

# **An Investigation of Mutant p53 Function**

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B. Science

B. Health Science (Hons)

Thesis submitted for the Degree of Doctor of Philosophy

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July 2011

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## ABSTRACT

The *TP53* tumour suppressor gene is mutated in approximately 50% of all human cancers. The majority of these mutations are missense mutations resulting in the expression of a mutated form of the full-length p53 protein. This mutant protein exhibits a loss of tumour suppressive activity, dominant-negative activity to inactivate functional p53 and gain-of-function properties to drive tumour progression and metastasis. Investigation into mutant p53-mediated oncogenic pathways and the mechanisms through which they are controlled plays an integral role in identifying new therapeutic targets for a range of mutant p53-expressing tumours.

To model the initial events that occur in cancer following sporadic p53 mutation, an isogenic panel of cell lines was established in the p53 null, H1299 lung cancer cell line, expressing wild-type or various p53 hotspot mutants under the control of an inducible promoter. These cell lines were harnessed to investigate a range of wild-type and mutant p53 functions. The induced wild-type p53 protein is demonstrated to be transcriptionally and biologically active, and its function can be further mediated by DNA damaging agents or expression of regulatory proteins. Conversely, induced mutant p53 exhibits a loss of the majority of the normal wild-type transcriptional activity while mediating gain-of-function, oncogenic phenotypes in H1299 cells. This system is demonstrated to provide an important platform with which to investigate both wild-type and mutant p53 function.

Mutant p53 is reported to function as an aberrant transcription factor, re-programming the cellular transcriptome to enhance oncogenic pathways. The mechanisms

underlying this were specifically examined through expression microarray analysis, which identified a number of mutant p53-regulated targets. Surprisingly, these targets were predominately also direct targets of wild-type p53. A novel mechanism for mutant p53 activity is subsequently suggested, whereby mutant p53 is recruited to the DNA through its interaction with p63.

A key function of mutant p53 is its ability to drive tumourigenesis through the initiation of a range of oncogenic pathways. Through utilising the inducible system, mutant p53 is demonstrated to influence mitotic pathways, resulting in multinucleation, and enhance the invasive and migratory properties of cancer cells. Importantly, an endogenous protein, ANKRD11, is identified with the capacity to suppress the oncogenic properties of mutant p53 and provide a potential target for the development of new cancer therapeutics.

The role of mutant p53 in driving the invasive and metastatic potential of breast cancer cells was further explored and a relationship between mutant p53 and a micro-RNA (miR-155) established. Mutant p53 expression is shown to correlate with miR-155 expression, with miR-155 target genes involved in invasive pathways. ZNF652 is specifically identified as a target of miR-155 and loss of ZNF-652 is correlated with increased invasion and poor prognosis in breast cancer.

Collectively, these studies identify key mechanisms through which mutant p53 functions to enhance tumourigenesis and importantly identify novel targets, ANKRD11, miR-155 and ZNF652, for the development of cancer therapies.



## DECLARATION

I, Jacqueline Elise Noll, certify that this work contains no material which has been accepted for the award of any other degree or diploma in any university or other tertiary institution and, to the best of my knowledge and belief, contains no material previously published or written by another person, except where due reference has been made in the text.

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## LIST OF PUBLICATIONS

**Noll JE**, Jeffery J, Al-Ejeh F, Kumar R, Khanna KK, Callen DF and Neilsen PM. (2011) Mutant p53 drives multinucleation and invasion through a process that is suppressed by ANKRD11. *Oncogene*. Submitted. Second Revision.

Muller PAJ, Trinidad AG, Timpson P, Morton J, Nixon C, Karim S, Caswell P, **Noll JE**, Coffill CR, Lane DP, Sansom O, Neilsen PM, Norman JC and Vousden KH. (2011) Mutant p53 induces c-Met signalling to drive cell scattering and invasion by inhibiting TAp63 and Dicer. *Nature Cell Biology*. Submitted, Second Revision

Chee JLY, Saidin S, Lane DP, Leong SM, Phua YT, **Noll JE**, Neilsen PM, Gabra H and Lim TM. (2011) Wild type and mutant p53 mediate cisplatin resistance through interaction and inhibition of caspase-9. *Carcinogenesis*. Submitted

Neilsen PM\*, **Noll JE\***, Tay BS, Bracken C, Schulz R, Lim S, Gregory P, Kumar R, Goodall G and Callen DF. (2011) Mutant p53 drives invasion in breast tumors through a pathway involving miR-155 and ZNF652. *Text in Manuscript*

\* These authors contributed equally to this work

## **ABBREVIATIONS**

**ANK** – Ankyrin

**ANKRD11** – Ankyrin repeat domain 11

**BCA** – Bicinchoninic acid

**CBP** – CREB binding protein

**CIP** – Calf intestinal phosphatase

**cDNA** – Complimentary DNA

**ChIP** – Chromatin immunoprecipitation

**DBD** – DNA binding domain

**DMEM** – Dulbecco's modified eagle medium

**DMSO** – Dimethyl sulfoxide

**DN** – Dominant negative

**DNA** – Deoxyribonucleic acid

**ECL** – Enhanced Chemiluminescence

**EDTA** – Ethylenediaminetetraacetic acid

**EI** – Ecdysone inducible

**EMT** – Epithelial to mesenchymal transition

**FACS** – Fluorescence-activated cell sorting

**FBS** – Fetal bovine serum

**GFP** – Green fluorescent protein

**GOF** – Gain of function

**HRP** – Horseradish peroxidase

**LOF** – Loss of function

**MDM2** – Murine double minute 2

**mRNA** – Messenger RNA

**ORF** – Open reading frame

**PBS** – Phosphate buffered saline

**PCR** – Polymerase chain reaction

**PonA** – Ponasterone A

**P/CAF** – p300/CBP associated factor

**RE** – Response element

**RNA** – Ribonucleic acid

**RT** – Room temperature

**RT-PCR** – Reverse transcription PCR

**SDS** – Sodium dodecyl sulfate

**SDS-PAGE** – Sodium dodecyl sulfate polyacrylamide gel electrophoresis

**shRNA** – Short hairpin RNA

**SNP** – Single nucleotide polymorphism

**ssDNA** – Salmon sperm DNA

**SV40** – Simian virus 40

**UTR** – Untranslated region

## **ACKNOWLEDGEMENTS**

To my supervisors – Prof. David Callen, Dr. Paul Neilsen and Dr. Raman Sharma. Thank you for providing me with such a wonderful opportunity and for guiding me along the way: to study, to learn and to develop my skills and understanding.

To the members of the Cancer Therapeutics Laboratory/Breast Cancer Genetics Group – Thank you for providing such a fun and supportive atmosphere to work in. I would particularly like to thank Renee; no matter where I was at you were there – to lift my spirits or to celebrate with me. Your support and friendship over the past few years has been immensely important to me and I will never forget it.

To my friends and family – Thank you for always showing an interest in what I was doing, even when you didn't understand it. And most importantly thank you for believing in me and my abilities, even when I doubted myself.

To my amazing husband, Ben – Thank you for your constant love and support. I know I can achieve anything while I have you by my side. Your continuing and never-ending positive outlook on life helped me focus my attention to where it mattered most. Whether I needed a shoulder to cry on or someone to celebrate with you were there, and knowing that made all the difference. I couldn't have made it through without you. Love you, babe.

NOTE:

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