. Pewarchuk¹, David E. Cohn¹, Wendy P. Robinson^{1,2,3}, Wan L. Lam¹

¹British Columbia Children's Hospital Research Institute, Vancouver, British Columbia, Canada

²British Columbia Cancer Research Centre, Vancouver, British Columbia, Canada

³Department of Medical Genetics, University of British Columbia, Vancouver, British Columbia, Canada





The placenta is the fundamental organ required throughout pregnancy. Precise gene regulation in the placenta is crucial in maintaining a healthy pregnancy¹.

Small non-coding RNAs (sncRNAs) are known regulators of gene expression, with certain microRNAs already been associated with placental disorders ^{2,3} – however, the global sncRNA profile of healthy placentas remains underinvestigated.

Impact

SncRNAs have recently been introduced as diagnostic / prognostic markers in several diseases and disorders

Elucidating changes in sncRNA expression within the placenta throughout pregnancy would aid in early and better-informed intervention and care of pregnant women.

Hypothesis

The expression of sncRNAs changes dynamically throughout pregnancy in the healthy human placenta

Methods

30 control placentas at stages of gestation

Trimester 1 = 5, Trimester 2 = 16, Trimester 3 = 9

Fetuses: 17 Females, 13 Males

Total RNA extracted and sequenced

linear model:

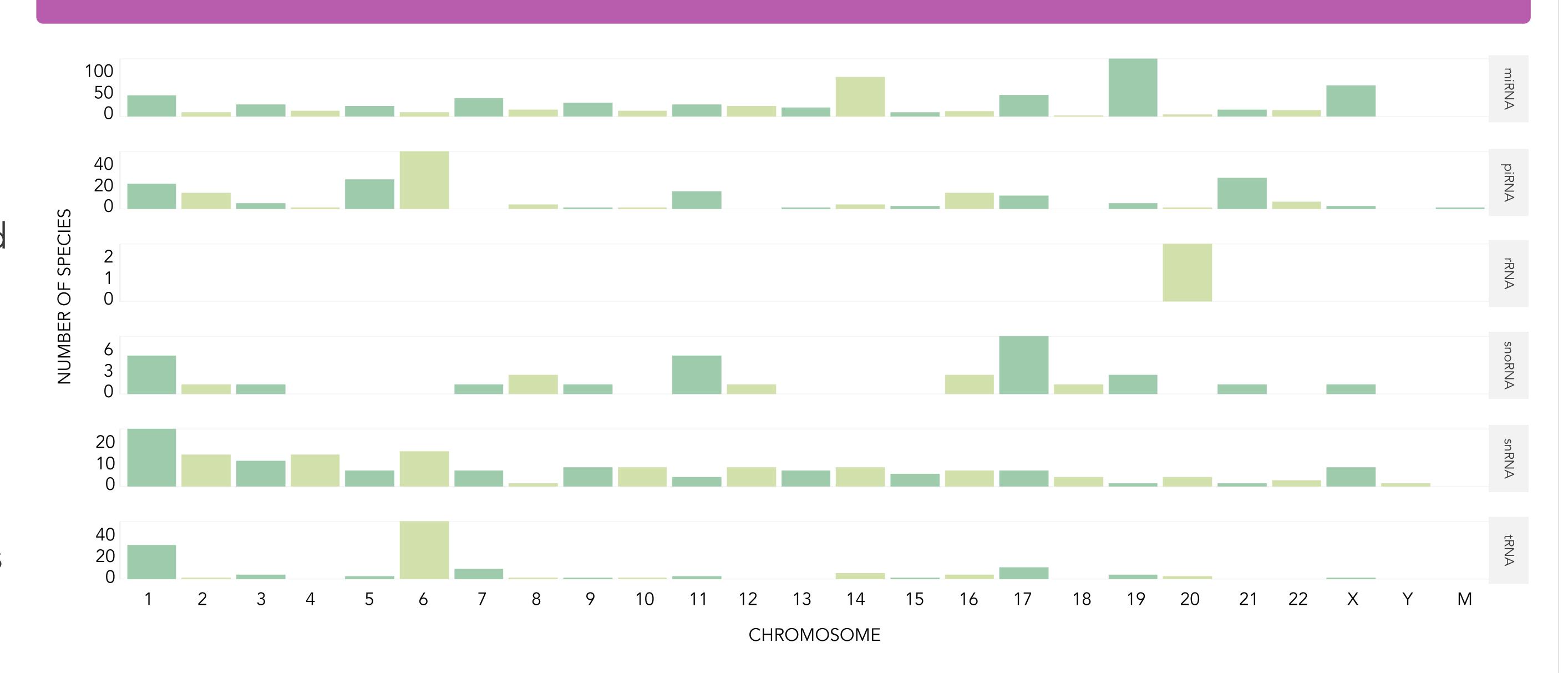
~sncRNA + Trimester + Sex + Flow Cell

Conclusions

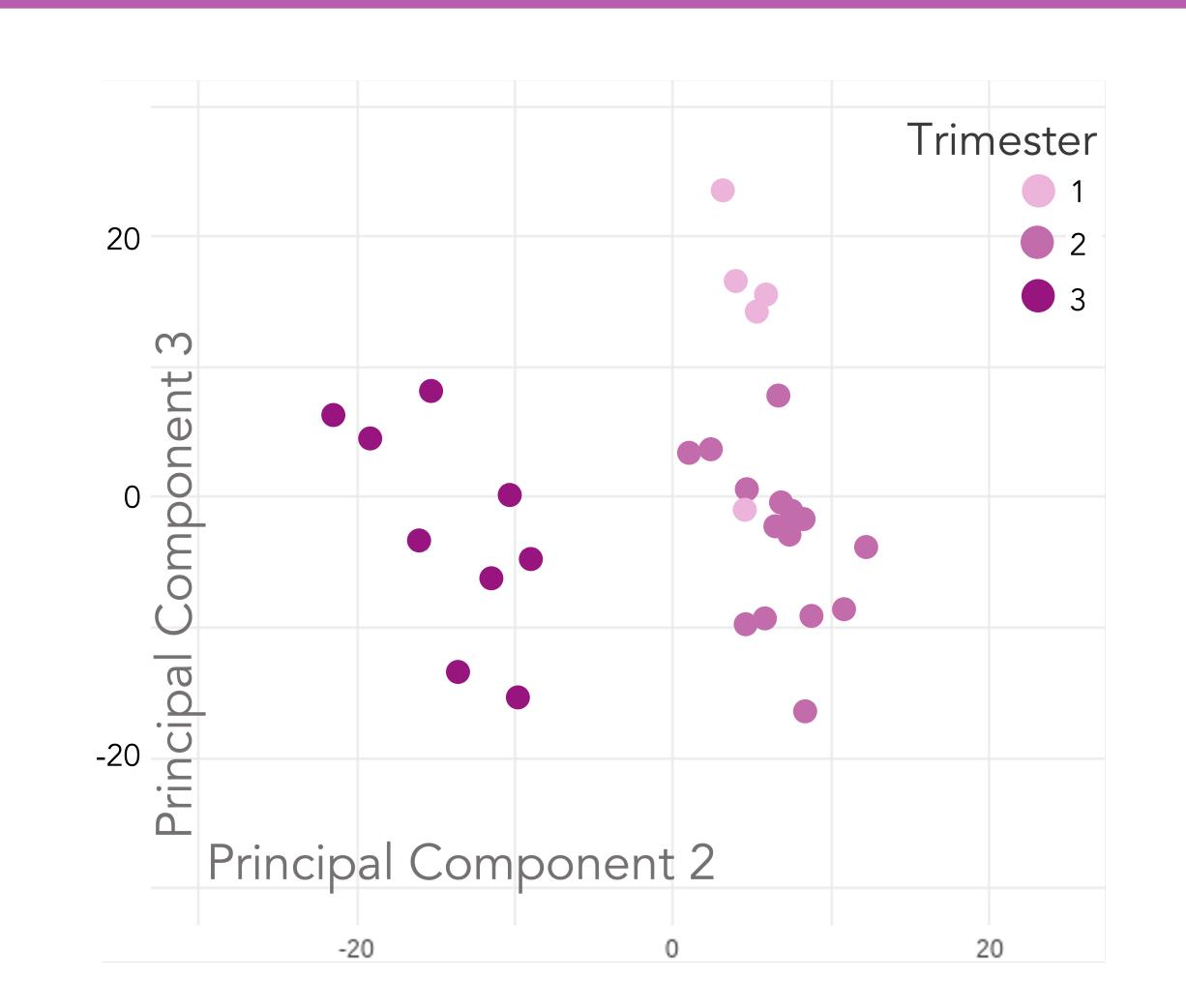
The human sncRNA profile changes with gestational age, indicating timely and fixed regulation of gene expression.

Technical factors can significantly contribute to the varied expression patterns, and need to be accounted for.

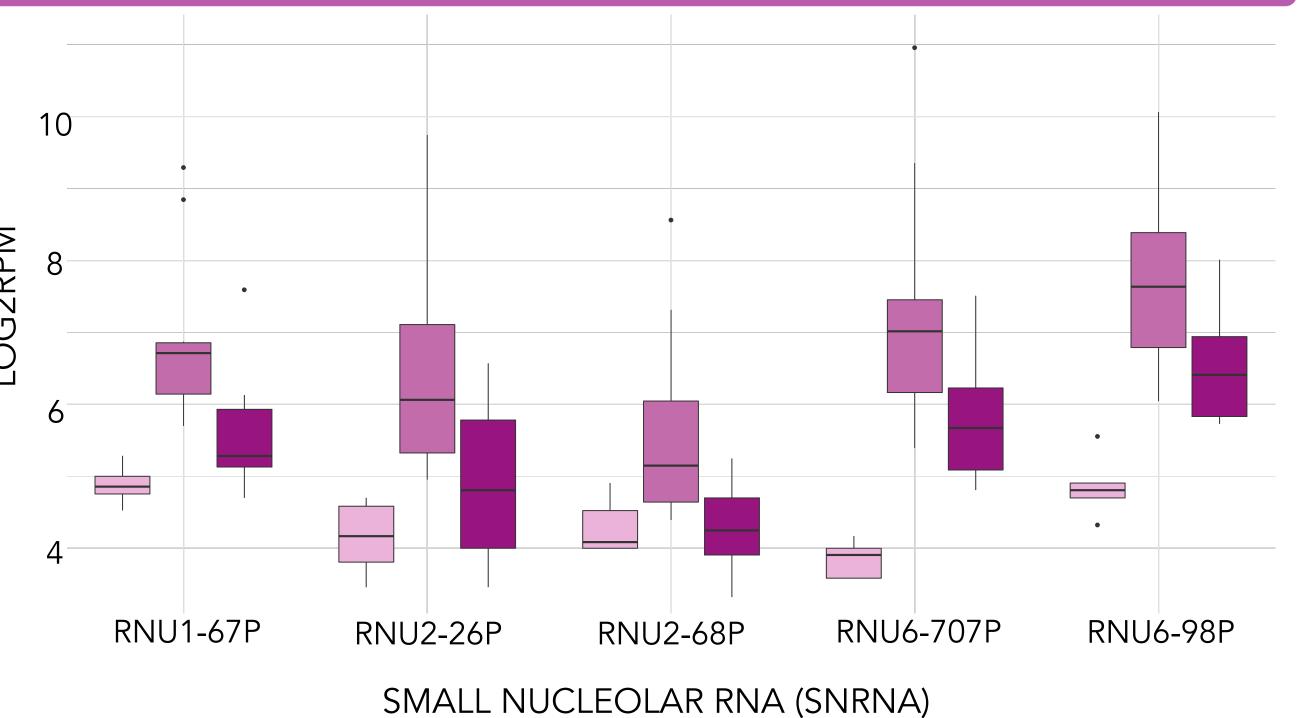
Placental differentially-expressed sncRNAs are present genome-wide



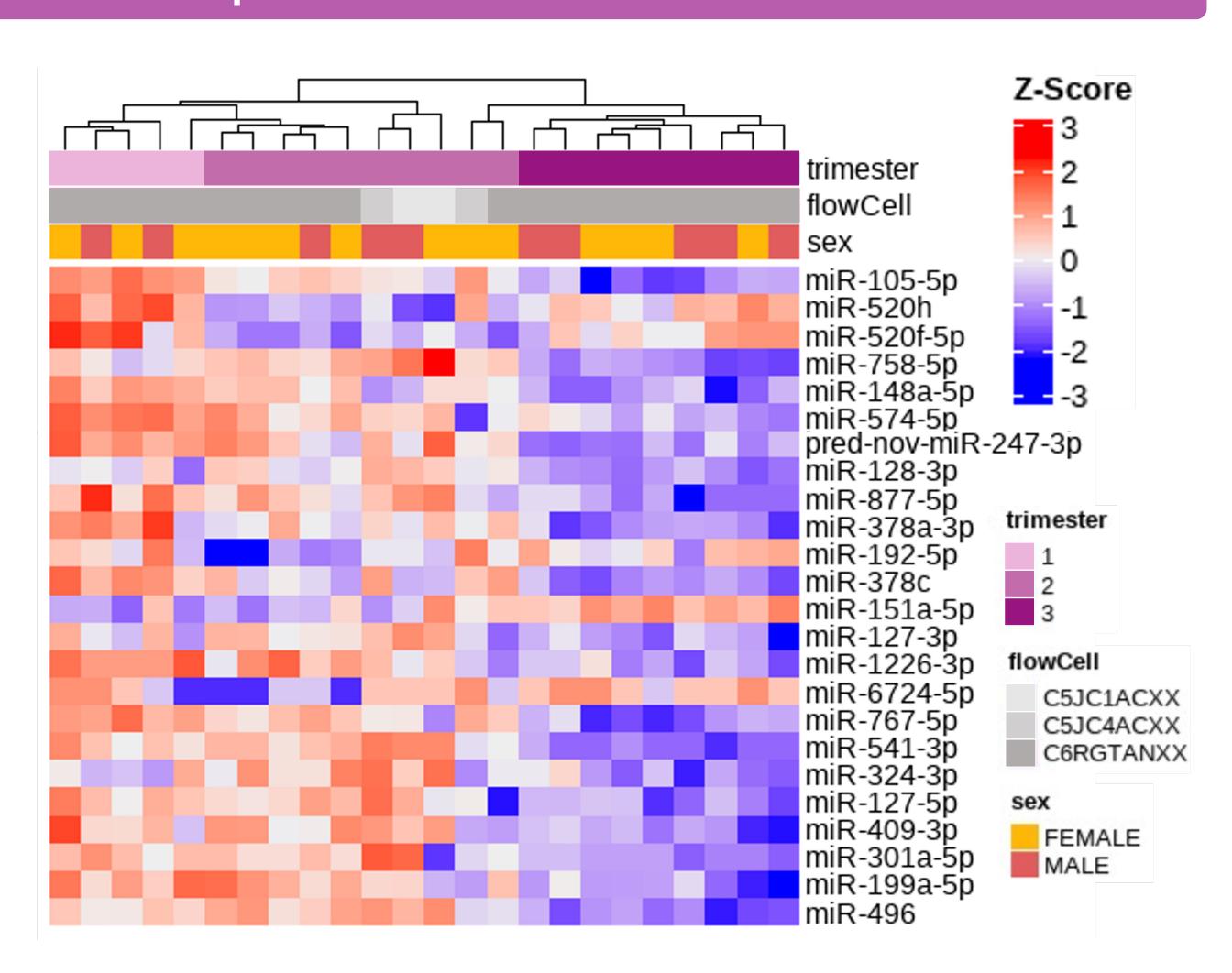
Samples separate by trimester



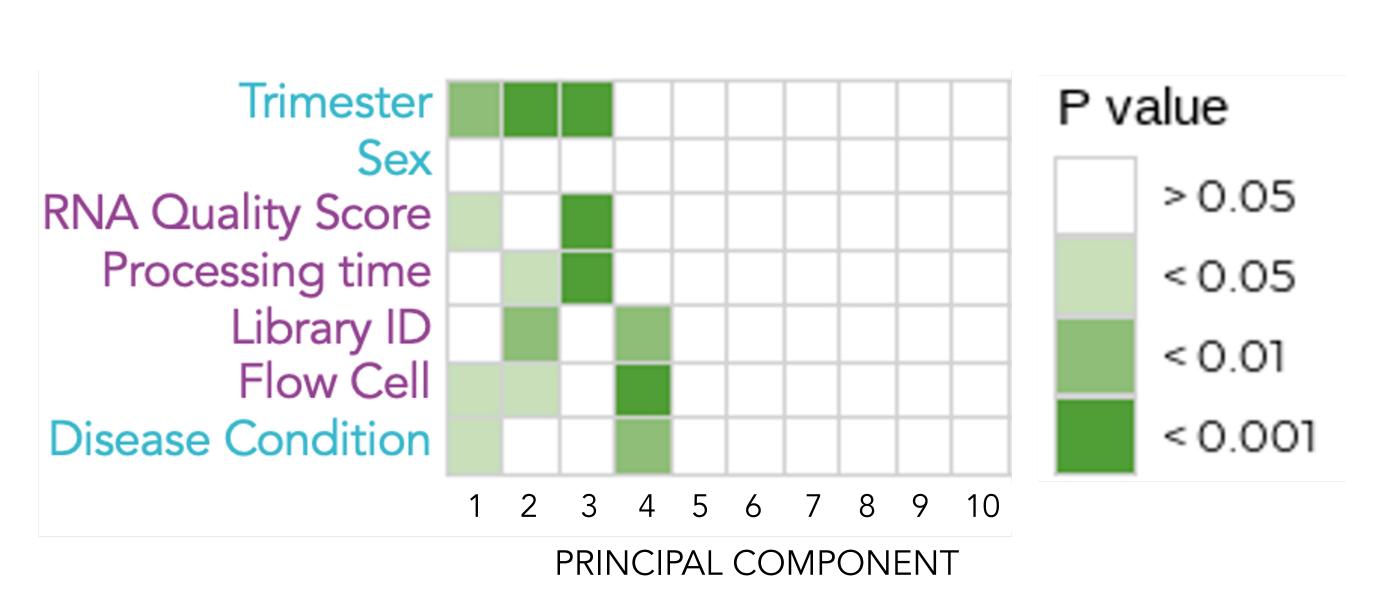
Chromosome X shows sncRNAs with trimester-dependant expression



24 miRNAs (1 novel) show changes in expression within all 3 trimesters



Non-biological factors may also alter expression



1. B. Cox, et al., (2015). 2. J.-F. Mouilletet al (2015). 3. D. M. Morales-Prieto, et al., (2013).