



Uploaded to the VFC Website

▶▶▶ 2016 ◀◀◀

This Document has been provided to you courtesy of Veterans-For-Change!

Feel free to pass to any veteran who might be able to use this information!

For thousands more files like this and hundreds of links to useful information, and hundreds of "Frequently Asked Questions, please go to:

[Veterans-For-Change](#)

If Veterans don't help Veterans, who will?

Note:

VFC is not liable for source information in this document, it is merely provided as a courtesy to our members & subscribers.



Epigenetics and women's health research: an interview with Professor Steve Conlan, Swansea University

Published on February 24, 2015 at 6:04 AM



Please can you give a brief introduction to your research?

My name is Steve Conlan. I'm a professor of cellular microbiology in the College of Medicine at Swansea University. The College of Medicine is a very new college in the UK. We're ten years old this year, and we work at the interface with colleagues in hospitals and very closely with businesses.

Our research focus is on women's health, which involves gynaecological investigations into infertility in women and into gynaecological oncology where we're most interested in uterine cancer, the lining of the uterus, the endometrium, and also ovarian cancer.

Our research into gynaecological oncology focuses around understanding mechanisms of how genes are regulated or how they become dysregulated in a disease; and also the effects that has on the surface of the endometrium and also the function of the ovaries.

How do you try to understand these mechanisms?

We take a conventional approach using molecular and cell biology and we also use the application of nanotechnology into the investigations.

On the molecular biology side, we're very interested in the epigenetic mechanisms that can go wrong in cancer, and also how the transcription mechanisms can go wrong.

We don't see a difference between transcription and epigenetics. We think these are completely intertwined, and we focus on them both at the same time.



When did you start to focus on epigenetics?

It was about five years ago when we started to push our epigenetics investigations. We realised that the technologies that were available were very limiting.

We've been working over that time with a company – Porvair – to develop disruptive technologies that are going to make epigenetic investigations much simpler, much more effective, faster, and aid us in pushing forward our research.

Please can you give an overview of Chromatrap?

We've developed a technology called Chromatrap, which essentially is moving from bead based methods of DNA enrichment to a spin column.

We've also cut down experimental times from several days to being able to do the whole process in a day, and that just allows us to tackle more samples.

How important is high-throughput analysis in your research?

We work very closely with the hospital. We have patient samples coming in every day so the more samples we can get through, the more robust our data analysis is and a better understanding of the disease processes we can elucidate from that.



What types of modification do you focus on and why?

We're very interested in how epigenetic markers, DNA methylation, histone acetylation are changed in cancers. If we can understand that, then we can begin to think what drugs can we employ in these cancers to make the cells in those patients return to normal?

We focus on histone modifications. There are a multitude of modifications that we can look at, and we need specialized antibodies against each of those. If we can understand that, we understand the picture of how a cancer may progress.

Then we target enzymes that are involved in this process such as histone deacetylases using drugs that are available for treating leukemias and seeing if we can we apply those to solid tumor cancers.



What impact do you think the Chromatrap system will have?

Epigenetic analysis has been complex; it's been a difficult technique. The system that we set up – the Chromatrap system – allows us to tackle high-throughput analysis. This is going to lead to the idea of using epigenetic targets in personalised medicine.

The approach that we've developed lends itself to high-throughput automation, and this is where we need to be going if we're going to analyse the epigenome of every patient.

For example, being able to target drug use in patients. We're going to need to know their epigenetic profiles and simple disruptive technologies that can be embedded into pathology labs are going to enable us to do that.

What do you think the future holds with regards to companion diagnostics?

Companion diagnostics is a rapidly emerging area, and there are a few examples out there of how tests are essential before you can give a drug, maybe because it's too toxic if you don't or maybe it just won't benefit the patient if you don't.

At the moment, our epigenetic research is about aligning drugs with biomarkers. It's at a very early stage, but it is an evolving area, and it will make an impact in the future.

Where can readers find more information?

www.chromatrap.com



About Professor Steve Conlan

Steve Conlan is Professor of Molecular and Cell Biology, head of Reproductive Biology and Gynaecological Oncology research and Director of the Centre for NanoHealth at Swansea University, Wales.

Prof Conlan works closely with the life science and healthcare industrial sector, and has active collaborations with SMEs and multinational companies. As Director of Strategic Partnerships, and member of Swansea University's Internationalisation Strategy Group, he leads strategic collaborative initiatives with Grenoble - France, Houston - US and Suzhou and Wuhan – China.

Prof Conlan holds an honorary consultant position in the Abertawe Bro Morgannwg NHS Board, working with clinical colleagues in Singleton Hospital Swansea and The Princess of Wales Hospital Bridgend. He is a Senior Affiliate Member of The Methodist Hospital Research Institute, Houston Texas, and Distinguished Professor at Xi'an Jiaotong University, Suzhou Academy, China.

Chromatrap® - Porvair Sciences Ltd



Clywedog Road South
Wrexham Industrial Estate
Wrexham LL13 9XS
United Kingdom
PH: 44 (1978) 666239
Email: info@chromatrap.com
Visit [Chromatrap® - Porvair Sciences Ltd](#) Website

Company Background

Chromatrap® is a product of Porvair Sciences, a wholly owned subsidiary of Porvair plc. We are one of the largest manufacturers of Ultra-Clean microplates, 96 well well filtration plates and Microplate handling equipment for life science and synthetic chemistry. With offices and Class VIII clean room manufacturing located in the UK, combined with a world-wide network of distributors and dedicated distribution hub in the USA, we pride ourselves on our continuous innovation, research and flexibility to meet customer demands. We offer OEM production and contract manufacturing through our North Wales facility.

Our porous polymeric material, BioVyon™, whose chemical functionalisation can endow it with internal surface properties individually configured to capture and separate target species out of difficult mixtures, has opened up many possibilities in the field of BioSciences where molecules of interest such as DNA, RNA, proteins etc can be selectively pulled out of complex mixtures of biological origin. The materials have proven to be a remarkably good substrate for accepting novel chemistries such as the organically bound Protein A and Protein G in Chromatrap®.

Using our 25 years experience of microplate manufacturing, Porvair Sciences has now developed a high-throughput bead-free ChIP assay based on our filtration plates containing our Chromatrap chemistry. Chromatrap-96 enables large scale epigenetic screening to become a reality in many laboratories and eliminates many of the long and laborious steps previously undertaken in such work.