

# Pathophysiological impact of targeting the ROS-p53 axis

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Cover illustration: “The Great White Void”

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*To boldly go where no one has gone before*



# ABSTRACT

The goal of this PhD thesis was to define the importance of the interplay between reactive oxygen species (ROS) and their activation of the tumor suppressor p53 in development and disease. We addressed this question using molecular biology and biochemical techniques together with mouse genetics and bioinformatics.

We have made two important discoveries:

First, we show that antioxidant supplementation accelerates lung cancer progression in mice and the growth of human lung cancer cell lines. By reducing the levels of ROS and DNA damage, antioxidants deactivate the p53 protein and help cancer cells to evade growth arrest.

Second, we show that the transcription factor zinc finger protein 148 (Zfp148) is a potent suppressor of p53 activation under oxidative conditions. During lung development, suppression of p53 prevents growth arrest of pulmonary cells and permits prenatal lung maturation. However, in the *Apc<sup>Min/+</sup>* model of colorectal cancer and in the *ApoE<sup>-/-</sup>* model of atherosclerosis, suppression of p53 promotes tumor development and atherosclerosis, respectively. Thus Zfp148 suppression of p53 plays important roles in both physiological and pathological contexts.

We conclude that:

- 1) Antioxidant supplementation may stimulate the growth and progression of undiagnosed lung tumors and should be used with caution. The risk of developing lung cancer in patients with chronic obstructive pulmonary disease (COPD) who take the antioxidant acetylcysteine to break down mucus should be carefully evaluated.
- 2) Therapeutic targeting of Zfp148 may have beneficial effects in cancer and atherosclerosis by increasing p53 activity.

**Keywords:** ROS, p53, Antioxidants, Zfp148, cancer and atherosclerosis

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# SAMMANFATTNING PÅ SVENSKA

Målet med denna doktors avhandling har varit att definiera betydelsen av reaktiva syreföreningar (ROS) och deras samverkan och aktivering av tumörsuppressorer p53 i hälsa och ohälsa. Vi adresserar denna fråga med hjälp av molekylärbioologiska och biokemiska tekniker tillsammans med mus genetik och bioinformatik .

Vi har gjort två viktiga upptäckter :

- 1) Vi visar att antioxidant tillskott accelererar lungcancer progression hos möss och tillväxten av humana lungcancer cellinjer . Genom att sänka nivåerna av ROS och DNA-skada , inaktiverar antioxidanter p53-protein och hjälper cancerceller att kringgå tillväxtestopp.
- 2) Vi visar att transkriptionsfaktorn zink finger protein 148 ( Zfp148 ) är en potent suppressor av p53 aktivering under oxidativa förhållanden . Vi visar även att denna suppressor aktivitet är betydelsefull för hämning av kolorektal cancer och ateroskleros i mus modeller. För mycket av denna suppressor aktivitet leder dock till en lung utvecklings defekt följt av respiratorisk distress i samband med födsel av möss. Således har vi visat att Zfp148's hämning av p53 spelar en viktig roll i både fysiologiska och patologiska sammanhang .

Vi drar slutsatserna att :

- 1 ) Antioxidant tillskott stimulerar tillväxt och utveckling av framför allt tidiga icke diagnostiserade tumörer. Därför bör antioxidant tillskott användas med försiktighet . Risken att utveckla utvecklings lungcancer hos patienter med kronisk obstruktiv lungsjukdom ( KOL ) som tar antioxidanter acetylcystein att bryta ner slem bör utvärderas vården noggrant .
- 2 ) Terapeutisk hämning av Zfp148 kan ha gynnsamma effekter på utveckling av cancer och åderförkalkning genom att öka p53 –aktivitet.

## LIST OF PAPERS

This thesis is based on the following studies, referred to in the text by their Roman numerals.

- I. Sayin VI, Nilton A, Ibrahim MX, Agren P, Larsson E, Petit MM, Hulten LM, Stahlman M, Johansson BR, Bergo MO and Lindahl P.  
**Zfp148 deficiency causes lung maturation defects and lethality in newborn mice that are rescued by deletion of p53 or antioxidant treatment**  
*PLoS One* **2013** 8(2):e55720
  
- II. Nilton A\*, Sayin VI\*, Bondjers C, Agren P, Bergo MO and Lindahl P  
\*Equal Contribution  
**Zfp148 deficiency reduces tumor formation in APCMin/+ mice in a p53-dependent manner**  
*In Manuscript*
  
- III. Sayin VI, Ibrahim MX, Larsson E, Nilsson JA, Lindahl P and Bergo MO.  
**Antioxidants accelerate lung cancer progression in mice**  
*Science Translational Medicine* **2014** 6(221):221ra15
  
- IV. Sayin VI, Khan OM, Pehlivanoglu LE, Staffas A, Ibrahim MX, Asplund A, Agren P, Nilton A, Bergström G, Bergo MO, Borén J and Lindahl P.  
**Loss of one copy of Zfp148 reduces lesional macrophage proliferation and atherosclerosis in mice by activating p53**  
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## ABBREVIATIONS

ApoE	Apolipoprotein E
APC	Adenomatosis polyposis coli
CMR	Chylomicron remnant
COPD	Chronic obstructive pulmonary disease
CRE	Cyclic recombinase
FAP	Familial adenomatous polyposis
Floxed	Flanked by loxP
FAP	Familial adenomatous polyposis
Floxed	Flanked by loxP
GSH	Glutathione
GT	Gene trap
LDLr	Low-density lipoprotein receptor
LSL	Lox stop Lox
MEF	Mouse embryonic fibroblast
Min	Multiple intestinal neoplasia
NAC	N-acetylcystein
Nrf2	Nuclear factor like 2 also known as NFE2L2
ROS	Reactive oxygen species

VLDL	Very low-density lipoprotein
Zfp148	Zinc finger protein 148
WT	Wild-type
mRNA	Messenger Ribonucleic acid
p53	Protein coded by Trp53 in mice and Tp53 in humans
P1	Postnatal day 1
TUNEL	Terminal deoxynucleotidyl transferase dUTP nickend labeling

# 1 INTRODUCTION

In this thesis I will explore the impact of targeting reactive oxygen species (ROS) and their interplay and activation of p53 (hereafter denoted the ROS-p53 axis) in cancer, vascular disease and development. To achieve this we applied two major strategies. Firstly, we knocked out Zinc finger protein 148 (Zfp148) in mice, a transcription factor that interacts with p53. Secondly, we treated cancer prone mice with ROS scavengers, more widely known as antioxidants. In this section, I will introduce and provide a background to the major concepts of this thesis including ROS, antioxidants, P53 and Zfp148 as well as highlight knowledge gaps coupled to these key factors.

## 1.1 Reactive Oxygen Species

Our present ecosystem is the product of a series of dramatic events. One such event, termed the great oxidation, started around 2.4 billions years ago as cyanobacteria begun to harness the power of photosynthesis. As a byproduct of photosynthesis, our atmosphere became rich in free oxygen (1, 2). This process nearly wiped out all life on earth, since it was previously obligate anaerobic. As a consequence, the great oxidation sparked the evolutionary adaptation of early life to the presence of oxygen. Whether or not high levels of oxygen were required for higher animals to evolve remains debated. Indeed, recent discoveries of multicellular organisms living their whole lives in an oxygen free environment suggest that this might not be the case (3, 4). Nevertheless, the majority of life forms on earth including all mammals are at present totally dependent on oxygen, a highly toxic agent. Once oxygen was harnessed by respiration in the final step of the electron transport chain, aerobic life became able to generate exponentially more energy molecules per nutrient molecule. However, this boost in energy production came with a

price in the form of byproducts of oxygen metabolism called reactive oxygen species (ROS) which are toxic and cause damage to macromolecules including proteins, lipids and DNA (5-8). In order to sustain long life, an adaptive response was evolved. Nuclear factor like 2 (Nrf2 or NFE2L2), the master transcriptional regulator of the endogenous antioxidant response, maintains homeostasis and redox balance in response to oxidative stress (increased ROS levels) within each cell (9). Naturally, an adaptive antioxidant defense system, which is activated by oxidative stress, will always be one step behind. Hence it is not surprising that ROS are implicated not only in cancer and cardiovascular disease, but also in physiological aging (10).

## 1.2 Antioxidants

Antioxidants are electron donating molecules that neutralize ROS and other free radicals that may otherwise cause oxidative damage to DNA, proteins and lipids and promote cancer and cardiovascular disease (11, 12). Antioxidants are produced endogenously in cells to balance ROS levels. During oxidative stress, Nrf2 translocates to the nucleus and induces the expression of glutathione (GSH) and other antioxidants (13). Essential antioxidants, including vitamins, carotenes and minerals are found naturally in food. Sufficient intake of antioxidants from food is important, and is underscored by the dramatic symptoms of vitamin deficiency (14-20). Additionally, gene targeting experiments of endogenous antioxidants in mice have led to a range of disease phenotypes including cardiovascular disease and increased susceptibility to cancer development (21-23).

Consequently, popular wisdom—supported by numerous cellular and preclinical studies—holds that antioxidant supplements protect against cancer (24-26) and cardiovascular disease (27-31). However, large randomized

clinical trials have produced inconsistent results related to the effect of supplementation with antioxidants on cancer (32-34) and cardiovascular disease (35-38). Importantly, several studies show that antioxidants may even increase the risk of developing cancer (39-42) and all-cause mortality (43). Nevertheless, there is a big discordance between clinical outcomes on antioxidant supplementation and the use of antioxidant supplements in the population. Indeed, over 10% of random populations take antioxidant supplements on a daily basis (44). One explanation for the misuse of antioxidants is the lack of a mechanistic understanding of how antioxidants may accelerate cancer progression.

## 1.3 p53

Endogenous antioxidants are the first line of defense against cancer; however, a second more crucial line of defense is kept by tumor suppressors. The tumor suppressor p53 is the most frequent mutated gene in cancer (45, 46) and may be the most widely studied gene overall with over 73 000 publications on the topic. P53 induces DNA repair, senescence, or apoptosis in response to a wide range of stressors including oxidative stress (46, 47). Whether p53 activation leads to DNA repair, senescence or apoptosis depends on a complex interplay between co-factors that control p53 stability, activation and transcriptional outcome (48). The E3 ubiquitin ligase Mdm2 negatively regulates p53 by targeting it for degradation. Mdm2 null mice die during early embryogenesis in a p53-dependent manner, which demonstrates that physiological stress alone triggers p53-dependent senescence or apoptosis in the absence of appropriate control mechanisms (49, 50). In a similar fashion, mice lacking Mdm4, a structural homologue of Mdm2 that lacks ubiquitination activity but binds to the N-terminus of p53 and directly represses transcriptional activity, die during embryogenesis in a p53-dependent manner (51, 52). The interaction between p53 and Mdm2 has been conserved across 2.4 billion years of evolution (53, 54), and is traced back to the beginning of multicellular life. Interestingly, the evolutionary timing of p53 and MDM2 interaction happens to correlate with the early phases of the great oxidation, hinting on a possible evolutionary link.

ROS and p53 have a complex and context dependent relationship, dating back to the great oxidation (55). It is well established that increased levels of ROS (oxidative stress) activates p53, and that p53 in turn can increase levels of ROS by enhancing the transcription of pro-apoptotic genes (56, 57). However, p53 can also reduce levels of ROS by transcriptional induction of antioxidant genes and this function may contribute to the tumor suppressor

properties of p53 (25). Moreover, p53 itself undergoes redox regulation due to redox sensitive cysteine residues (58). There are two clusters of cysteine in the DNA binding domain of p53 which are essential to the specific binding of p53 to target genes (59). Furthermore, exposing p53 to oxidants causes Cysteines-124,141 and 182 on p53 to form disulfide bonds with GSH, effectively changing the DNA binding activity of p53 (60, 61). Notably, the change in DNA binding of p53 after oxidation can be reversed by antioxidants (60, 61). Thus, there is a complex crosstalk between ROS and p53.

When cells are exposed to stressors like increased ROS, the most important function of p53 is to regulate expression of downstream target genes. In response to DNA damage, p53 represses the expression of cell cycle-related genes involved in the G2/M phase transition (62). In tumors lacking functional p53, these genes are expressed at high levels and correlate with increased malignancy and poor clinical outcome (63). Thus, repression of cell cycle-related genes is a crucial tumor suppressor function of p53. The mechanisms underlying p53 regulation of cell cycle genes are not fully understood but involve both direct binding of p53 to regulatory elements and indirect interactions with other transcription factors (63).

## 1.4 Zinc Finger Protein 148

The transcription factor Zinc finger protein 148 (Zfp148) (also known as: ZBP-89, BFCOL, BERF1) contains four krüppel type zinc finger domains and binds to GC-rich DNA sequences (64-67). The protein is predominantly localized in the nucleus and is expressed ubiquitously in tissues of adult mice (65, 67). Zfp148 harbours putative repressor and transactivation domains in the N- and C-terminal regions respectively (66, 67), and is capable of recruiting co-activators and co-repressors to promoter regions (68-71).

Zfp148 has been linked to a number of target genes, but there are no obvious functional connections between them (65-69, 71-78). However, four arguments suggest that Zfp148 plays a role in cell cycle control. First, Zfp148 interacts physically with p53 (79). Second, overexpressing Zfp148 in cancer cell lines increases p53 levels in the nucleus and induces growth arrest or apoptosis (76, 79, 80). Third, Ataxia telangiectasia mutated (ATM), one of the two central regulators of the DNA damage response (81), phosphorylates Zfp148 at the zinc finger domains (82). Moreover, ATM together with Zfp148 and p300 binds to the promoter of the cyclin dependent kinase inhibitor 1a (p21) (69). And finally, silencing of Zfp148 in the NCI-H460 cell line induces senescence through induction of Ink4a (71). Notably, there is discordance between the outcomes in the cell cycle related studies; however, collectively these studies implicate Zfp148 in cell cycle control.

The physical interaction between Zfp148 and p53 suggests a potential role for Zfp148 in tumor suppression beyond cell cycle control (79). Binding studies have shown that the DNA binding zinc finger domains of Zfp148 are required for the binding to p53 (79). Another study shows that mutations in the p53 transactivation domains are dispensable for the interaction between Zfp148 and p53 (83). However, the same study show that hotspot mutations



spanning amino acids 175-281 in the DNA binding domain of p53 abolish the binding between p53 and Zfp148 (83). Furthermore, cells with mutant forms of p53 evade the apoptotic function of Zfp148 (80, 84). Nevertheless, the physiological relevance of the physical interaction between p53 and Zfp148 remains a knowledge gap.

The physiological function of Zfp148 remains unclear. Three gene targeting experiments on Zfp148 in mice have produced inconsistent results (85-87). In the first, Takeuchi et al. showed that Zfp148 heterozygote males suffer from sertoli cell-only syndrome, lacking germline cells, making propagation of the strain impossible. In the second study, Zfp148 deficient mice died during embryogenesis with neural tube defects and anaemia. In the final study, Zfp148 exon 4 knockout mice were generated that showed partial postnatal lethality and dextran sulphate induced colitis. The inconsistency between the gene targeting experiments could be a result of different targeting strategies. Nevertheless, the conflicting gene targeting experiments shows that the physiological role of Zfp148 remains unclear.

*"Vision is the art of seeing things invisible." Jonathan Swift*

## 2 AIM

The initial aim of this thesis was to define the impact of *Zfp148* deficiency on development, health and disease.

The specific aims of the four papers included in this thesis were:

**I. *Zfp148* Deficiency Causes Lung Maturation Defects and Lethality in Newborn Mice That Are Rescued by Deletion of p53 or Antioxidant Treatment**

The aim of the first study was to generate, validate and phenotype *Zfp148* deficient mice.

**II. *Zfp148* deficiency reduces tumor formation in *APC*<sup>Min/+</sup> mice in a p53-dependent manner**

The aim of the second study was to breed the *Zfp148* deficient mice on to the *APC*<sup>Min/+</sup> model of intestinal cancer to test whether *Zfp148* plays a role in colorectal cancer.

**III. Antioxidants accelerate lung cancer progression in mice**

The aim of the third study was to define the impact and mechanism of antioxidant treatment on lung cancer progression.

**IV. Loss of one copy of *Zfp148* reduces lesional macrophage proliferation and atherosclerosis in mice by activating p53**

The aim of the final paper of this thesis was to define the impact of *Zfp148* deficiency on the progression of atherosclerosis.

*"Measure what is measurable, and make measurable what is not so." -  
Galileo*

## **3 EXPERIMENTAL STRATEGIES AND CONSIDERATIONS**

In this section I provide a more general description of the genetic strategies and mouse models that are central to the work behind this thesis, why we chose to work with them and discuss their limitations.

Detailed descriptions of methods used in this thesis can be found in the materials and methods section of each enclosed paper.

### **3.1 The mouse as a model organism**

Since experimental research on humans is limited to highly controlled and regulated clinical trials, we depend on model organisms, like the mouse, for gaining insights into mechanisms of health and disease. The most important advantage of using mice as model organisms is their similarity to humans in anatomy, physiology, and genetics. In fact, more than 95% of the mouse genome is similar to our own, making mouse genetic research applicable to human biology and disease (88). Mouse models are cost-effective tools that speed up research and are crucial to the development and validation of targeted drug therapies. Mice are born in large litters, have a small size and a short lifespan, keeping the space, time, and costs required to perform research at reasonable levels.

### **3.1.1 Transgenic mice**

More than 30 years ago the first transgenic mouse was generated. Foreign DNA was introduced into the mouse genome, resulting in expression of the transgene (89-92). Shortly thereafter, through the use of homologous recombination, gain and loss of function experiments on mice became a golden standard for mapping gene function (93-98). The generation of knock-out and knock-in mice is still considered the highest level of evidence when validating gene function.

### **3.1.2 Modeling human disease in mice**

Even though there are striking similarities between men and mice, normal wild-type (WT) mice seldom face our most common diseases, including cancer and cardiovascular disease. Only after generations of back-crossing of laboratory mice into disease prone backgrounds have we been able to generate strains of mice where human disease could be studied (99). The introduction of transgenic techniques revolutionized the field of genetics and opened up for the generation of countless new mouse models of disease (100, 101).

Importantly, a lot of human hereditary or spontaneous genetic disorders manifest in a similar fashion in mice when the underlying genetic events are known and are introduced in the mouse genome (102-105). However, there are limitations. Some conditions are poorly mimicked or overly simplified in mouse models. (105, 106)

## 3.2 Cre-loxP System

Several of the mouse models used in this thesis utilizes the Cre-loxP System for conditional activation or conditional deletion of target genes, hence an introduction is warranted.

Cyclic recombinase (*Cre*) is an enzyme isolated from the bacteriophage P1 that recognizes and binds to specific *locus of x crossing over* (loxP) sites (107-109). LoxP sites consist of 34 base pair long DNA elements that include two 13 base pair inverted repeats flanking an 8 base pair spacer region. *Cre* cleaves DNA sequences that are flanked by two loxP sites (floxed) oriented in the same direction. The cleaved DNA is excised into a circular loop of DNA(110).

The transgenic introduction of loxP sites into mammalian cells (111) enabled the conditional targeting of genes in a time and cell type dependent manner. The deletion of floxed elements is controlled by the expression of *CRE* recombinase, which is in turn dictated by the experimental setup. The expression of *CRE* can be controlled in two ways. One way is with the help of exogenous insertion of vectors expressing *CRE*, like plasmids or virus particles. Another way is with transgenic insertion of *CRE* behind tissue specific promoters.

In this thesis, expression of *CRE* is (In paper III) *CRE* expression is controlled by adenoviral vectors that are delivered to the lung epithelia of mice through inhalation of a calcium phosphatase precipitate solution (112).

The Cre-loxP system is a powerful tool for conditional gene modifications in animals and has achieved widespread acceptance over the last decade.

However, there are four limitations to consider when interpreting the results from Cre-loxP experiments.

First, endogenous expression of *CRE* is dependent on the promoter region of the inserted transgene which dictates the tissue specificity and temporal control of the *CRE* expression (113). Even though tissue specific promoters are supposed to be fairly specific, depending on the nature of the promoter, it can have leaky expression or just naturally be expressed elsewhere in a developmental stage or context dependent fashion (114-118)

Second, loxP sites have no functionality in mammalian genomes and should therefore not be present. However; a recent genome wide study identified frequent cryptic loxP sites that can promote illegitimate DNA recombination and cause damage in cells and tissues that express *CRE* (119). For example, this is a problem in heart tissue where *CRE* expression alone causes fibrosis (120) and dilated cardiomyopathy (121).

The third potential problem with the Cre-loxP approach is partial recombination, i.e. that some floxed alleles are left undeleted. This becomes evident when more than one floxed allele is targeted for simultaneous deletion, especially if the *CRE* expression is transient (122, 123).

And finally, on rare occasions, the targeted gene may still be expressed from the excised circular DNA fragment. In high proliferative tissues like intestinal epithelium and tumor cells the circular DNA fragment would be diluted over time. However, in non-dividing cells like neurons the presence of this extra chromosomal DNA fragment has been a problem. (124)



### 3.3 Zfp148 deficient mice

The generation and validation of Zfp148 deficient mouse is well described (paper I) and will be further described in the Results section. The Zfp148 deficient mice were generated with the help of gene-trapping technique. The gene-trap (gt) vector is designed as a false exon with a splice acceptor and a transcriptional stop. Integrating the gt vector between exon 4 and 5 in *Zfp148* renders the major part of the translated region (exon 4 to 9) un-transcribed (exon 5 to 9). Instead a fusion protein containing exon 4 and the gt vector is obtained containing a reporter element (bgal) and a neomycin resistance element. Of note, the Zfp148 gt allele is not a null allele but rather a hypomorph. The transcriptional machinery occasionally makes mistakes which results in a small degree of leakage of WT transcripts.

## **3.4 Mouse models of cancer**

Cancer is a complex genetic disease, where gain of function mutations of oncogenes and loss of function mutations of tumor suppressor genes result in the transformation of a normal cell into a malignant proliferative tissue, a tumor. The tumor is like an overheating wound that eventually evolves in to a metastasizing disease. If left unchecked or unsuccessfully treated, the cancer leads to major organ failure and subsequent death (125). In Sweden the number of deaths from cancer related causes was 22 904 in 2012 and in the United States, 585 720 cancer deaths are projected in 2014 (126).

Mouse models have contributed significantly to our understanding of the origin, pathogenesis and biology of cancer. The most common mutations found in tumors from human patients, when introduced into the mouse, indeed led to initiation and progression of cancer. However, the genetically altered mouse models are challenged with being overly simplified, lacking passenger mutations and having a less complex mutational landscape, in some aspects casting doubt over the translational potential between cancer in genetic mouse models and patients. (127-129)

### **3.4.1 p53 knockout mouse**

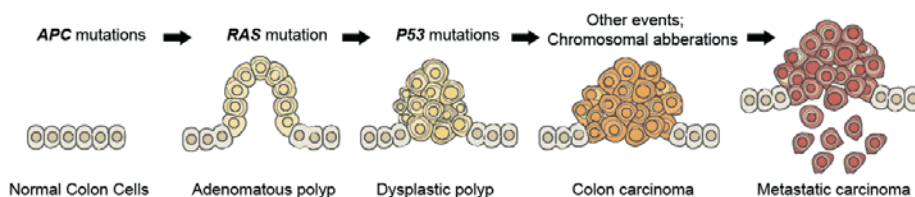
There are numerous p53-deficient or mutant mouse models. The p53 knockout model used in this thesis was generated in Tyler Jacks laboratory (130). All mice on a p53 knockout background succumb to spontaneous tumors and become moribund on average around 20 weeks of age. The tumor spectrum consists predominantly of lymphomas (>70%), but sarcomas and carcinomas are also common (130).

More relevant for human disease, mice heterozygous for p53 totally recapitulate the Li-Fraumeni syndrome, a hereditary disease where the patients are prone to develop cancers (sarcomas in particular). Similar to the p53 heterozygote mice, Li-Fraumeni patients lack one functional allele of p53 (98, 131, 132). The p53 heterozygote mice are prone to develop cancers (sarcomas in particular) with an average onset of disease around the age of 60 weeks (130, 133, 134). These mice are also more sensitive and prone to develop carcinogen and radiation induced tumors (135-139).

In papers I, II and IV the p53 knockout allele is used for genetic interaction studies, to test if our conclusions that the downstream effects of losing *Zfp148* are dependent on p53 activation. Notably, we don't allow them to age to the point where they develop spontaneous tumors.

### 3.4.2 *APC<sup>min/+</sup>* mouse model

The *Apc<sup>min/+</sup>* mouse used in paper II is an intestinal cancer model generated through a forward genetic screen with an inactivating point mutation in the tumor suppressor gene *APC* (adenomatosis polyposis coli) (140, 141). It was found that the multiple intestinal neoplasia (min) phenotype in these mice co-segregated with a mutation in the *Apc* gene, which had been found mutated in patients with colorectal cancer and in familial adenomatous polyposis (FAP) (142). Studies on the *Apc<sup>min/+</sup>* model stand as part of the foundation to the Vogelstein model (143), depicting cancer as a disease with linear progression that evolves and becomes progressively more invasive as clones with new mutations emerge within the tumor. Briefly, a transformed cell grows to become a small neoplasia, evolving into an adenoma, which mutates into an adenocarcinoma and finally becomes an invasive metastatic cancer. In this model, loss of the gene APC is an initiating event, mutation of RAS an early event, and loss of p53 a later event (143, 144) (Figure 1).



**Figure 1.** Vogelstein model depicting linear progression in colorectal cancer

One distinct difference between humans and mice concerning the role of *Apc* is that humans with mutant *APC* primarily develop tumors in the colon whereas mice develop tumors in the small intestine.(145)

### 3.4.3 *Kras*<sup>LSL-G12D</sup> mice

The *Kras*<sup>LSL-G12D</sup> mouse was generated by the Tyler Jacks laboratory. This mouse carries a latent point-mutant allele, G12D, immediately preceded by a LoxP flanked STOP cassette resulting in a null mutation that renders the mouse heterozygous for *Kras*. Cre-mediated recombination leads to deletion of the lox-stop-lox (LSL) sequence and expression of the constitutively active oncogenic protein K-Ras<sup>G12D</sup>. In lung epithelial cells the expression of K-Ras<sup>G12D</sup> leads to transformation and initiation of lung tumorigenesis (146). The transformed lung cells proliferate and progress to atypical adenomatous hyperplasia (AAH), which progress to adenomas that in turn progress to invasive adenocarcinomas (147). *CRE*-mediated recombination of *Kras*<sup>LSL-G12D</sup> in cells *in vitro* leads to increased proliferation and loss of contact inhibition, which are hallmarks of transformation and cancer (148-150).

### 3.4.4 *Braf*<sup>CA-V600E</sup> mice

The *Braf*<sup>CA-V600E</sup> mouse, generated in the Martin McMahon laboratory, carries, similar to the *Kras*<sup>LSL-G12D</sup> mouse, a latent point mutation that after *CRE* mediated recombination leads to expression of oncogenic B-Raf<sup>V600E</sup>. In contrast to the *Kras*<sup>LSL-G12D</sup> allele the *Braf*<sup>CA-V600E</sup> allele expresses WT B-Raf levels before *CRE* recombinase mediated conditional activation. Expression of oncogenic B-Raf<sup>V600E</sup> leads to similar transformation of lung cells *in vivo* and cells *in vitro* as expression of K-Ras<sup>G12D</sup>. However, one important difference in the progression of disease is that B-Raf<sup>V600E</sup> induced lung tumors do not progress to adenocarcinomas, due to oncogene induced senescence (151).

### 3.4.5 *p53*<sup>flox/flox</sup> mice

This mouse carries a conditional knockout allele for p53 and was generated by the Tyler Jacks laboratory (152). The conditional allele makes it possible with the help of the Cre-loxP system to knock out p53 in a temporal and spatial manner. We have combined this allele with the *Braf* and the *Kras* oncogenic alleles to generate *p53*<sup>flox/flox</sup>; *Kras*<sup>LSL-G12D</sup> and *p53*<sup>flox/flox</sup>; *Braf*<sup>CA-V600E</sup> mice. These mice combine constitutively activating mutations in *Kras* or *Braf* with the loss of two alleles of p53 after inhalation of *CRE* adenovirus, which result in a more aggressive lung tumor model that eventually develops invasive metastasis (153). We used these four lung cancer models described above in paper III.

## **3.5 Mouse models of atherosclerosis**

Atherosclerosis is a complex disease of the vascular wall where cellular changes progress for decades before they manifest in acute cardiovascular events, such as stroke or acute myocardial infarction. Defining the pathogenesis and mechanisms of atherosclerosis in humans is challenging for several reasons including the complexity of the disease, the slow progression in each individual patient, and the lack of noninvasive detection techniques. Hence, we depend on animal models of atherosclerosis to define mechanistic pathways. Over the last two decades mouse models have come to dominate the field of experimental atherosclerosis, providing a tool to uncover the pathogenesis and molecular mechanisms of atherosclerosis.

### **3.5.1 C57BL/6 strain**

As a species the mouse is highly resistant to atherosclerosis. Therefore, normal WT mice, irrespective of diet, do not spontaneously develop atherosclerosis. However, some inbred strains, especially the C57BL/6 mice, can develop atherosclerotic lesions when fed a high-fat and cholesterol rich diet that promotes hyperlipidemia (154). Nevertheless, the atherosclerotic lesions in C57BL/6 mice remain small and confined to the aortic roots and do not progress beyond the earliest phases of the disease, even after prolonged periods on high-fat diet and cholesterol rich diet (155, 156).

### **3.5.2 Apolipoprotein-E and Low-density lipoprotein receptor knockout mice**

To better recapitulate human atherosclerotic development, several mouse models targeting lipoprotein metabolism have been backcrossed into the C57BL/6 genetic background (157). The most well established and widely

used models are the Apolipoprotein E (ApoE) knockout model (158, 159) and the low-density lipoprotein receptor (LDLr) knockout model (160). In this thesis we choose to work with the ApoE knockout mice on a C57BL/6 N background, even though the LDLr knockout mice has a more human like lipoprotein profile. The ApoE knockout mouse is a more flexible model that progresses faster and also has an inflammatory component (161).

ApoE knockout mice develop spontaneous hypercholesterolemia which subsequently leads to the development of atherosclerosis independent of diet (159). Atherosclerotic lesions are formed throughout the whole aorta but are most profound in the aortic root and the aortic arch. The progression of atherosclerosis in the ApoE knockout mice can be accelerated with the use of high-fat and cholesterol rich diet. The use of high-fat diet shortens the time to develop advanced lesions to around 20 weeks, in contrast to a year on normal chow diet. In ApoE knockout mice, plasma cholesterol is packaged in chylomicron remnant (CMR) particles and very low-density lipoproteins (VLDL), in contrast to LDLr knockout mice and humans in which the cholesterol is packaged in LDL particles (156). However, there are humans with familial ApoE deficiency (162), and their lipid profiles are similar to the profile in ApoE knockout mice.

### **3.5.3 Bone-marrow transplantations**

We used bone-marrow repopulation experiments in paper IV to narrow down the effector cells that caused the atherosclerosis phenotype in *Zfp148*<sup>gt/+</sup> mice on ApoE knockout background (paper IV). Lethal irradiation followed by bone-marrow transplantation leads to a repopulation of the host (recipient) hematopoietic stem cell pool from the graft (donor) (163). If the transplanted bone marrow cells are from a transgenic background and the recipient is a WT, the recipient after successful engraftment will be a WT mouse with a

transgenic bone-marrow with all the hematopoietic derived cells including macrophages and t-cells becoming transgenic. Bone marrow transplantation is a widely used method to study the role of hematopoietic cells in atherosclerosis. (164)

## **3.6 Ethical considerations**

All animal experiments were approved by the Research Animal Ethics Committee in Gothenburg. The principle of 3R, reduce, replace & refine has been part of the experimental planning.

*"Nothing is particularly hard if you divide it into small jobs." –Henry Ford*



## 4 RESULTS & DISCUSSION

In this section I will summarize and discuss some of the key findings in each enclosed paper.

### **Paper I: Zfp148 Deficiency Causes Lung Maturation Defects and Lethality in Newborn Mice That Are Rescued by Deletion of p53 or Antioxidant Treatment**

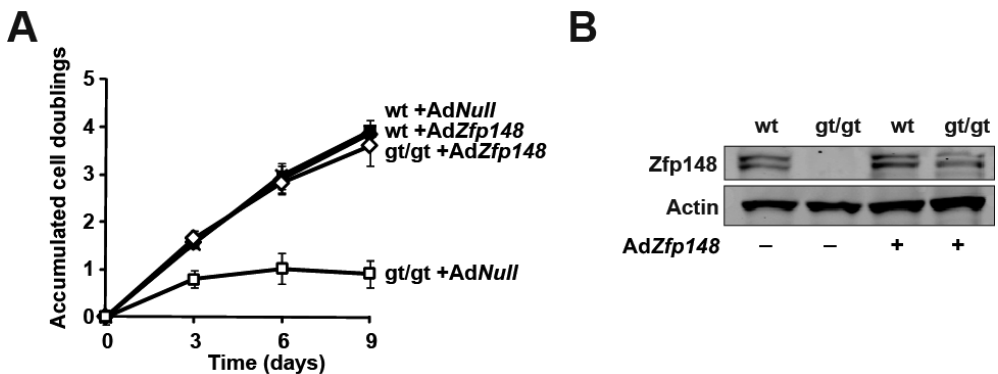
Our original rationale for knocking out Zfp148 was based on a bioinformatics screen for new transcriptional regulators of the vascular wall (165). However, the biology that unfolded took us elsewhere into cell cycle control, tumor suppression and oxidative stress.

#### **Generation and validation of Zfp148 deficient mice**

*Zfp148*-deficient mice were generated from a gene-trap (gt) ES-cell clone. Analysis confirmed that the gt vector was incorporated between exon 4 and 5 of *Zfp148*, thereby disrupting nearly 90% of the coding sequence including all four zinc finger domains. ES-cells were injected into C57Bl/6 blastocysts to achieve germline transmission of the *Zfp148<sup>gt</sup>*-allele. Analysis of mRNA levels from tissues of *Zfp148<sup>gt/gt</sup>* mice revealed that *Zfp148<sup>gt/gt</sup>* mice are hypomorphic, even though the protein is undetectable on western blots. Importantly, we have now established viable *Zfp148*-deficient mice, opening up for studies on the *in vivo* and cellular importance of endogenous Zfp148.

*Zfp148<sup>gt/gt</sup>* Mouse embryo fibroblasts (MEF) exhibited increased sensitivity to oxidative stress and slower proliferation at early passages and subsequently senesced prematurely. The sensitivity of *Zfp148<sup>gt/gt</sup>* MEFs to oxidative stress was coupled to increased activation of p53 independent of DNA damage. Importantly, reducing oxidative stress by the addition of the

antioxidant n-acetylcystein (NAC) or reducing oxygen pressure to physiological levels (3%) prevented the *Zfp148<sup>gt/gt</sup>* MEFs from entering premature senescence and reduced the activation of p53. Furthermore, the premature senescence did not manifest in *Zfp148<sup>gt/gt</sup>* MEFs if one or two alleles of p53 were deleted. As an important control, adding back *Zfp148* exogenously with viral vectors to *Zfp148<sup>gt/gt</sup>* MEF made them escape premature senescence, confirming that phenotypes observed in *Zfp148<sup>gt/gt</sup>* MEFs and mice are not caused by off-target effects of the initial gene targeting approach (**Figure 2**).



**Figure 2.** Ad back of *Zfp148* rescues *Zfp148* deficient MEF cells from premature senescence.

(A) MEF cell growth curve. (B) Protein levels of *Zfp148* in MEF cells from (A).

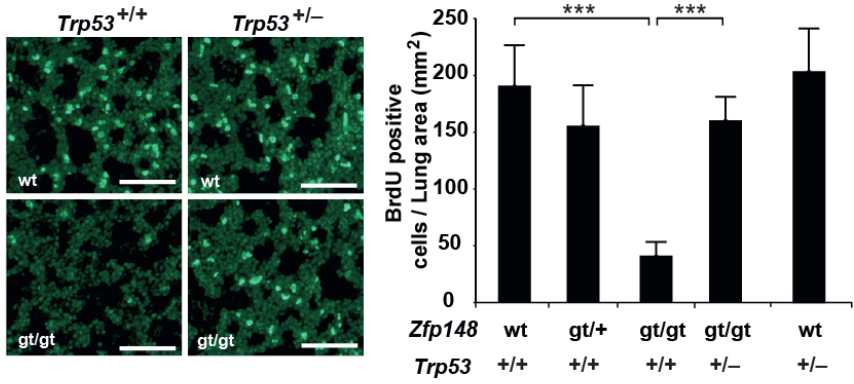
*Zfp148<sup>gt/gt</sup>* mice that were conceived through heterozygous intercrosses, develop normally and are found according to the expected Mendelian ratio throughout development. However, the observed ratio was halved postnatal day1 (P1), indicating that half of the *Zfp148<sup>gt/gt</sup>* mice died shortly after birth.

A closer examination of the P1 *Zfp148<sup>gt/gt</sup>* mouse revealed a lung maturation defect, suggesting that *Zfp148<sup>gt/gt</sup>* mice at P1 die of respiratory distress. A closer phenotypic analysis of the embryonic and P1 lungs of *Zfp148<sup>gt/gt</sup>* mice showed that *Zfp148*-deficiency disrupts cell proliferation at the saccular stage of lung development. Strikingly, loss of one allele of p53 in *Zfp148<sup>gt/gt</sup>* mice restored cell proliferation in the developing lung and prevented respiratory distress and neonatal lethality. Furthermore, NAC treatment of pregnant females also prevented respiratory distress and neonatal lethality in *Zfp148<sup>gt/gt</sup>* mice, strongly suggesting that the phenotype is triggered by oxidative stress (**Figure 3**).

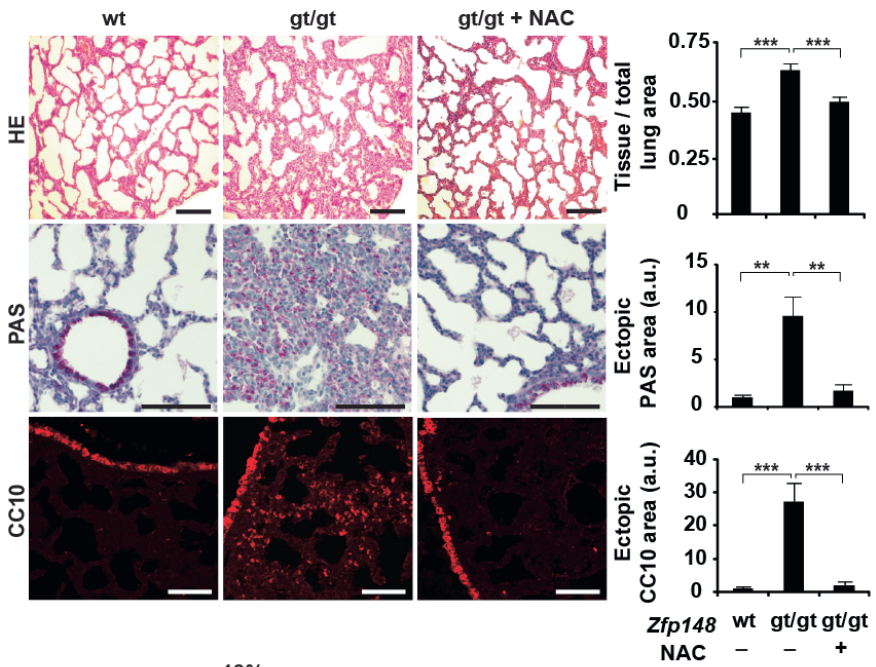
**Figure 3.** *Deletion of one copy of p53 or antioxidant treatment rescue lung maturation defect in Zfp148-deficient mice.*

(A) Proliferation defect in E.19.5 lungs is restored in *Zfp148*-deficient mice after loss of p53. (B, C) Lung maturation defects (B) and postnatal day 1 lethality (C) is prevented by antioxidant treatment in *Zfp148<sup>gt/gt</sup>* mice.

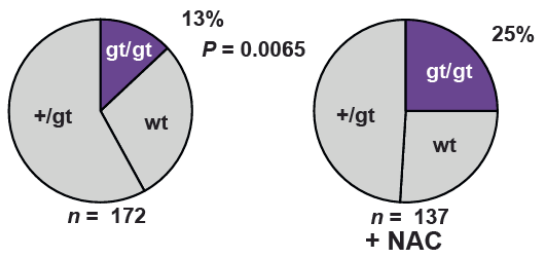
**A**



**B**



**C**

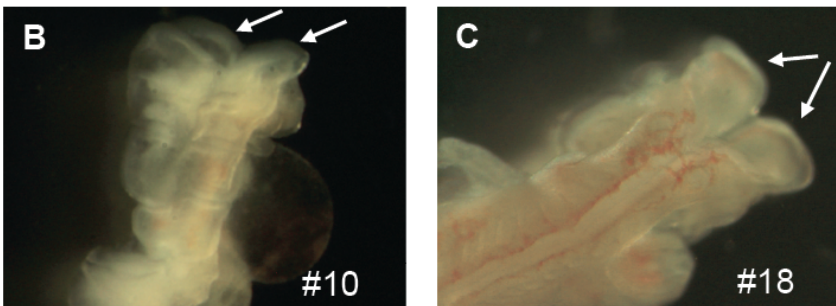


The results from paper I clear up some of the discrepancies that emerged from earlier gene-targeting attempts of *Zfp148*. Takeuchi et al suggested that *Zfp148* heterozygous mice developed a phenotype similar to the sertoli only-cell syndrome (87). However, this phenotype was not reproduced in our study or in any of the three other studies that inactivate *Zfp148* in mice (not counting conditional inactivation in specific tissues) (85, 86, 166). Moreover, there has been no follow up since the original publication dating 11 years back (87). Altogether, this suggests that the phenotype observed by Takeuchi et al. was not caused by deletion of *Zfp148*.

Another publication is in direct conflict with our findings. Woo et al. showed that *Zfp148*-deficiency caused death at embryonic day 8.5-10.5 with unclosed neural tubes and anemia (86). Interestingly, these *Zfp148*-deficient mice were generated by the same gene-trap ES-cell clone as we used to generate our *Zfp148*-deficient mice. Importantly, we observed the same phenotype (unclosed neural tubes and anemia) in embryos of the first generation (F1) intercrosses. However, we showed that this phenotype was caused by a second gene-trap that was inserted in another locus. After the second gene-trap was bred out in subsequent intercrosses, the embryonic lethal phenotype with unclosed neural tubes in *Zfp148<sup>gt/gt</sup>* mice disappeared. Thus, the early embryonic death was not caused by *Zfp148* deficiency (Figure 4).

**A**

Litter #	Embryo #	Phenotype	LacZ PCR	X-Gal staining	Zfp148 <sup>wt</sup> PCR
A	1	-	+	++	-
A	2	-	+	++	-
A	3	-	+	+	+
A	4	-	-	-	+
A	5	-	+	+	+
B	6	-	+	-	+
B	7	-	+	-	+
B	8	-	-	-	+
B	9	pale	-	-	+
B	10	unclosed neural tube, ectoderm defect, somite defect, unturned	+	N.D.	+
B	11	-	+	-	+
B	12	-	+	-	+
B	13	resorbed	N.D.	N.D.	N.D.
C	14	dead, very small, pale	+	+	+
C	15	-	+	+	+
C	16	-	+	+	+
C	17	abnormal, posterior defect, pericardial edema	+	N.D.	+
C	18	unclosed neural tube	+	+	+
C	19	-	+	+	+
C	20	-	+	-	+
C	21	unclosed neural tube, somite defect, posterior defect, pericardial edema	+	-	+



**Figure 4.** Integration of second gene-trap caused unclosed neural tubes and lethality in embryos from the F1 generation

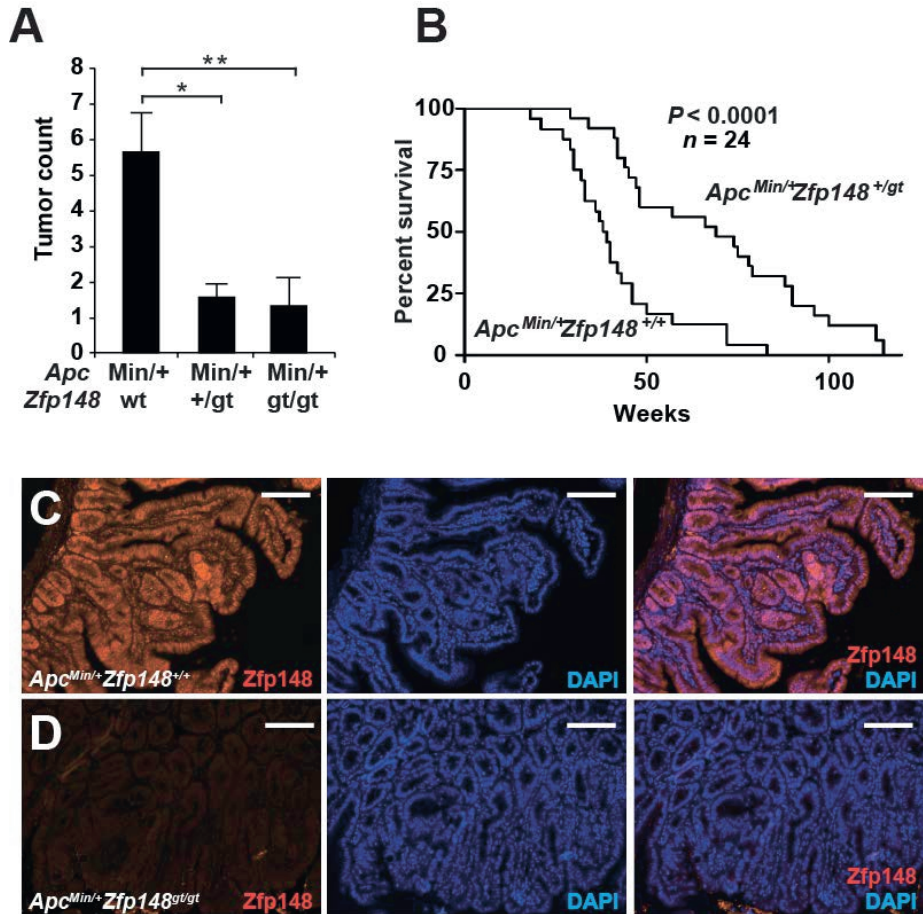
(A) Dissection of 21 E9.5 embryos of F1 generation intercrosses identified a proportion of embryos with unclosed neural tubes. Importantly, these mice had at least one intact Zfp148 allele. Moreover, the gene-trap vector was propagated to 79% of the brown offspring (F1 mice) of crosses between chimeric mice and C57Bl/6 mice, which is consistent with the presence of two gene-trap alleles in the injected ES-cells. (B, C) Dorsal view of E9.5 embryos exhibiting unclosed neural tubes (arrows).

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**Paper II: *Zfp148* deficiency reduces tumor formation in *Apc<sup>Min/+</sup>* mice in a p53-dependent manner**

In paper II, we show that *Zfp148*-deficiency reduces tumor formation in *Apc<sup>Min/+</sup>* mice. We hypothesized that *Zfp148*-deficiency would increase p53 activation and thus reduce intestinal tumor development. Indeed, the loss of either one or two alleles of *Zfp148* reduced the numbers of tumors in *Apc<sup>Min/+</sup>* mice. Additionally, *Zfp148*-deficient mice survived markedly longer in *Apc<sup>Min/+</sup>* background compared to *Apc<sup>Min/+</sup>* controls (**Figure 5**).

Mechanistically, we were able to show that the reduced tumor formation in *Zfp148<sup>gt/+</sup> Apc<sup>Min/+</sup>* mice depended on p53. However, we did not observe any differences in proliferation or apoptosis in tumors from *Zfp148<sup>gt/+</sup> Apc<sup>Min/+</sup>* mice compared to *Apc<sup>Min/+</sup>* controls, suggesting that the tumor protective effect of losing *Zfp148* occurs during tumor initiation and not during tumor progression. In support of this argument, tumors from *Zfp148<sup>gt/+</sup>; Apc<sup>Min/+</sup>* mice were equally invasive and showed similar histopathological grade at the moribund stage as tumors from *Apc<sup>Min/+</sup>* controls.



**Figure 5.** : *Zfp148* deficiency reduces tumor formation in *Apc<sup>Min/+</sup>* mice

(A) Tumor counts (B) Two year survival study. (C, D) Intestinal tumor stained for Zfp148 from (C) *Zfp148<sup>+/+</sup> Apc<sup>Min/+</sup>* mice and (D) *Zfp148<sup>gt/gt</sup> Apc<sup>Min/+</sup>* mice



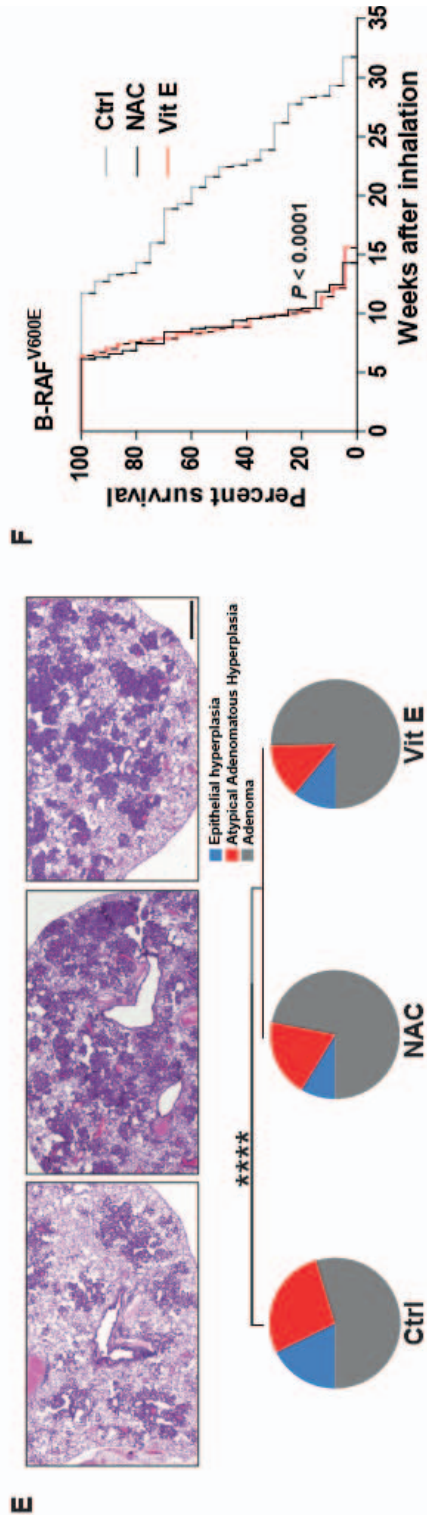
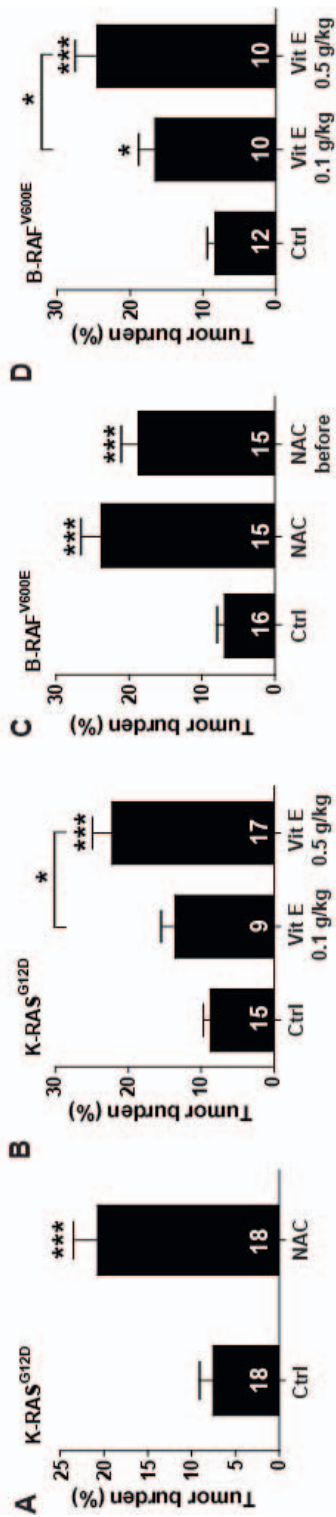
### **Paper III: Antioxidants accelerate lung cancer progression in mice**

This study was initiated serendipitously. In a natural continuation of Paper I, where the role of Zfp148 was studied in lung cancer progression, the antioxidant NAC was given with the hope that it might lower tumor progression or rather have no effect. To our great surprise we saw an acceleration of tumor progression.

We show that, supplementing the diet with the antioxidants NAC and vitamin E markedly increases tumor progression and reduces survival in mouse models of B-RAF- and K-RAS-induced lung cancer (Figure 7). Moreover, the increase in tumor burden after antioxidant treatment was dose dependent (Figure 7). Extensive mechanistic analysis revealed that antioxidants reduced levels of ROS, DNA damage and p53 activation in both mouse and human tumor cells resulting in increased tumor cell proliferation and progression of disease. Additionally, the presence of WT p53 was required for the pro-tumorigenic effect of antioxidants. Because p53 inactivation occurs late in tumor progression, antioxidants may accelerate the growth of early tumors or precancerous lesions.

#### ***Figure 7. : Antioxidants accelerate lung cancer progression in mice***

NAC and vitamin E increase tumor growth (A–E; size, number, and stage) and reduce survival (F) in mice with K-RAS<sup>G12D</sup>- and B-RAF<sup>V600E</sup>-induced lung cancer



One important question is if the doses administered to the mice are relevant for humans. It is well established that extrapolation of doses used in mice to a human equivalent dose by a simple conversion based on body weight is not accurate (167). We calculated the appropriate NAC dose for our mouse study with the help of body surface area conversion (167) so it would match prescription to chronic obstructive pulmonary disease (COPD) patients. For our vitamin E studies, we adjusted the dose in our mice studies to match the relative fold increase found in vitamin supplements compared to recommended daily intake (RDI). The standard vitamin E supplement exceeds RDI 20 times and can range up to 80 times RDI. Our vitamin E doses were adjusted to be 5 (low dose) and 50 (high dose) times higher than RDI for mice found in normal laboratory mouse chow. Hence, doses used in our study are relevant in a human context.

Another important question is whether the effects of NAC and vitamin E on lung cancer progression can be generalized to other antioxidants? NAC and Vitamin E have distinct molecular properties. Vitamin E is fat-soluble, regulates enzymatic activities, and is used as a dietary supplement, whereas NAC is water-soluble, participates in glutathione metabolism, and is used as a mucolytic agent (168, 169). Despite distinct molecular properties, unbiased whole-transcriptome sequencing of K-RAS<sup>G12D</sup> tumors revealed that the transcriptional changes induced by NAC and vitamin E were highly coordinated, indicating that they have a common mechanism of action. Moreover, the mechanism we propose is coupled to the drugs antioxidant properties. By lowering the levels of ROS and DNA damage, the extra antioxidants help cancer cells evade p53 activation. Collectively, these arguments indicate that antioxidants in general accelerate lung cancer progression.

Recent genomic analyses of cancers show a high frequency of gain of function mutations in the antioxidant master regulator Nrf2 (170-173), suggesting that decreasing levels of ROS promotes tumor growth. Consistent with this notion, targeting Nrf2 signaling and subsequently lowering endogenous antioxidants inhibits tumor progression (174-176). Furthermore, a recently published study shows that overexpression of Peroxiredoxin 6, an endogenous antioxidant enzyme, accelerates the progression of pre-existing skin tumors through reduction of ROS (177). In addition to confirming our data, this study shows that the pro-tumorigenic effects of antioxidants are not confined to lung tumors. Collectively, these studies are in line with our, showing that tumors thrive in the presence of increased antioxidants.

Antioxidants accelerate tumor progression. Our findings from Paper III published earlier this year have sparked a worldwide debate among scientists and physicians (178-182) about whether it can still be claimed that there is no harm in use of antioxidant supplements. A recent review in the New England Journal of Medicine (183) translates our findings to the clinical setting, highlighting it to primary care physicians.

Collectively, our data suggests that antioxidants may stimulate the growth and progression of undiagnosed lung tumors and should be used with caution, especially in individuals with increased risk of developing lung cancer including smokers and COPD patients.

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## **Paper IV: Loss of one copy of *Zfp148* reduces lesional macrophage proliferation and atherosclerosis in mice by activating p53**

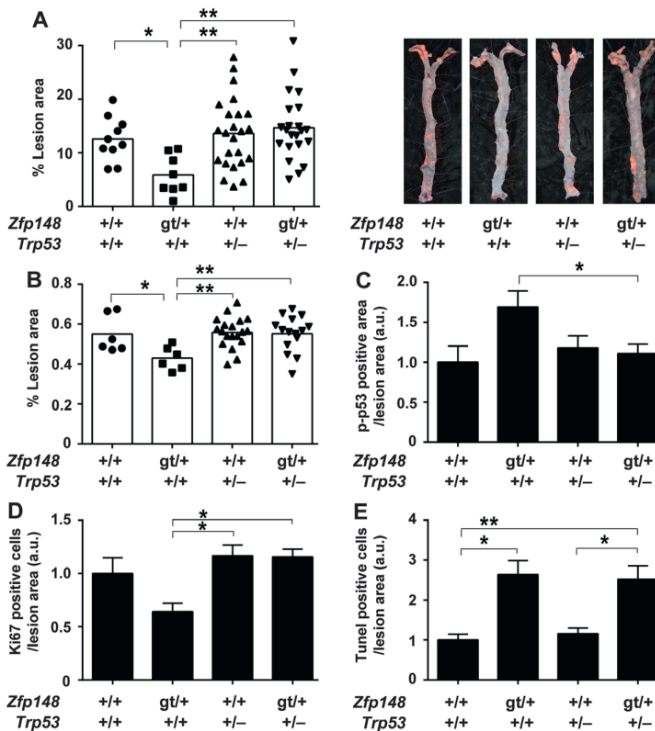
It has been previously established that loss of p53 and other tumor suppressors accelerates the progression of atherosclerosis (184-186). However, whether increased activation of p53 can protect against atherosclerosis remains unknown. Here we test the hypothesis that targeting *Zfp148* would reduce the progression of atherosclerosis in mice by activating p53.

We show that loss of one copy of *Zfp148* reduces atherosclerosis in *ApoE*-deficient mice independent of diet and without any apparent effects on lipid metabolism. With bone marrow transplantation experiments, we could pin down the effector cell responsible for reduced atherosclerosis in *Zfp148*<sup>gt/+</sup> *ApoE*<sup>-/-</sup> mice to the hematopoietic system. Mechanistic experiments show that lesional macrophages from *Zfp148*<sup>gt/+</sup> *ApoE*<sup>-/-</sup> mice exhibited reduced proliferation compared to controls, and that plaques from *Zfp148*<sup>gt/+</sup> *ApoE*<sup>-/-</sup> mice had less proliferating cells, increased apoptosis and increased p53 activation compared to controls. Importantly, there was no difference in atherosclerosis between *Zfp148*<sup>gt/+</sup> *ApoE*<sup>-/-</sup> mice and controls on a p53 heterozygote background, and there was no difference in p53 activity or cell proliferation. These results show that loss of *Zfp148* reduces atherosclerosis through p53 activation (Figure 8)

Whether the reduced atherosclerosis in *Zfp148*-deficient mice is mediated by oxidative stress is currently unknown. Interestingly, two studies revealed that deleting Nrf2, the master regulator of the endogenous antioxidant response, in *ApoE*<sup>-/-</sup> mice reduced atherosclerosis (187, 188). As shown in paper I and III, increased ROS levels can activate p53. It is therefore possible that increased p53 activity and decreased macrophage proliferation are part of the

underlying mechanism in the Nrf2 studies, suggesting a potential link to our study.

Drug therapies against atherosclerosis in clinical use or in late-stage clinical trials target inflammation or lipid metabolism. However, despite the success of cholesterol lowering drugs, disease progression is seen in many patients, emphasizing the need for alternative therapies that target other pathways (189). Paper IV suggests that therapeutic targeting of macrophage proliferation through the actions of Zfp148 and p53 might be one such alternative approach. Since Zfp148 reduced atherosclerosis independently of lipid metabolism, drugs targeting Zfp148 could have synergistic effects with lipid lowering drugs.



**Figure 8.** : Activation of p53 reduces atherosclerosis and lesional cell proliferation in Zfp148<sup>gt/+</sup> ApoE<sup>-/-</sup> mice.

**A**, Quantification of subendothelial lipid accumulation in aorta as percentage of Sudan IV-stained area per total area. **(B to E)** Quantification at the level of the aortic root for subendothelial lipid accumulation **(B)**, p-p53<sup>Ser18</sup> positive area/lesion area **(C)**, proliferating Ki67-positive cells/lesion area **(D)**, and terminal deoxynucleotidyl transferase dUTP nickend labeling (TUNEL) positive cells/lesion area **(E)**.

*"The ancient Oracle said that I was the wisest of all the Greeks. It is because I alone, of all the Greeks, know that I know nothing." – Socrates*

## 5 CONCLUSIONS

We conclude that:

### **Paper I**

- Zfp148 is required for structural maturation of the prenatal lung by preventing oxidative stress-dependent p53 activity during the saccular stage of lung development. The result demonstrates for the first time that Zfp148 plays a critical role for cell cycle progression in vivo, and establishes Zfp148 as a novel factor in mammalian lung development.

### **Paper II**

- Zfp148 is a modifier of the APC locus and that Zfp148 deficiency protects against intestinal adenomas by unleashing p53 activity. The results raise the possibility that therapeutic targeting of Zfp148 may protect against colorectal cancer by increasing p53 activity.

### **Paper III**

- Antioxidants NAC and vitamin E markedly increase tumors and reduce survival in mouse models of B-RAF- and K-RAS-induced lung cancer. NAC and vitamin E, which are structurally unrelated, produce highly coordinated changes in tumor transcriptome profiles, and increase cell proliferation by reducing ROS, DNA damage, and p53 expression in mouse and human lung tumor cells. Inactivation of p53 increases tumor growth to a similar degree as antioxidants and abolishes the antioxidant effect. Thus, antioxidants accelerate tumor growth by inactivating the ROS-p53 axis.



## Paper IV

- *Zfp148* deficiency reduces atherosclerosis in the *ApoE*<sup>-/-</sup> model without affecting lipid metabolism. We further show that the effector cell is of hematopoietic origin and that *Zfp148* deficiency confers protection against atherosclerosis by increasing p53 activation, thus reducing proliferation of lesional macrophages.

## 6 GENERAL DISCUSSION

Here I will discuss general findings that span beyond the individual papers.

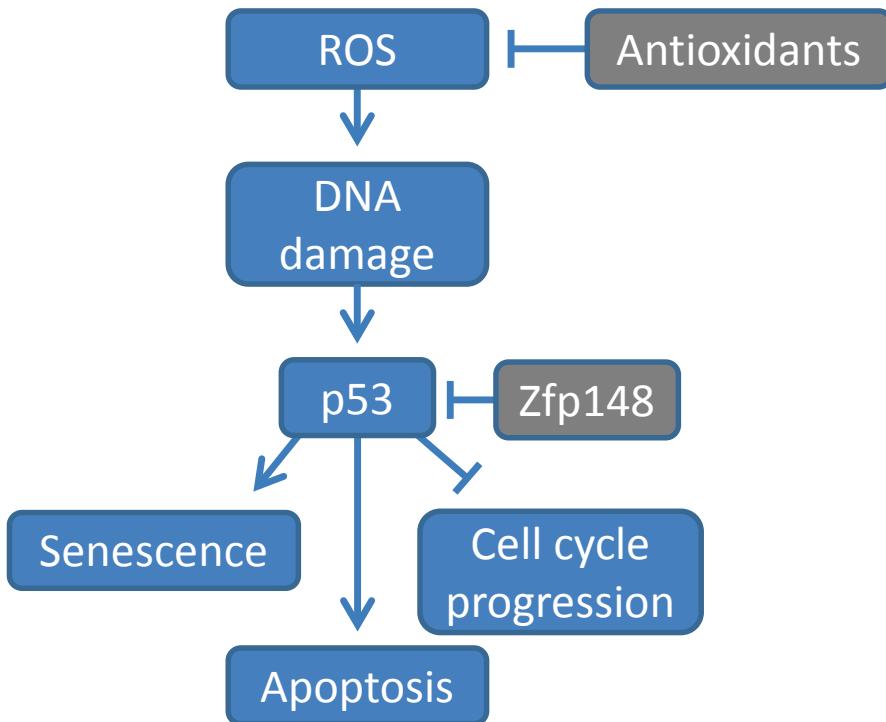
As we have seen throughout the thesis (paper I-IV), adjustments of the ROS-p53 axis can have major impact on both physiological and pathological processes. In some contexts, the adjustments of the ROS-p53 axis can be beneficial to the host organism (paper II + IV) and in others it can be harmful (paper I and III). There is a huge body of preclinical evidence which suggests that targeting the ROS-p53 axis can have beneficial effects on disease progression. However, it remains to be seen if we will be able to therapeutically manipulate the ROS-p53 axis to successfully treat human malignancies, including cancer and atherosclerosis.

In this thesis we establish Zfp148 as a potential target for increasing p53 activity.

Zfp148-deficient MEFs, lung cells and macrophages display a lowered threshold to p53 activation in response to oxidative stress. Although indirect mechanisms might be underlying the activation of p53, three lines of evidence support the possibility that Zfp148 primarily regulates p53. Firstly, there is a physical interaction between Zfp148 and p53, which is dependent on the zinc finger domains of Zfp148 and the C terminal parts of the DNA binding domain of p53 (79, 83). Secondly, in response to oxidative stress, Zfp148 deficient cells display increased p53 activation independent of DNA damage (paper I and IV). Finally, there is clear evidence of genetic interaction between Zfp148 and p53 (paper I, II and IV). However, to confirm the proposed mechanism of regulation, further biochemical analysis of Zfp148 is required.

Briefly, our data suggests that targeting Zfp148 and its downstream network might be beneficial in preventing intestinal cancer and atherosclerosis through activation of the ROS-p53 axis. However, our data also show that excessive activation of the ROS-p53 axis can lead to defects in lung development and shorten lifespan. Finally, we show that decreasing the ROS-p53 axis through antioxidant supplements can lead to increased tumor progression.

A graphical abstract depicting the ROS-p53 axis (blue boxes) and the consequences of our interventions throughout the thesis (grey boxes) is provided (Figure 9).



**Figure 9.** Graphical abstract of ROS-p53 axis, summarizing our findings from papers I-IV.

*"I never think about the future - it comes soon enough." –Albert Einstein*

## 7 FUTURE PERSPECTIVES

We are following up our findings in the papers included in this thesis.

- I. Loss of Zfp148 activates p53 and causes a lung maturation defect. We are identifying the downstream targets of Zfp148 by combining genome-wide expression analysis and with genome-wide transcription factor occupancy analysis. We are also genetically dissecting the molecular pathways causing the lung defect in *Zfp148*-deficient mice.
  
- II. *Zfp148*-deficiency reduces tumor formation in *Apc<sup>Min/+</sup>* mice. We are cross breeding the *Zfp148*-deficient mice into several tumor models, including lymphoma, sarcoma and lung cancer models. We will evaluate if targeting *Zfp148* in other tumor models will have similar effects as in *Apc<sup>Min/+</sup>* mice.
  
- III. We are evaluating the effect of Antioxidants on several cancer models, including colorectal cancer, melanoma and leukaemia. Also we are studying the effect of NAC on lung cancer incidence in COPD patients in a Swedish COPD registry with 7209 patients.
  
- IV. Experiments on LDLr knockout mice are ongoing to validate our findings in the ApoE knockout model. We are also working on translating our findings to the clinic by analysing *Zfp148* in the Biobank of Karolinska Endarterectomies (BiKE) containing more than 800 human atherosclerotic plaques and plasma samples along with clinical data from individual patients.

## ACKNOWLEDGEMENT

Through the course of my life, I have come across a lot of people that directly or indirectly have contributed to the making of this thesis. Without you this particular thesis wouldn't have existed.

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**It-Sven** and **Magnus**, the heart of Wlab? You guys make sure the clock ticks 24/7. There is not a problem that you cannot fix!

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