AUTOMATION OF THE ESTIMATION OF HUMAN OVARIAN RESERVE

February 2010

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My Background

- Senior Research Fellow in Maths & Computer Science
 - University of St Andrews, Scotland
- Constrained optimisation & minimisation
 - Roger Fletcher was my MSc supervisor
- Casimir forces
 - similar to van der Waals forces poorly understood,both in theory & practice
- Symmetry & search
 - where growth in both search space & number of symmetries is super-exponential



Medical/Biological Research

- Co-director of the Wallace Kelsey Research
 Foundation
 - www.wallacekelsey.org.uk
- We are interested in the late effects of cancer treatments on the survivors of childhood cancer
 - □ 70% 80% of children with cancer survive
 - □ The radio- and chemotherapies also damage healthy cells
- In particular, we concentrate on the fertility of survivors



There are two questions

- Can we quantify the effects of radio- and chemotherapies in order to predict early menopause?
 - □ We published papers on this in 2004 & 2005
 - We are working with the US CCSS on new techniques based on new data and new models
- Can we predict age of menopause for healthy women?
 - □ Reproductive biologists have no good results
 - But we need this as our baseline calculation

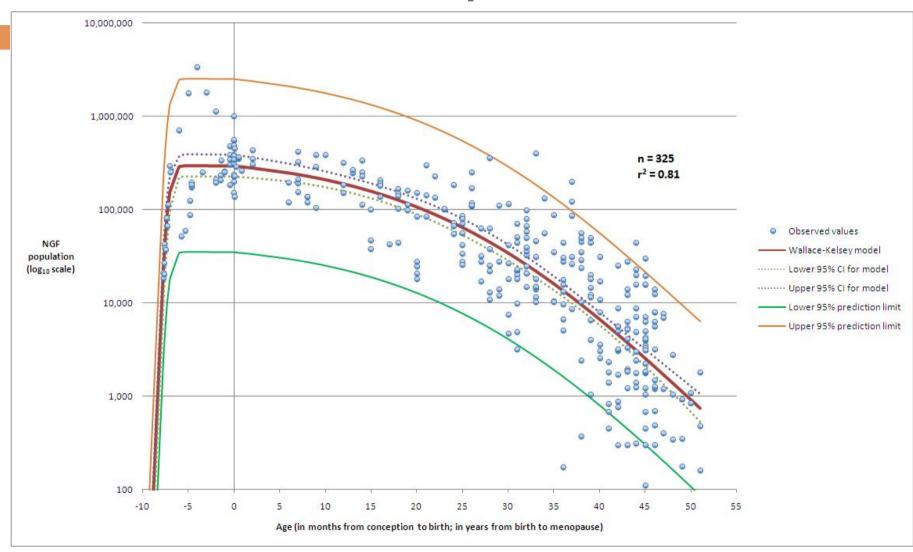


Non-growing follicles (NGF)

- The human ovary contains a population of nongrowing follicles
 - □ oocytes, ovocitos
- Some of these are recruited every 80-90 days
 - primary, secondary and antral follicles
 - most die off through atresia, but some become eggs
- □ When less than 1,000 remain, menopause occurs
 - □ so the population defines ovarian reserve in women
 - and the rate of decline is the "biological clock"



The Wallace-Kelsey Model



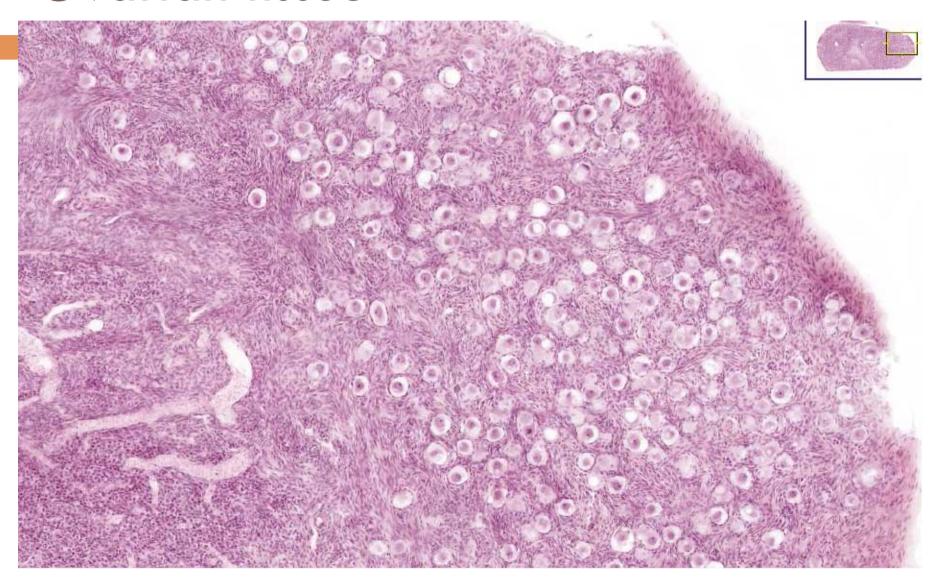


The Wallace-Kelsey Model

- Wallace WHB, Kelsey TW (2010) Human Ovarian Reserve from Conception to the Menopause.
 - □ PLoS ONE *5*(1): e8772. doi:10.1371/journal.pone. 0008772
 - ☐ The first model of ovarian reserve from conception
 - □ Agrees with separate studies on ages at menopause
- All the data was obtained by hand
 - Ovaries are sliced, stained & photographed
 - Sample population counts are integrated to get an estimate for the entire ovary



Ovarian tissue



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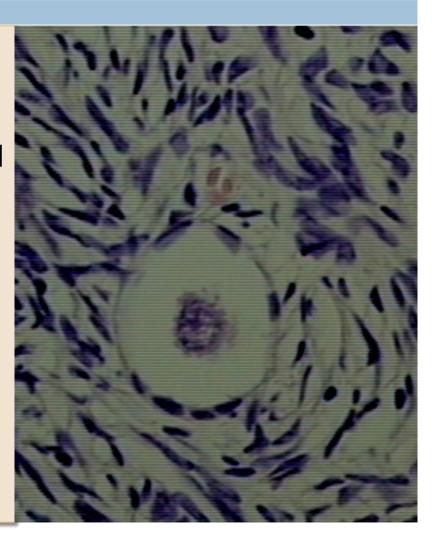


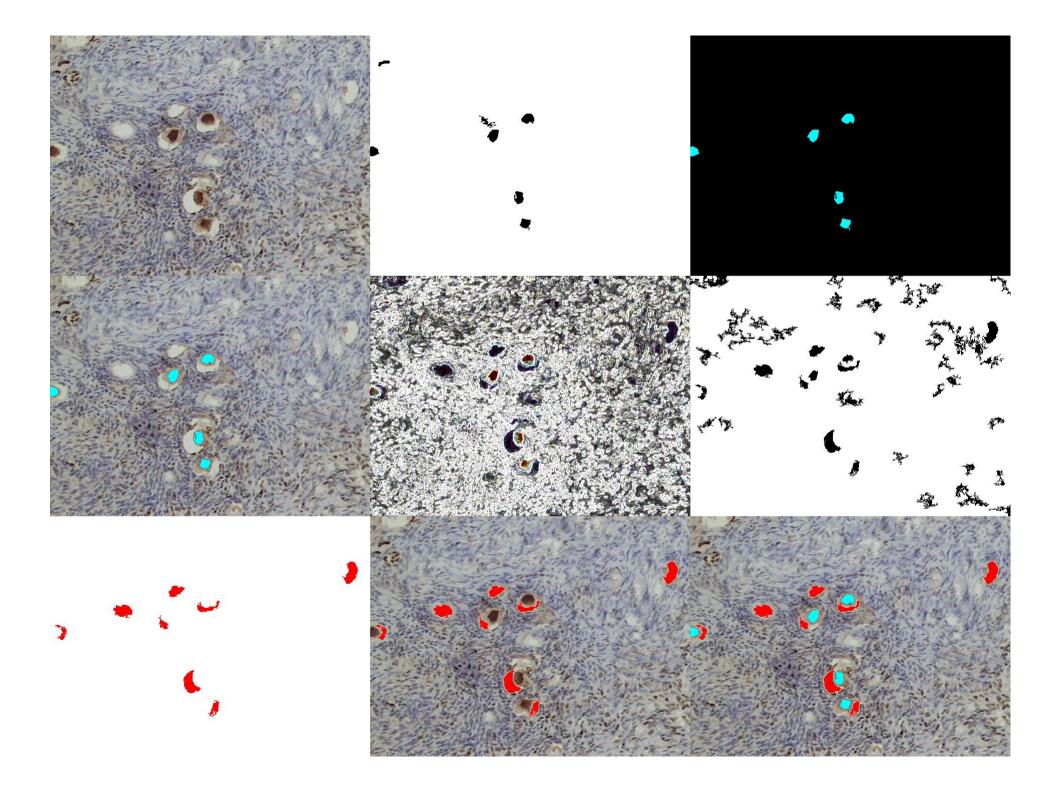
- Luis Castillo
 - Clinical Director, Fundacion Peluffo Giguens
 - Oncologist, also interested in late effects
- Pancho Coppola
 - Gynocologist at Perreira Rossell
- Benedicta Caserta
 - Histopathologist at Perreira Rossell
- We obtained ethical approval in 2008 for the study
 - □ I'm here to write up our results and submit to a journal



- The idea is to automate the population counts
 - □ input photos to a system that automatically identifies and counts the follicles that define ovarian reserve
 - more accurate than humans
 - we can process more of the ovary
 - and hence reduce errors at the integration stage
- It should be possible to count all the follicles in an entire ovary
 - but that's the next study
 - ☐ if we get funding

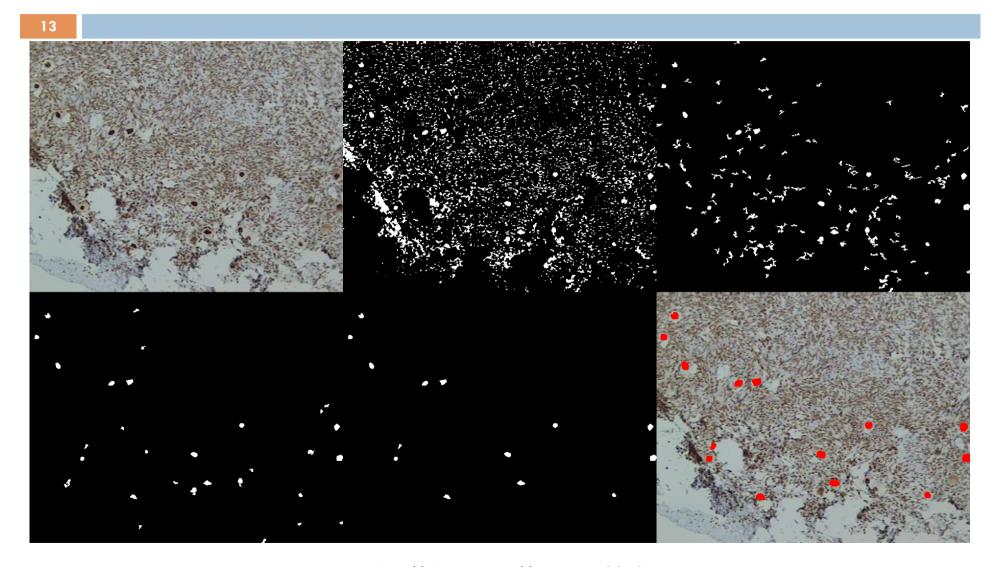
- Counting follicles at high magnification is easy
 - but you need more photos and more lab time
- Our main advance is finding a stain that allows us to identify follicles at low magnification
 - □ H & E is the standard stain
 - ☐ We use PCNA







Good results even at 100x



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- We can tune the code to be conservative...
 - ☐ few false positives
- □ ... or liberal
 - few false negatives
- The estimated true population is in the middle
- We still have some technical issues to resolve
 - what if we change the camera (or microscope, or technician)?
- But we will submit our paper for review in March

Muchas gracias, y he aprendido mucho

