



Assessing tumour antigens in archived liver samples: a clinical research experience.



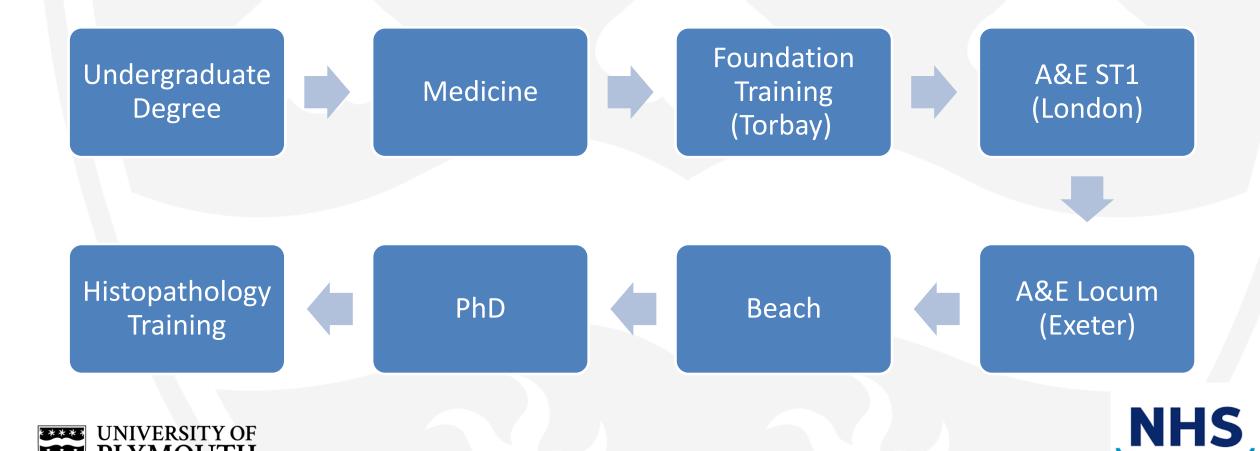
Ollie Rupar 10th June 2022

Oliver.Rupar@nhs.scot





My Timeline



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Hepatology Research Group

Dedicated to improving the understanding and treatment of liver disease

The hepatology research group (HRG) is an integral part of the Institute of Health and Care Research: infection, immunity and inflammation.

We utilise state of the art laboratory facilities based in the Derriford Research Facility, and the world class clinical research strengths of the Faculty of Health and Plymouth Hospitals NHS Trust (PHNT).

We work in unison with the South West Liver Unit, at Plymouth Hospitals NHS Trust, providing a full range and secondary, tertiary and community Hepatology services to the South West region, including assessment for liver transplantation, TIPS and liver cancer therapy.

The research team run several commercially sponsored clinical trials in hepatitis C therapy, non-alcoholic steatohepatitis (NASH), primary biliary cholangiopathy (PBC), alcoholic liver disease and liver failure amongst others, through the clinical research facilities of The Lind Research Centre at Derriford Hospital.

Hepatology Research Group Annual Reports

- Download the 2017 report
- Download the 2018 report
- Download the 2019 report
- Download the 2020 report

The proad themes of the hepatology research group are:

- Protection from hepatitis C virus infection (principal investigator) PI Professor Matthew Cramp
- Lipid metabolism and the pathogenesis and treatment of non-alcoholic steatohepatitis – PI Dr David Sheridan
- Alcoholic hepatitis Pl Ashwin Dhanda
- Molecular virology of hepatitis C and other her atitis viruses PI Dr Dan Felmlee





Health and Care Professionals ▼

Researchers -

Patients and the Public ▼

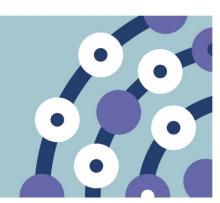
Partners and Industry 🔻

About us ▼

Q Search...

Research capability funding

Home > Researchers > Collaborations, services and support for your research > Research capability funding



NIHR research capability funding (RCF) is allocated to NHS organisations that undertake NIHR research, to help them maintain research capacity and capability.

What are the aims of research capability funding?

The aims of RCF funding are to:

- help research-active NHS organisations to act flexibly and strategically to maintain research capacity and capability
- support the appointment, development and retention of key staff undertaking or supporting people and patientbased based research
- contribute towards the costs of hosting NIHR-funded or 'adopted' research that are not currently fully covered across NIHR's programmes, and that are not met in other ways.

Documents

Research capability funding frequently asked questions

Research capability funding allocations 2022-23 (XLSX - Excel)

<u>Q&A on the outcome of the 2018 research capability funding review</u>

<u>Use of NIHR research capability funding in 2019-20</u>







BIOLOGY

nature



Century-Old Tumors Offer Rare Cancer Clues

DNA sequences from 100-year-old tumor samples could bolster childhood cancer research

By Heidi Ledford, Nature magazine on May 11, 2017



NHS Trust

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Credit: Ktsdesign Getty Images

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Heidi Ledford and Nature magazine

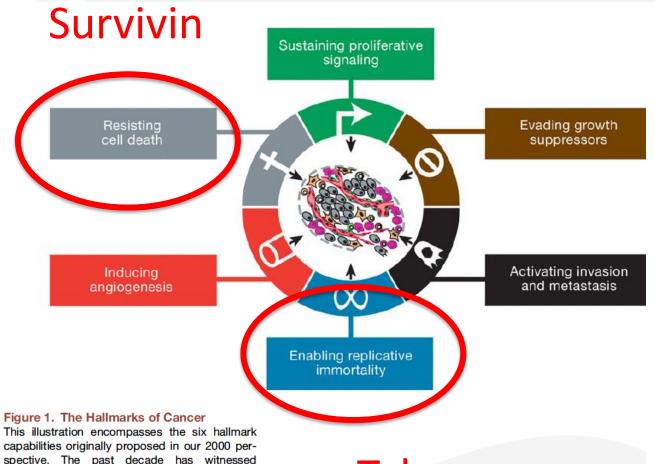
THE SCIENCES

Frederick Sanger, Father of DNA Sequencing, Dead at 95

Ewen Callaway and Nature News Blog







capabilities originally proposed in our 2000 perspective. The past decade has witnessed remarkable progress toward understanding the mechanistic underpinnings of each hallmark.

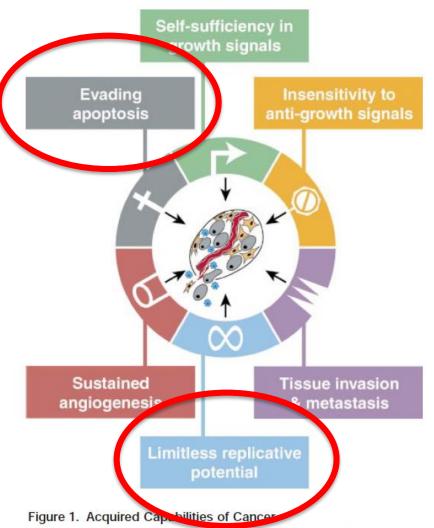
Telomerase



Hanahan & Weinberg, 2000 & 2011:

https://doi.org/10.1016/S0092-8674(00)81683-9,

https://doi.org/10.1016/j.cell.2011.02.013

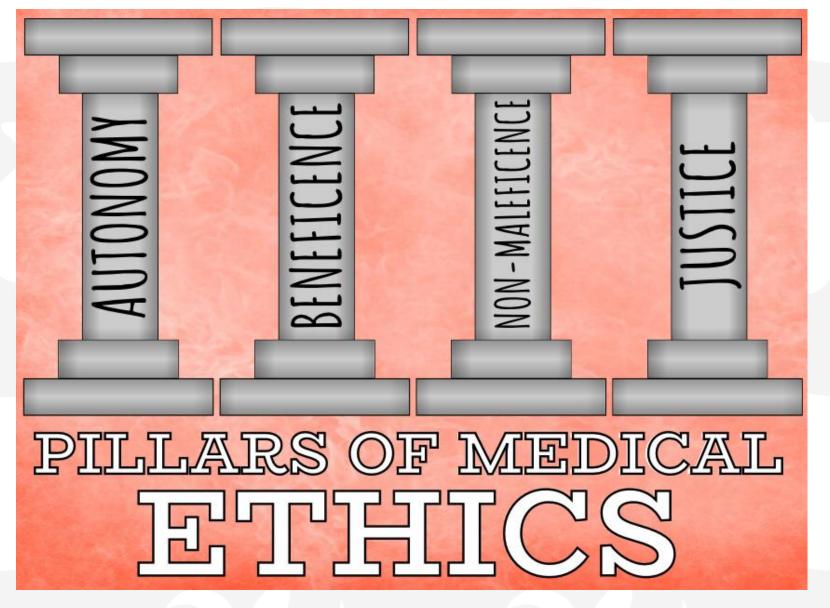


We suggest that most if not all cancers have acquired the same set

of functional capabilities during their development, albeit through

various mechanistic strategies.



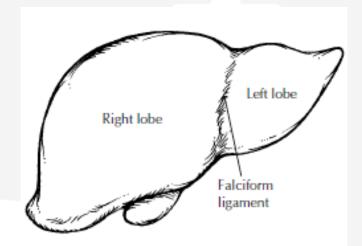






NHS Specimens

- Kept for 30 years
- Human Tissue Act
- Health Research Authority
- Research Ethics Committee (REC) approval
- Confidentiality Advisory Group (CAG) approval





Surgical Pathology Dissection: An Illustrated Guide, Second Edition. Ch 16, M Torbenson MD

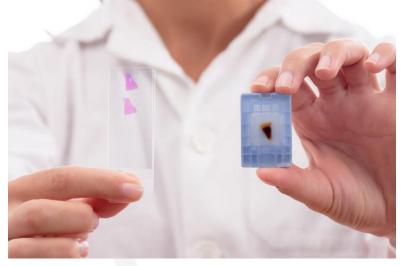
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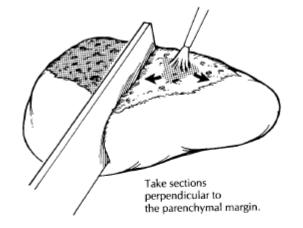
https://www.geneticistinc.com/blog/importance-of-ffpe-in-modern-research

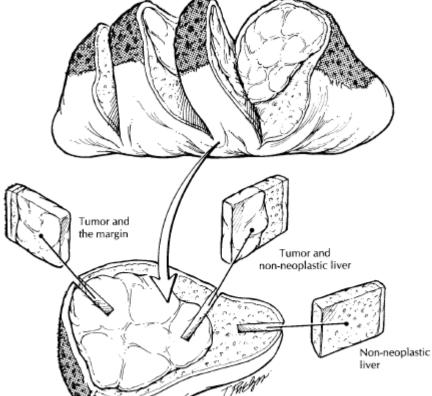
https://audubonbio.com/tissues/ffpe-tissue/







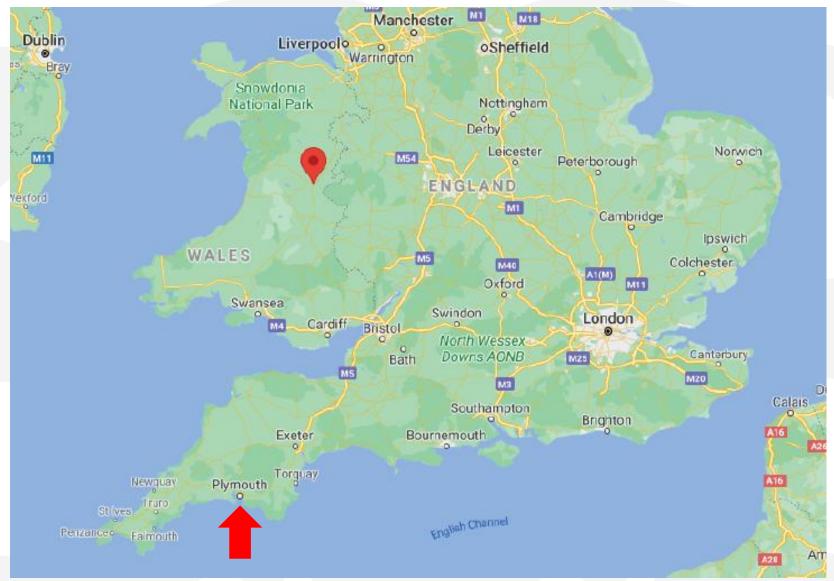
















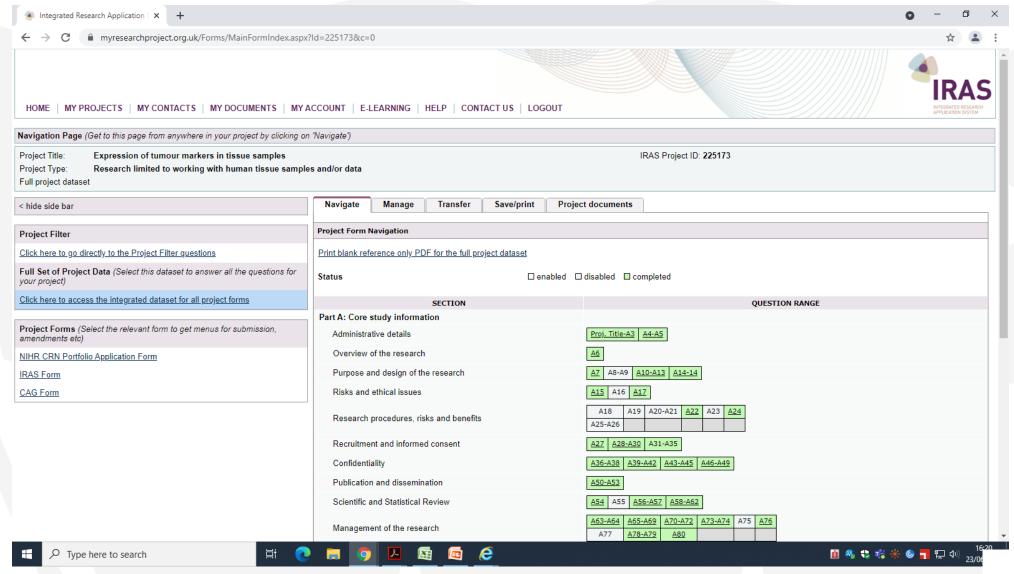
Integrated Research Application System

- Administration
 - CVs
 - Schedule of events
 - Statement of activities
 - Research Protocol
 - Sponsor (UoP)
 - Public liability
 - Professional indemnity
 - Insurance
 - Supporting document
 - British Liver Trust
 - Local patient interest group
- Confidentiality Advisory Group (CAG)
 - Caldicott Guardian input
- Research Ethics Committee (REC)
 - 17/NW/0457













Criteria

Inclusion

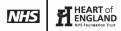
- 1. Tumour in the liver
- 2. Liver tissue at either UHP or KCH
- 3. >18 years old
- 4. Tissue surplus
- 5. Archived samples only
- 6. Research will not exhaust the archive

Exclusion

- 1. < 18 years old
- 2. Insufficient quantity of tissue
- 3. Explicit written
 evidence of the
 withdrawal of consent
 for research inclusion







Consent Form 1

Patient Agreement to Investigation or Treatment

Patient details (or pre-printed label)

Patient's surname/family name	I understand that you cannot give me a guarantee that a particular person will person will, however, have appropriate experience. I understand that I will have the opportunity to discuss the details of anaesthesia the procedure, unless the urgency of my situation prevents this (this only applies tranaesthesia). I understand that any procedure in addition to those described on this form will onecessary to save my life or to prevent serious harm to my health.	with an anaesthetist before o patients having general
Statement of health professional (to be filled in by health professional with appropriate	I have been told about additional procedures which may become necessary during here any procedures which I do not wish to be carried out without further discussi death:	ion, even if I become at risk of
I consent/do not consent to the removal of my tissue and/ I have explained th		is operation and
Research in connection with disorders and/or the fund Obtaining scientific or medical information about a live any other person (including a future person)	ctioning of the human body	ich may be relevant to
Patient's signature Any extra procedur Blood trans: Name (PRINT)		ier consent. Young people
Other procedures (please specify):	Name (FAINT)	
I have also discussed what the procedure is likely to involve, the benefits and risks of any available alternative treatments (including no treatment) and any particular concerns of this patient. The following leaflet/CD/DVD has been provided	Confirmation of consent (be completed by a health professional and the admitted for the procedure, if the patient has signed the form in advance) On behalf of the team treating the patient, I have discussed the treatment with the further questions or concerns. I have also confirmed with the patient that she/he hand wishes to go ahead.	ne patient and answered any
Name (PKINT)	Health Professional Signed	
Statement of interpreter (where appropriate)	Name (PRINT)	
I have interpreted the information above to the patient to the best of my ability and in a way in which I believe she/he can understand.	Patient	

Name (PRINT) .

Important notes: (tick if applicable)

See also advanced decision to refuse treatment/living will (e.g. Jehovah's witness form)

Patient has withdrawn consent, patient to sign and date here to confirm.

Statement of patient

any time, including after you have signed this form.

I agree to the procedure or course of treatment described on this form.

Please read this form carefully. If your treatment has been planned in advance, you should already have your own copy, which describes the benefits and risks of the proposed treatment. If not, you will be offered a copy

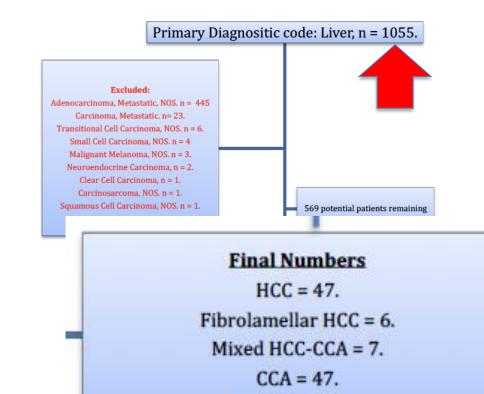
now. If you have any further questions, do ask - we are here to help. You have the right to change your mind at

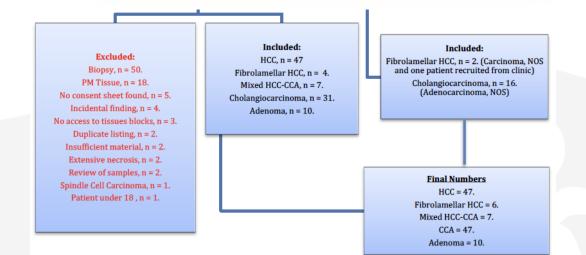


Signed .

Name (PRINT)..







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Figure 5. Eppendorf tubes containing tissue curls extracted from FFPE archived materials.





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Figure 6 Spin columns used for DNA extraction from FFPE tissues. The clear eluate in left image. Some tissue remains in the filter after processing in the right image.

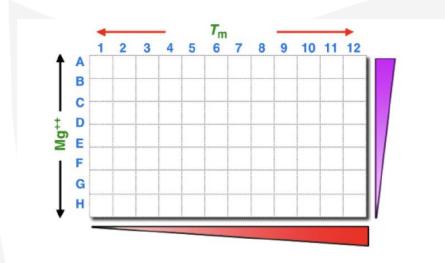


Figure 7. A grid pattern for altering two components of PCR simultaneously.

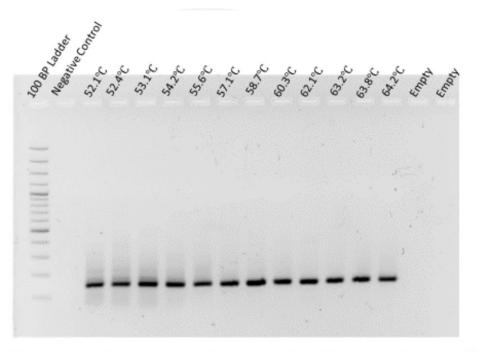
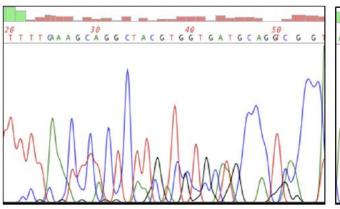


Figure 23. A 2% Agarose gel from a gradient PCR using the 'Surv F1R1' primers



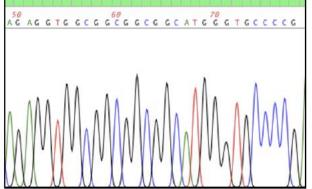




Figure 8. Chromatograms from MacVector with evidence of contamination (left) compared to an uncontaminated sequence. Note the low confidence (red bars, >1% error in the left image) compared to the high-confidence (green bars, <1% error in the right image) in base calls.



pTERT: Al	l Malignancies	N	Mutant	WT	P value
Gender	Female	36	1	35	0.001
	Male	65	18	47	
Outcome	Alive	41	6	35	0.444
	Dead	60	13	47	
Margin	R0	58	14	44	0.196
	R1	41	5	36	
Fibrosis	-	59	6	53	0.01
	+	41	13	28	
Cirrhosis	-	77	10	67	0.012
	+	23	9	14	
Tumour Stage	I & II	65	9	56	0.221
	III & IV	26	7	19	
Grade	Well Diff	11	2	9	0.401
	Mild/Mod Diff	66	14	52	
	Poorly Diff	17	1	16	
Vascular	-	42	7	35	0.794
Invasion	+	52	10	42	
Tumour Type	нсс	35	17	52	0.003
	FL-HCC	6	1	5	
	CCA	36	0	36	
	Mixed HCC-CCA	7	1	6	

Table 35. Clinical measures are compared between mutants and wild type Telomerase promoter sequences across all malignancies.

pTERT	: HCC only	N	Mutant	WT	P value
Gender	Female	12	0	12	0.005
	Male	40	17	23	
Outcome	Alive	23	4	19	0.043
	Dead	29	13	16	
Margin	R0	36	12	24	1.000
	R1	14	5	9	
Fibrosis	-	26	5	21	0.075
	+	26	12	14	
Cirrhosis	-	36	9	27	0.111
	+	16	8	8	
Tumour Stage	I & II	29	8	21	0.516
Stage	III & IV	17	7	10	
Grade	Well Diff	8	2	6	0.689
	Mild/Mod Diff	35	13	22	
	Poorly Diff	5	1	4	
Vascular	-	29	6	23	0.031
Invasion	+	19	10	9	

Table 37. Across traditional HCCs, clinical measures are compared between mutants and wild type Telomerase promoter sequences.

journal homepage: www.elsevier.com/locate/csbj

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Departmen of Allied He Lincoln, NE of Public I of Nebrask QuPath: The global impact of an open source digital pathology system



M.P. Humphries ^a, P. Maxwell ^a, M. Salto-Tellez ^{a,b,*}

^a Precision Medicine Centre of Excellence, The Patrick G Johnston Centre for Cancer Research, Queen's University, Belfast, UK ^b Integrated Pathology Programme, Division of Molecular Pathology, The Institute of Cancer Research, London, UK

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ABSTRACT

QuPath, originally created at the Centre for Cancer Research & Cell Biology at Queen's University Belfast as part of a research programme in digital pathology (DP) funded by Invest Northern Ireland and Cancer Research UK, is arguably the most wildly used image analysis software program in the world. On the back of the explosion of DP and a need to comprehensively visualise and analyse whole slides images (WSI), QuPath was developed to address the many needs associated with tissue based image analysis; these were several fold and, predominantly, translational in nature: from the requirement to visualise images containing billions of pixels from files several GBs in size, to the demand for high-throughput reproducible analysis, which the paradigm of routine visual pathological assessment continues to struggle to deliver. Resultantly, large-scale biomarker quantification must increasingly be augmented with DP. Here we highlight the impact of the open source Quantitative Pathology & Bioimage Analysis DP system since its inception, by discussing the scope of scientific research in which QuPath has been cited, as the system of choice for researchers.

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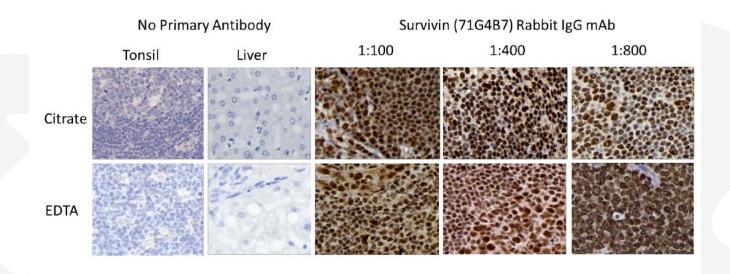
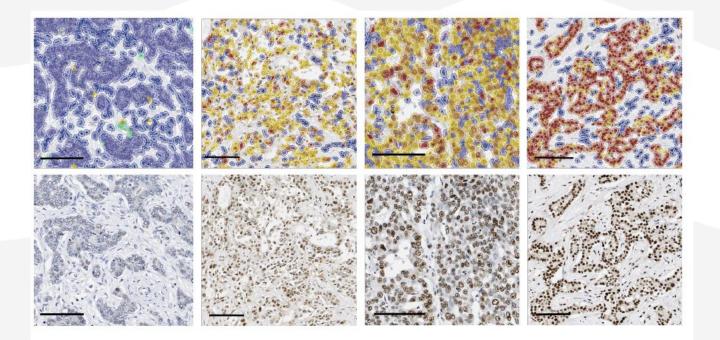


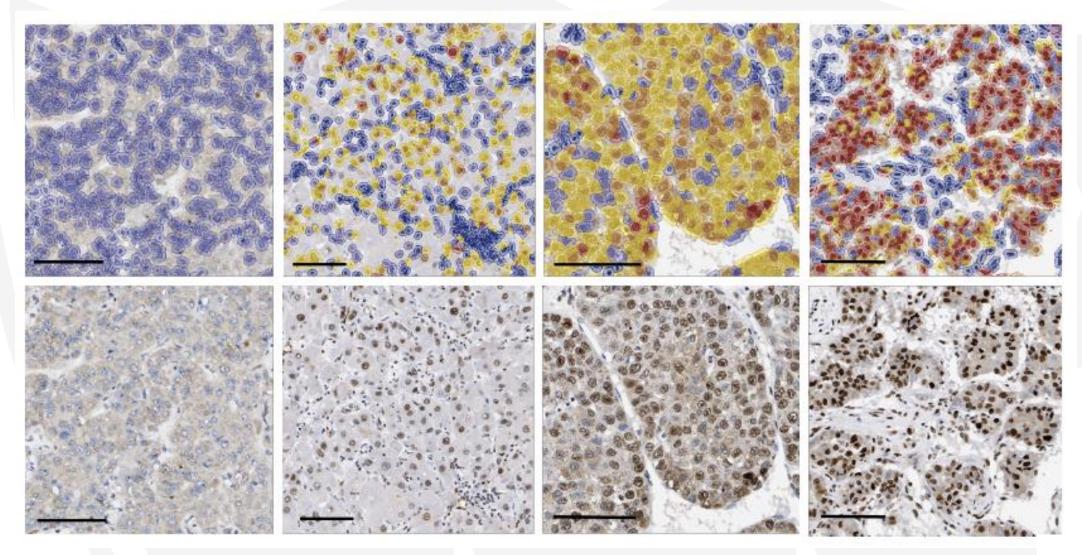
Figure 49. Optimisation steps for the Survivin primary antibody. Images shown at 200X magnification.





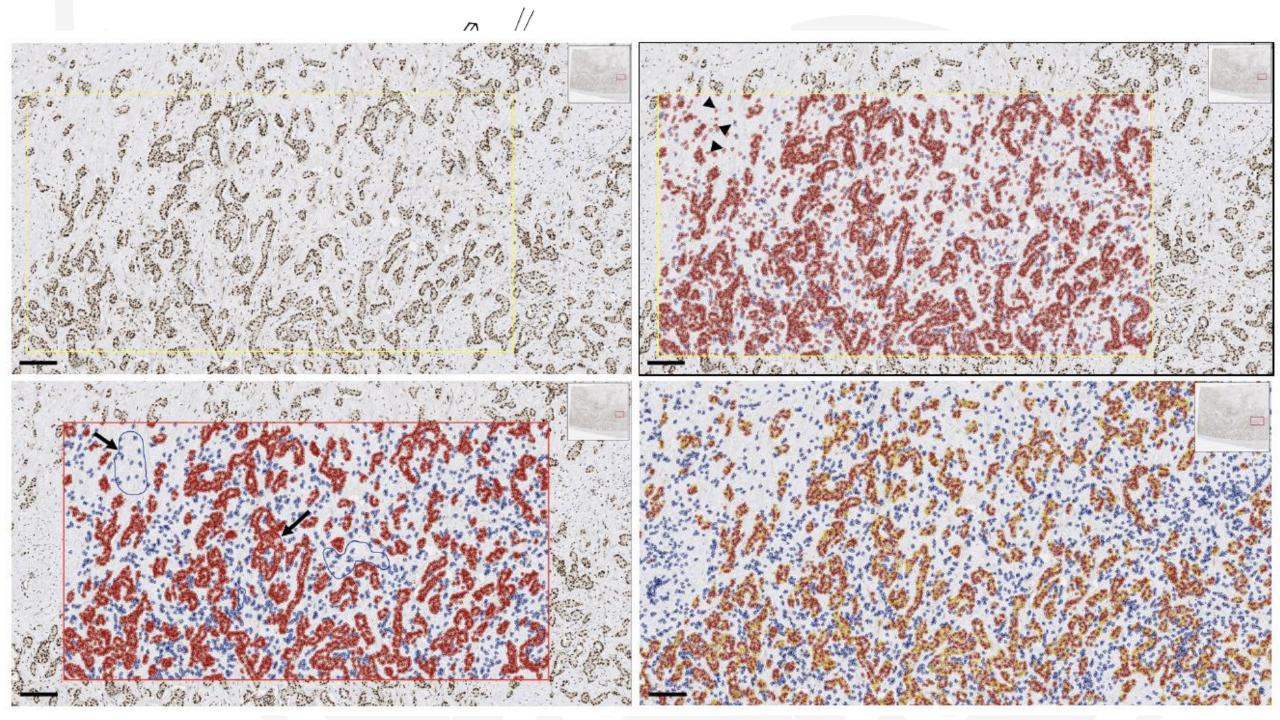
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Figure 51 CCA tumours with varying degrees of nuclear positivity for Survivin. Scale bar: 100 microns.

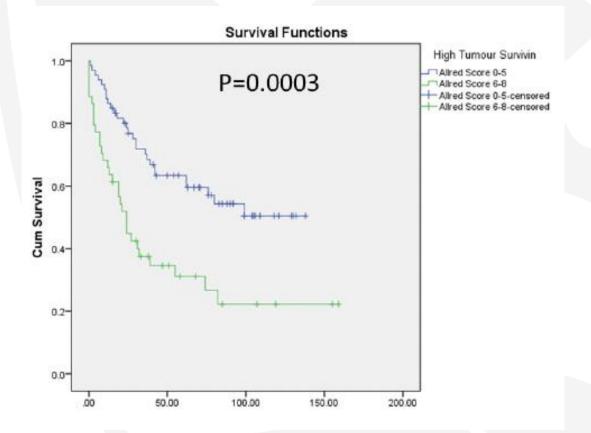




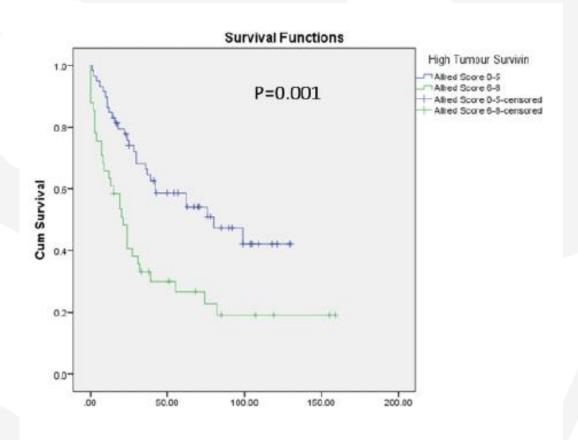




Neoplastic



Malignant







Malignancy - Survivin		N	Survivin ^{Low}	Survivin ^{High}	P-value
			% (n)	% (n)	
Outcome	Alive	41	75.5 (31)	24.4 (10)	0.007
	Dead	59	47.5 (28)	52.5 (31)	
Margin	R0	58	67.2 (39)	32.8 (19)	0.062
	R1	40	47.5 (19)	52.5 (21)	
Fibrosis	-	59	57.6 (34)	42.4 (25)	0.680
	+	40	62.5 (25)	37.5 (15)	
Cirrhosis	-	77	54.5 (42)	45.5 (35)	0.084
	+	22	77.3 (17)	22.7 (5)	
Tumour	I & II	65	67.7 (44)	32.3 (21)	0.029
Stage -	III & IV	25	40 (10)	60 (15)	
Tumour	I	30	83.3 (25)	16.7 (5)	0.007
Stage -	II	35	54.3 (19)	45.7 (16)	
-	TTT	10	2(0(7)	(2.0 (10)	

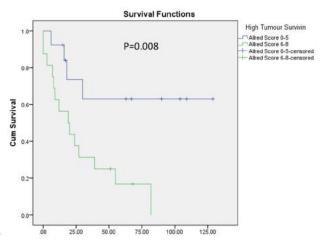




Figure 71. High Survivin expressing iCCA have a reduced cumulative survival.

			` '
GG	38	50 .0 (19)	50.0 (19)

Table 49. High tumour Survivin correlates with the clinic-pathological measures.

Independent-Samples Kruskal-Wallis Test

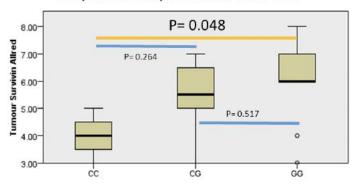


Figure 65. Survivin promoter variants differentially express Survivin protein In CCAs.

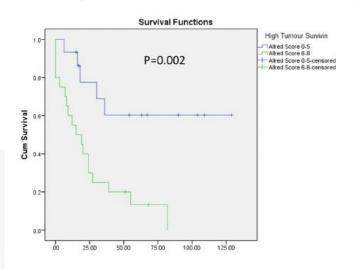


Figure 69. Cumulative survival differences based on Survivin expression in CCA.



Blood Work

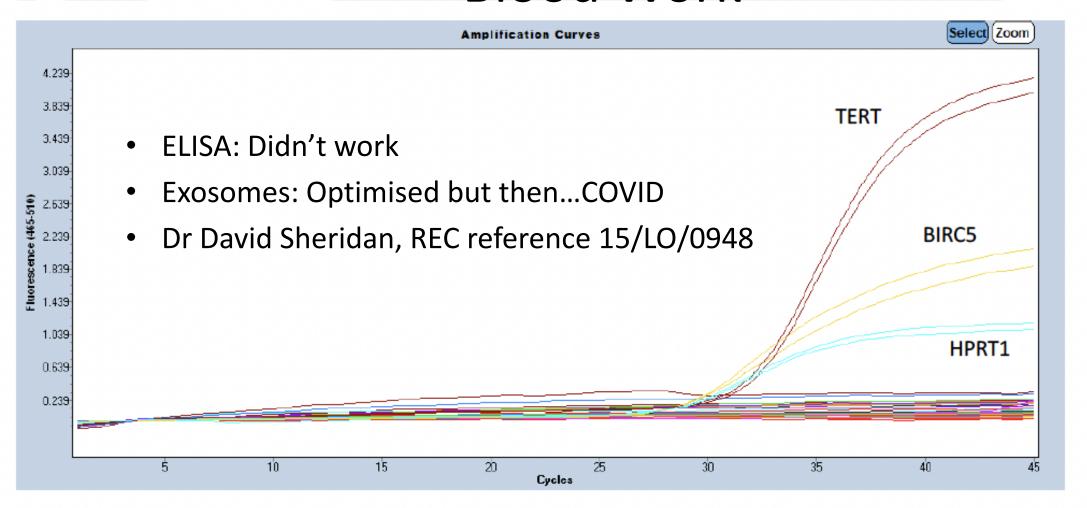


Figure 92. Quantitative PCR data from amplified Exosomal RNA. The calibrators (TERT and BIRC5) demonstrate the presence of this RNA from the Huh7.5 cell line and highlights that the reaction works. The internal control, HPRT1, is also detected.

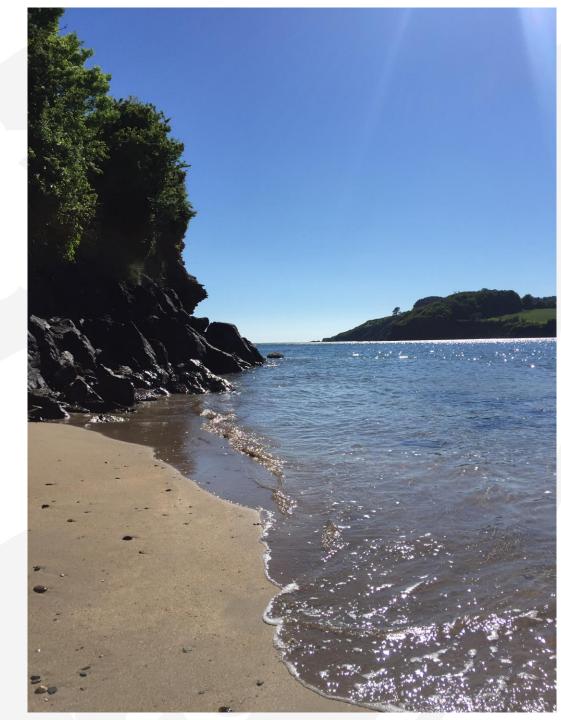


Conclusions

- It WORKED!
- Funding...
- Supervision...
- Time consuming, but worthwhile.
- Unimaginably large potential research resource







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Prof Matthew Cramp
Dr Michael Jarvis

Dr Jemimah Denson, Alec McLean, UHP Pathology.

The Hepatology Research Group, Plymouth, especially Dr David Sheridan.

Dr Jemma Dunn Dr Aisling Murphy

Pam Baxter



