

The environmental stress response as a target for therapeutic intervention

Jennifer Harbottle

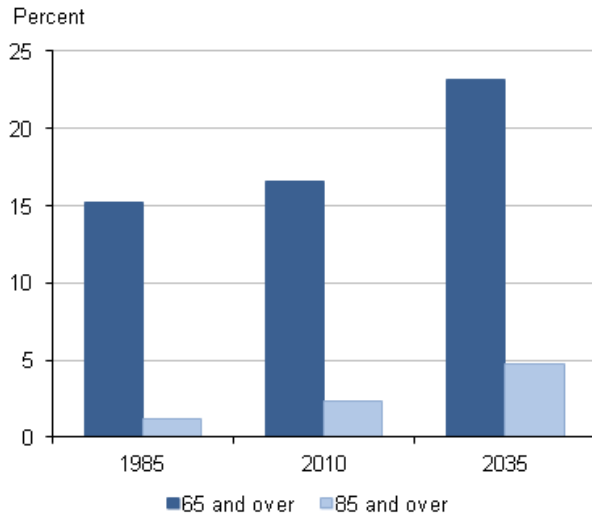
Supervisor: Dr Andreas Kolb

MRP Inter-institutional Post-Graduate Competition 2016

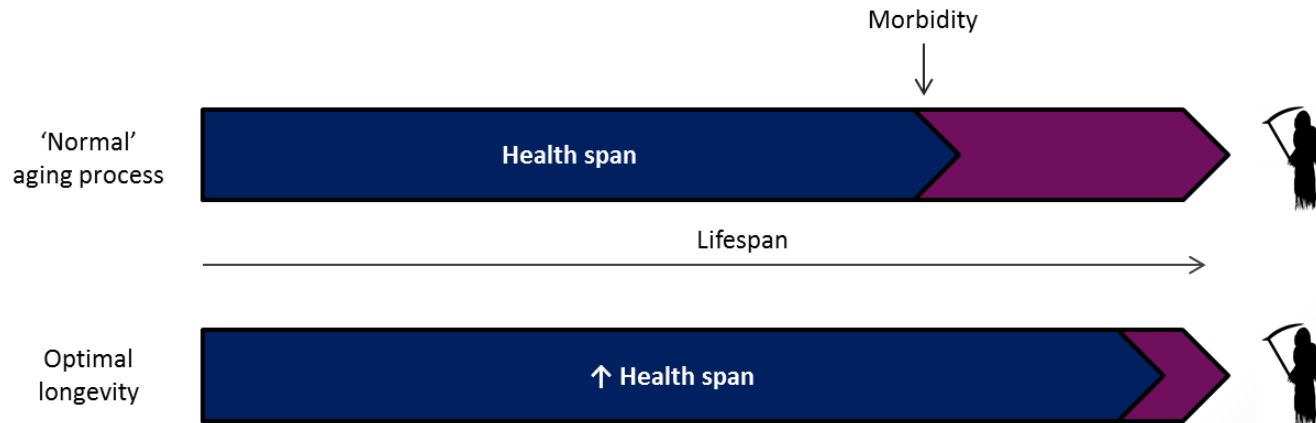
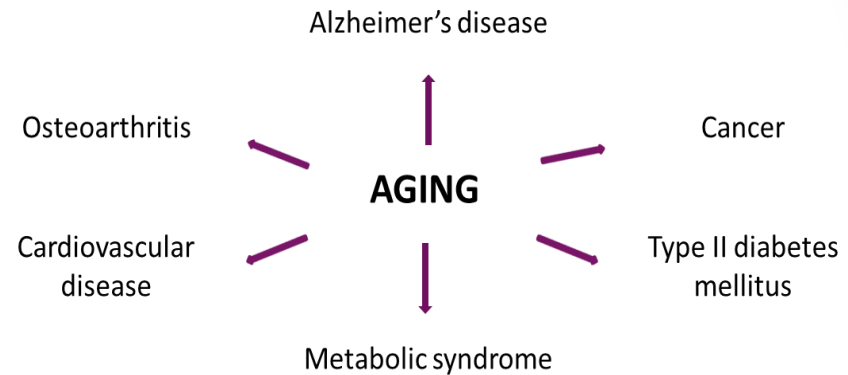
27-28 June 2016



Aging and health span



Office for National Statistics, 2012

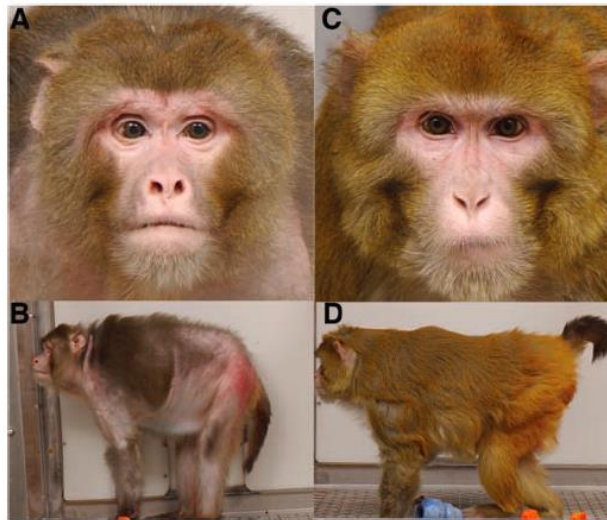


A common denominator



Normal diet

Calorie restriction



Stress resistance

Upregulation of the environmental stress response

Nrf2 drives the ESR

Nrf2, the “guardian of health span and the gate-keeper of species longevity” (Lewis et al. 2010)

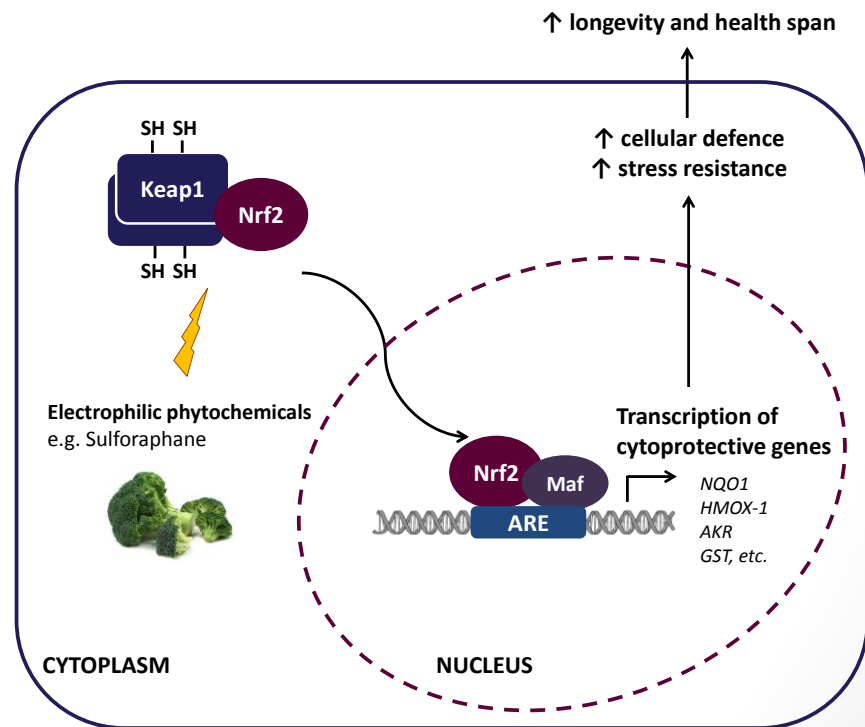
↓ Nrf2 signaling with age



↓ antioxidant defence
↓ stress resistance



↑ disease susceptibility



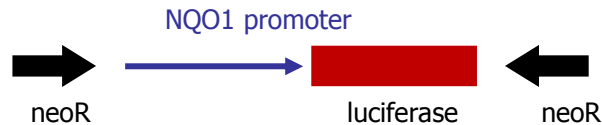
Phytochemicals activate the ESR



- Aim 1: Identify and characterise novel, natural compounds that induce the ESR.

A cell based assay system for the analysis of phytochemicals activating the ESR

Stable transfection of HepG2 cells & selection of HepG2 C1 clone

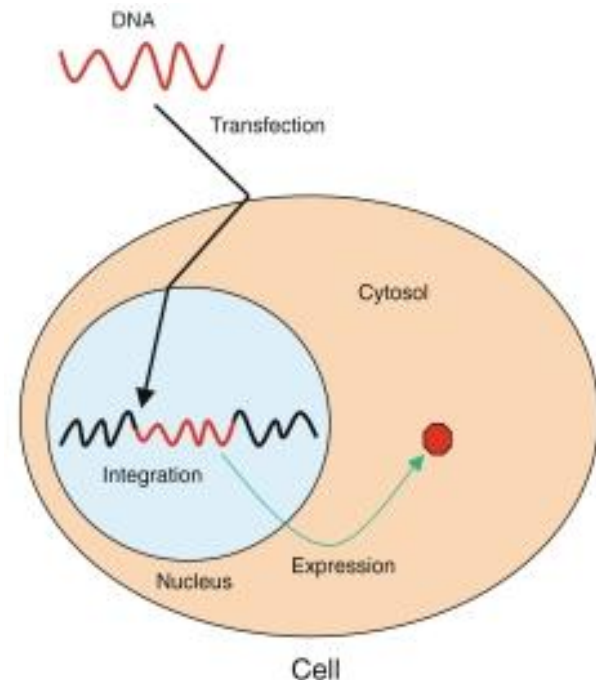


Validation of bioluminescent reporter system using sulforaphane

High throughput screening of natural chemical libraries

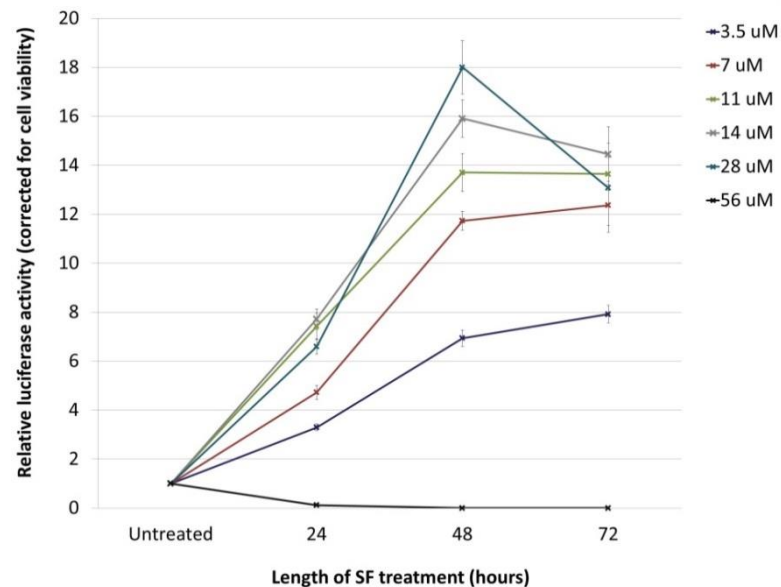
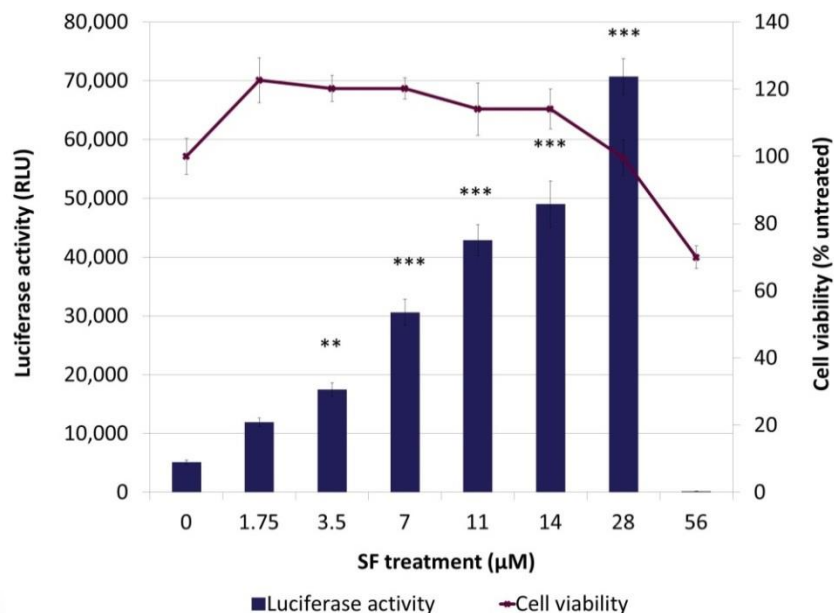
Identification and characterisation of compounds that activate the ESR

Stable transfection



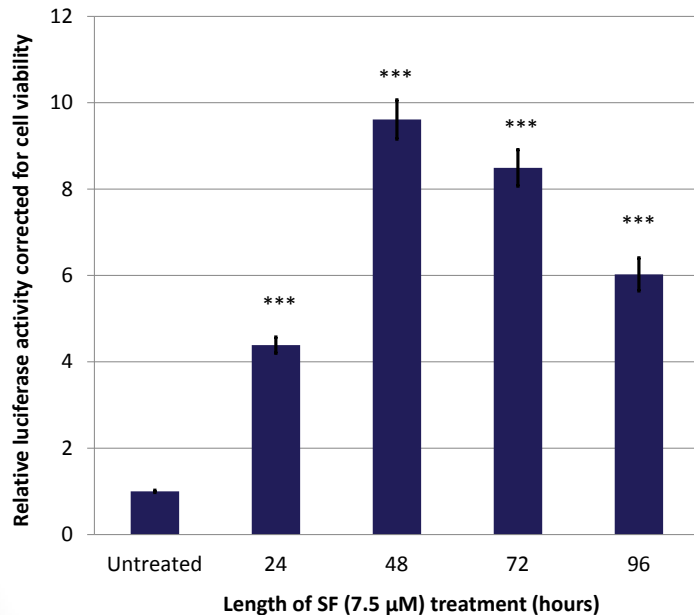
Validation with sulforaphane (1)

pNQO1-luc reporter induction

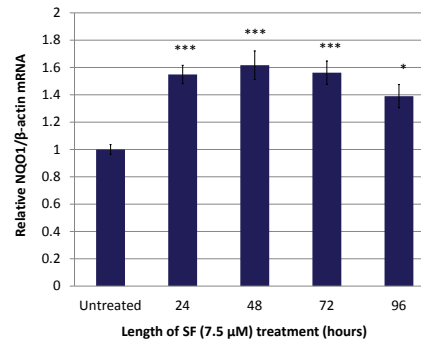


Validation with sulforaphane (2)

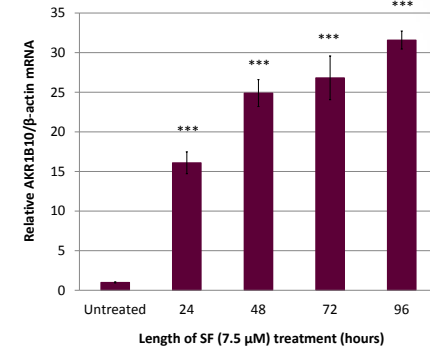
pNQO1-luc reporter induction



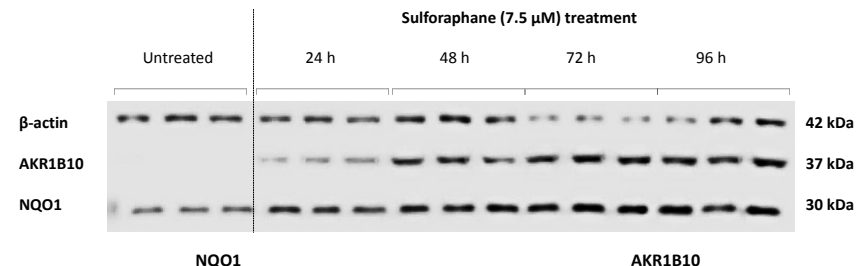
NQO1 gene



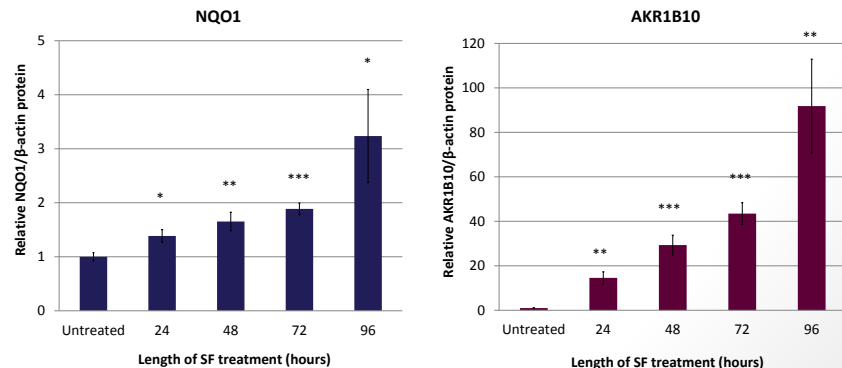
AKR1B10 gene



NQO1 protein

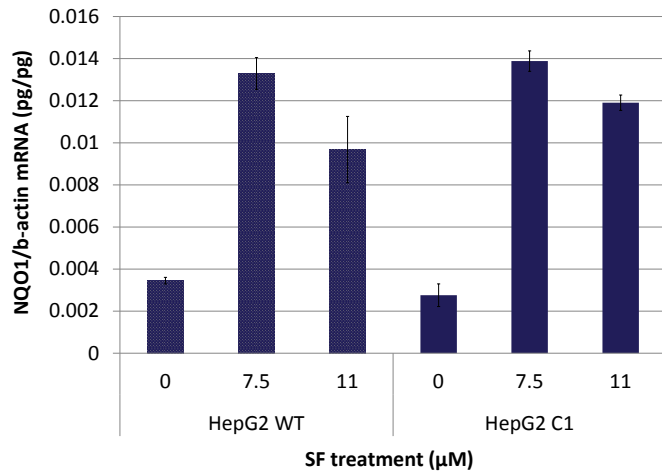


AKR1B10 protein



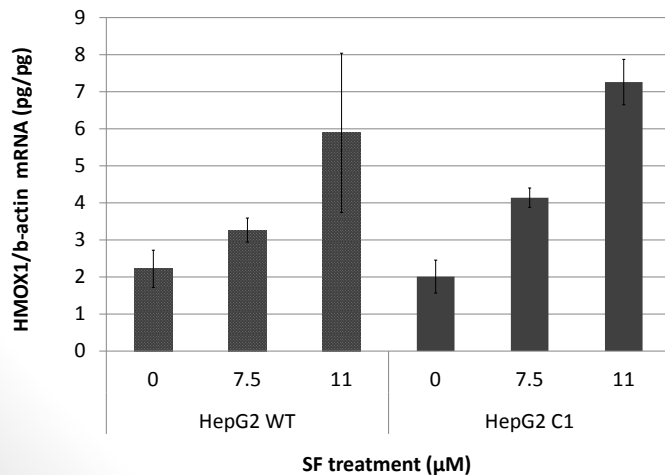
Validation with sulforaphane (3)

NQO1 gene

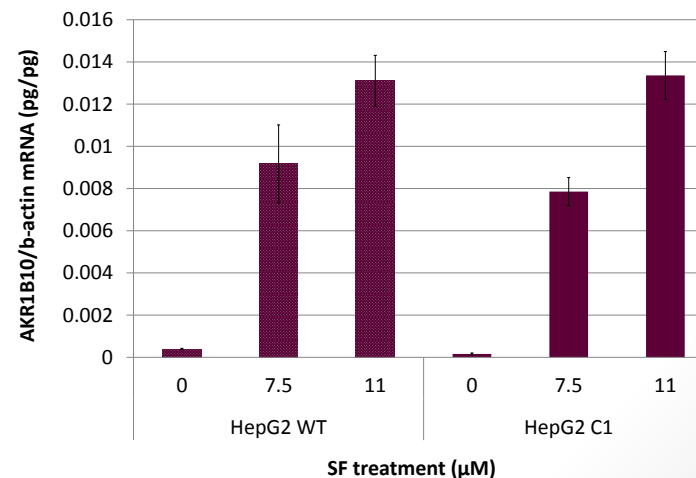


Endogenous ESR gene expression
in HepG2 WT 'vs' HepG2 C1

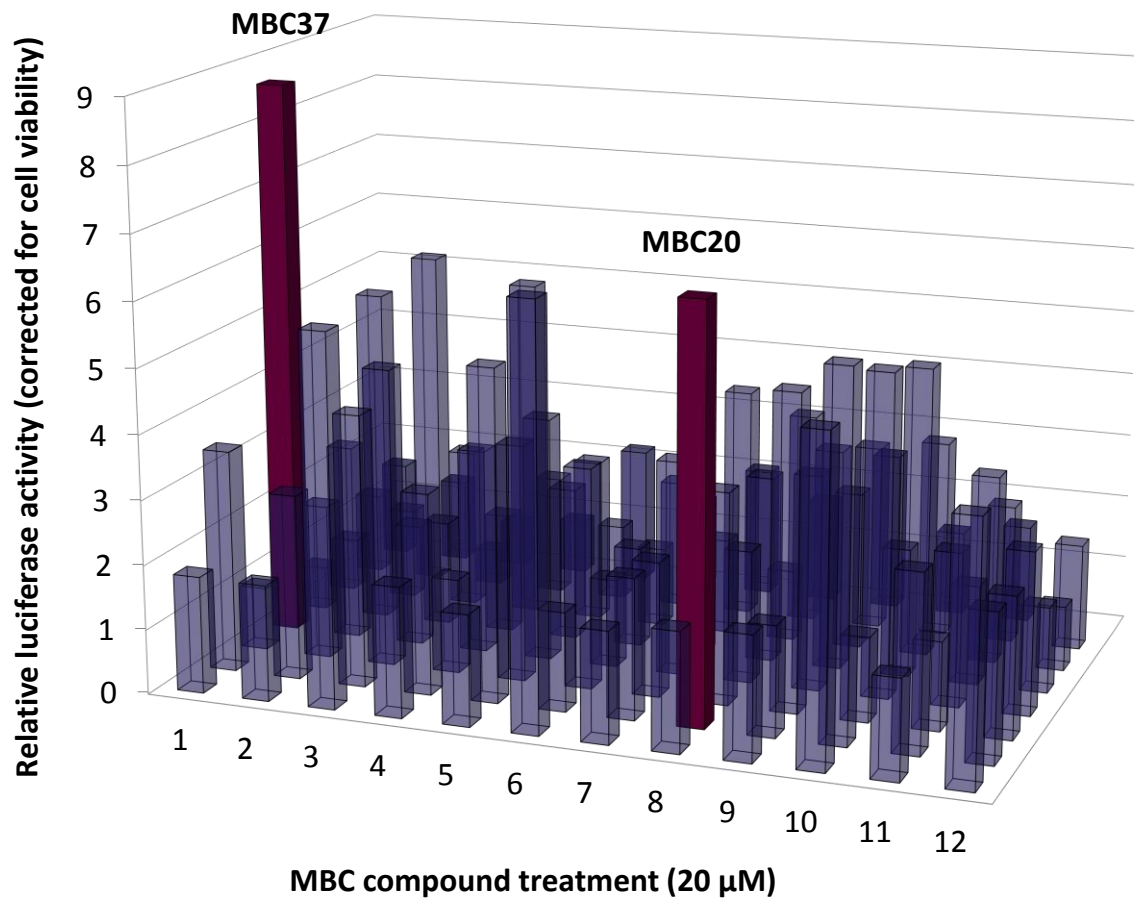
HMOX1 gene



AKR1B10 gene

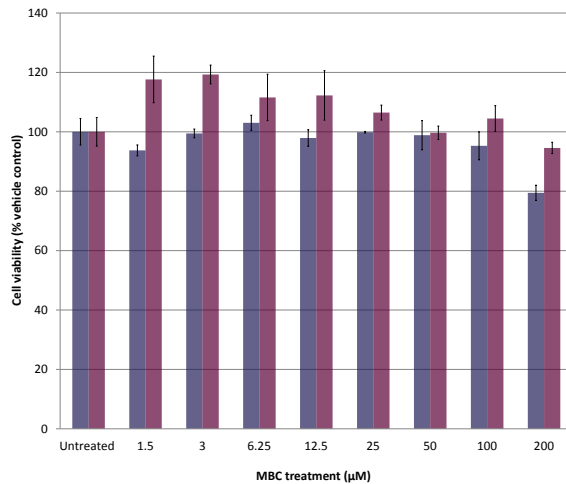


MBC library screening

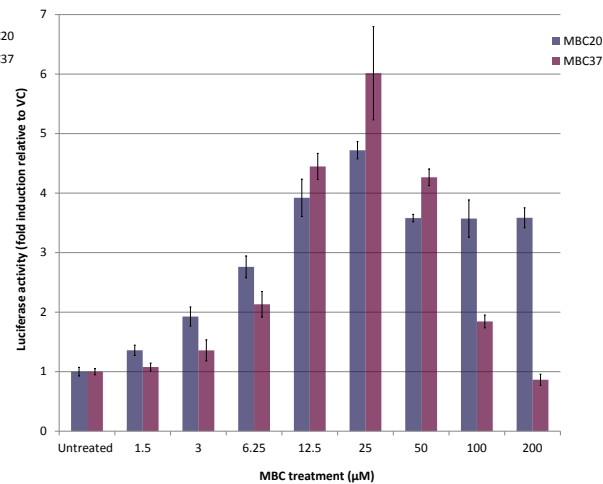


MBC20 and MBC37

Cell viability (0-200 μ M)

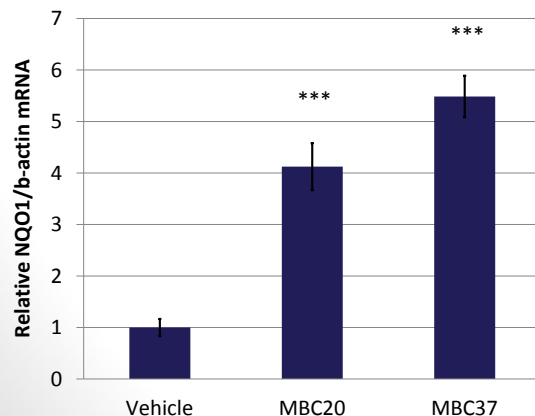


Reporter induction (0-200 μ M)



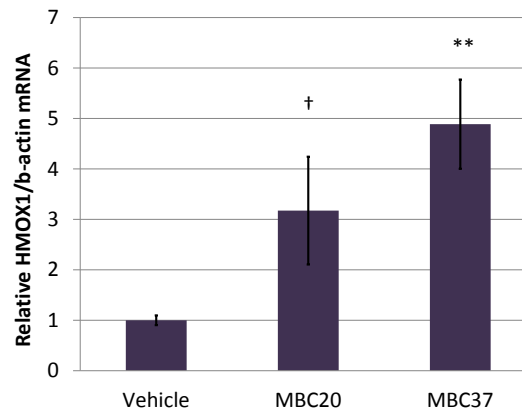
- No apparent cytotoxic effect
- Dose-dependent reporter activation

NQO1 gene



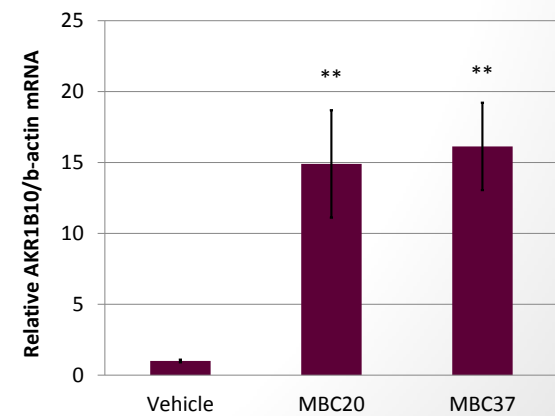
Cell treatment

HMOX1 gene



Cell treatment

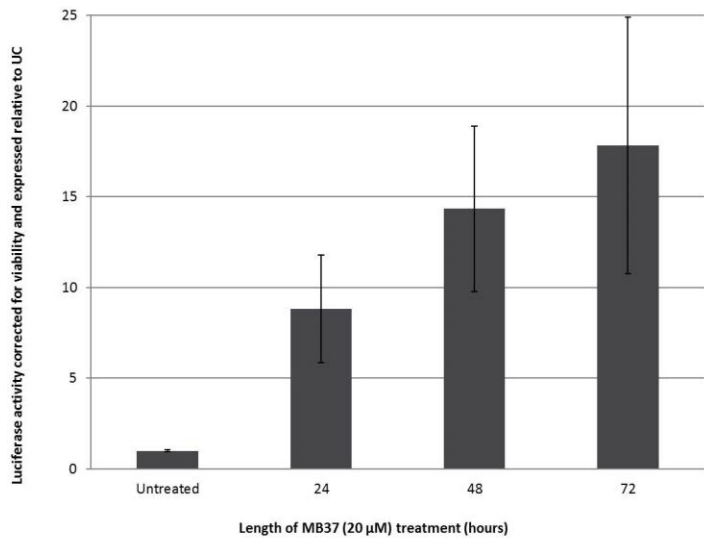
AKR1B10 gene



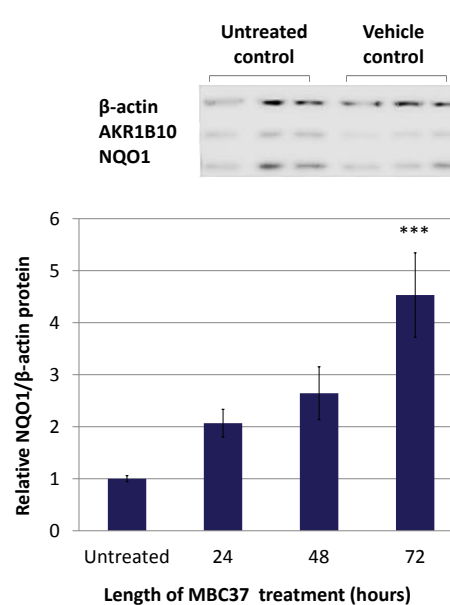
Cell treatment

MBC37 (0-72 h)

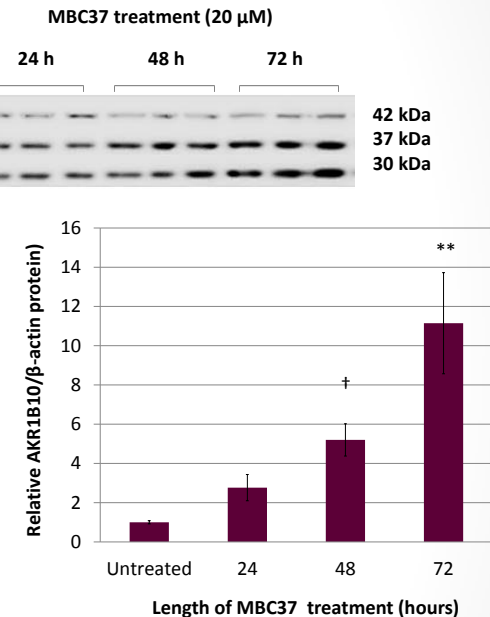
Reporter induction



NQO1 protein



AKR1B10 protein



- MBC37 ↑ ESR protein expression over time
- ↑ biopotency and ↓ cytotoxicity compared with SF

→ Successful screening

What now?

- Further characterisation of MBC20 and MBC37
→ Bioavailability, absorption, metabolism
- PhytoQuest library screening (MRCT)
→ Stereochemistry and ASR.



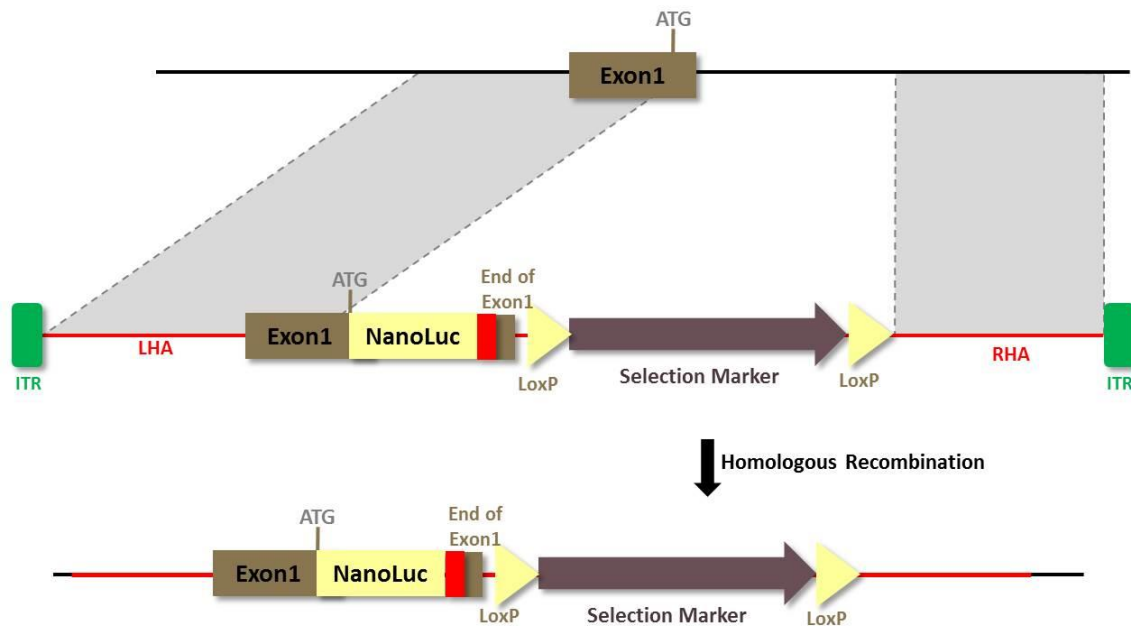
Limitations...



- **... Due to the nature of the transfection method.**
 - Random integration site;
 - Variable transgene copy number.
 - **... Due to the complexity of mammalian transcription .**
 - Regulatory elements (*cis*-acting enhancers/silencers).
- Simply inserting a promoter cannot capture the whole picture.

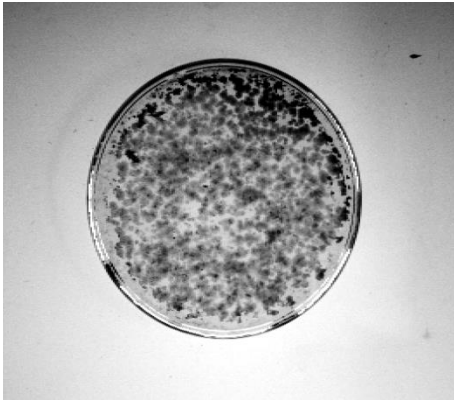
AAV-mediated integration

Aim 2: Site-specific integration of the luciferase reporter gene into the locus of the *HMOX1* gene.

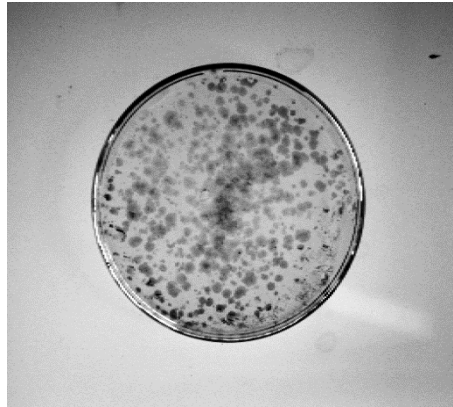


→ The NanoLuc reporter will reflect the expression pattern of endogenous *HMOX1*

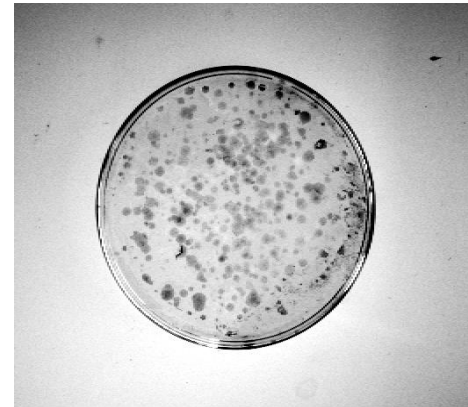
rAAV dilution range



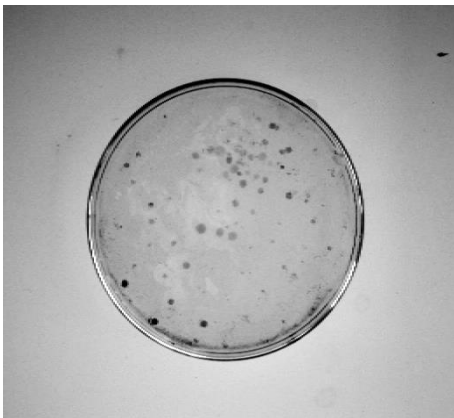
1×10^5 copies



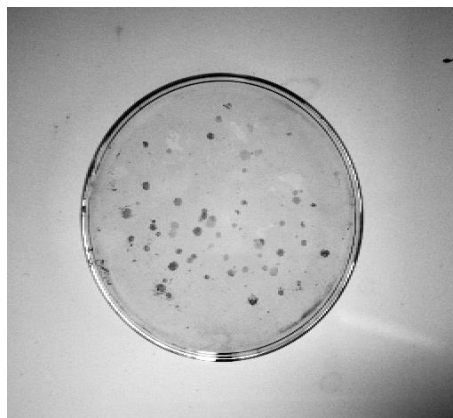
5×10^4 copies



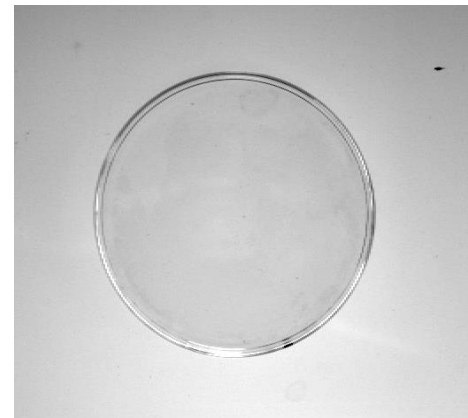
1×10^4 copies



5×10^3 copies

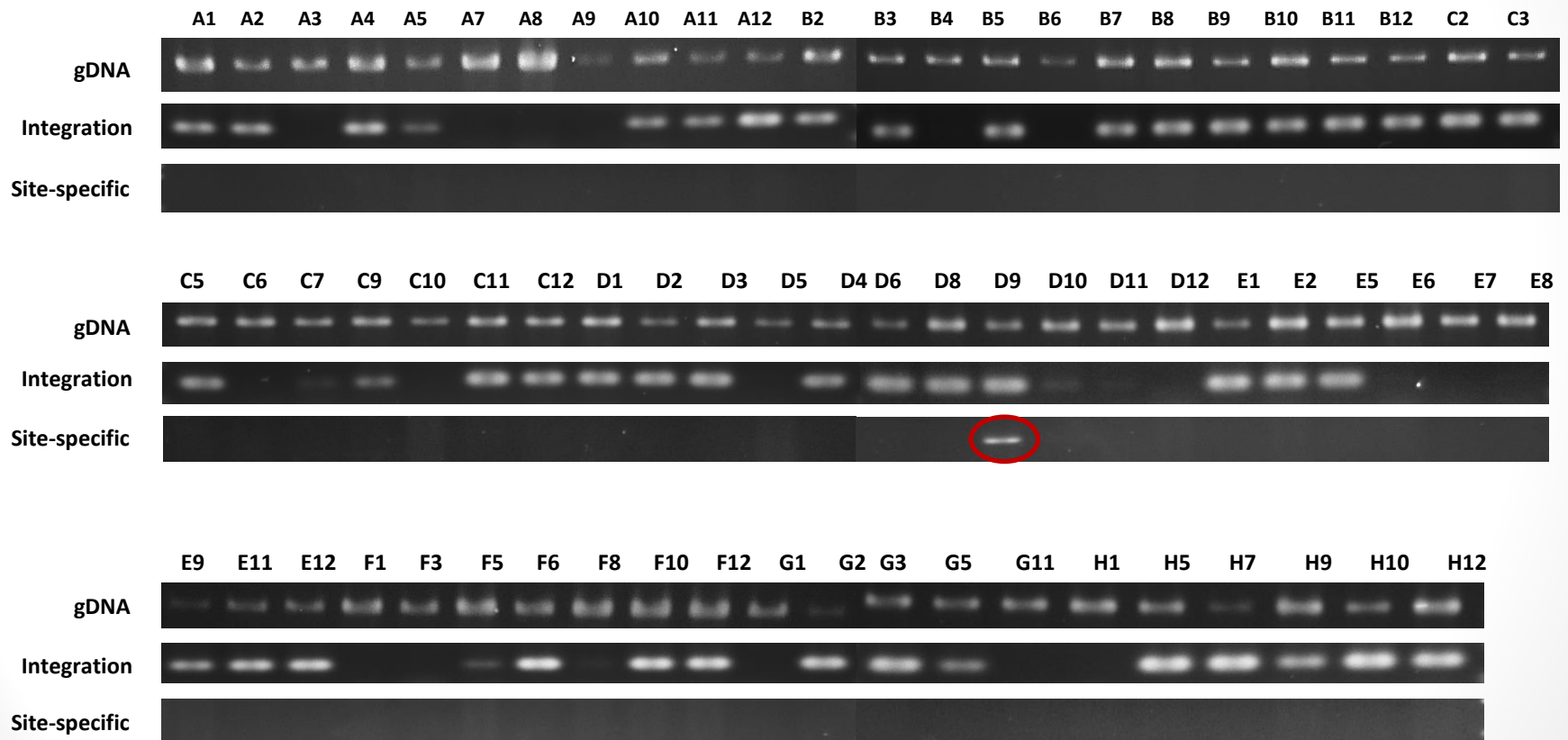


1×10^3 copies



Not infected

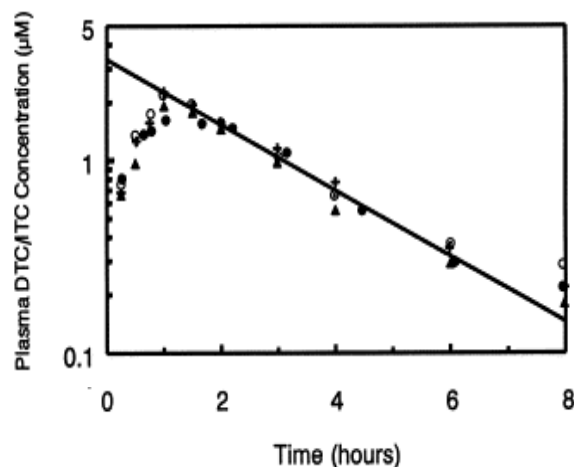
Screening



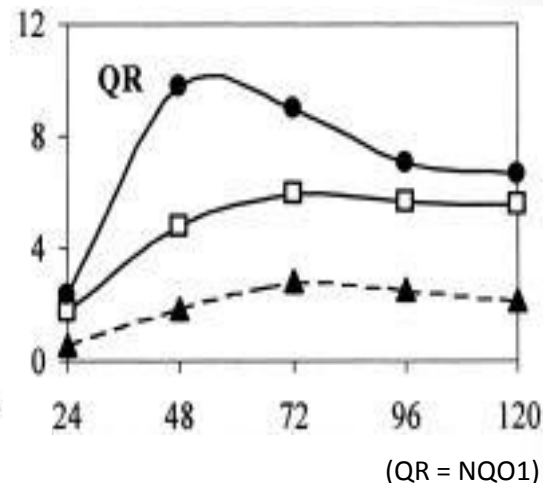
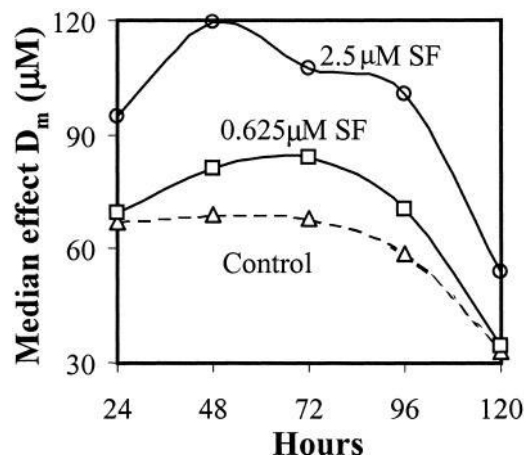


The ESR – A long term effect?

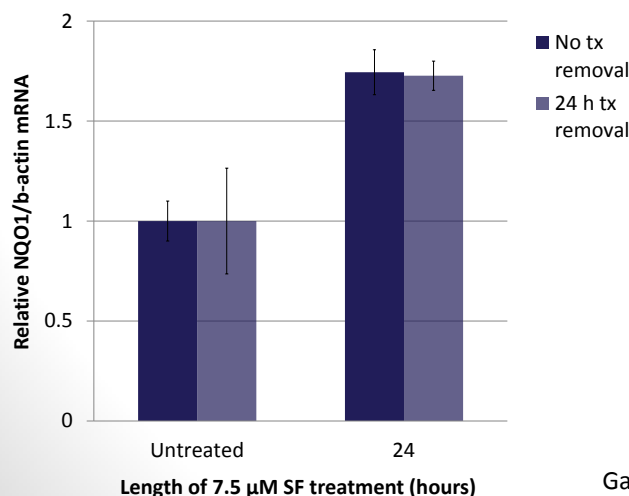
Plasma SF



Prolonged protection against menadione



NQO1 mRNA

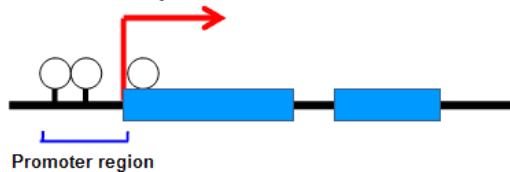


- Does SF evoke an antioxidant response that persists?
- Are epigenetic mechanisms involved?

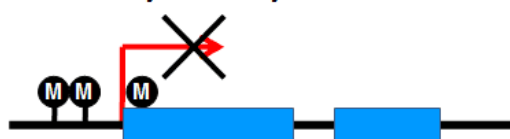
Phytochemicals and epigenetics

- Epigenetic silencing of Nrf2 and NQO1 with \uparrow age.
- Phytochemicals upregulate gene promoter demethylation
→ Facilitate gene transcription.

Genes that can be expressed



Genes inactivated by DNA methylation



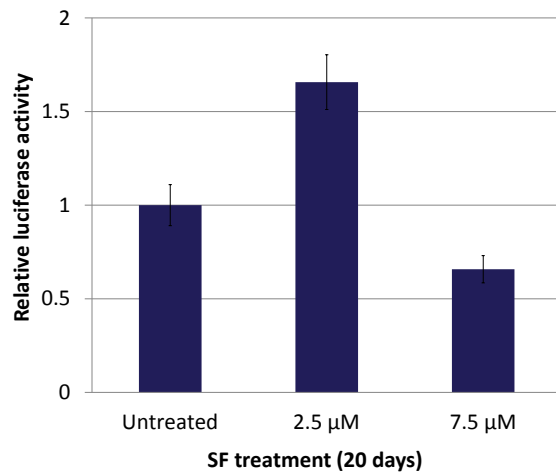
M Methylated

○ Unmethylated

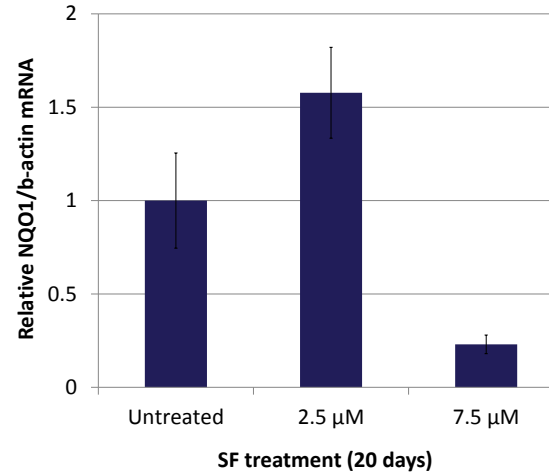


20-day SF treatment

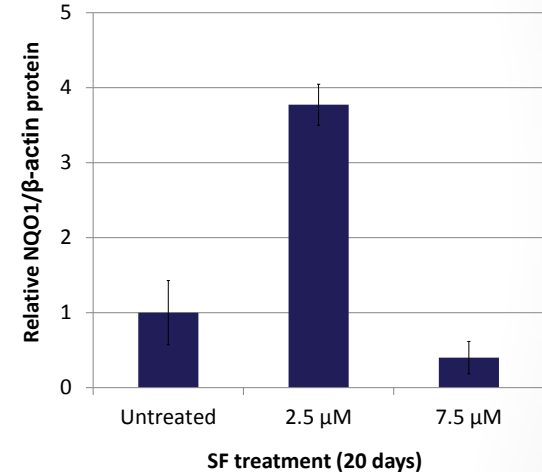
Reporter induction



NQO1 gene



NQO1 protein



Does SF (2.5 μ M) treatment demethylate ESR gene promoter?
If so, how stable is this epigenetic modification?

The big picture

- Identification and characterisation of compounds that elicit cytoprotection
 - Food fortification
 - Production of plants with enhanced nutritional quality
 - Optimise health span
- A better understanding of how food provides a conditioning environment that shapes the activity of the (epi)genome and determines the stress adaptive response.



Acknowledgements

Lab group supervision and support

Dr Andreas Kolb

Linda Petrie

Christopher Knowles

Patrikas Pultinevicius



Marine Biodiscovery Centre

Professor Marcel Jaspars

Dr Wael Houssen



Genomics

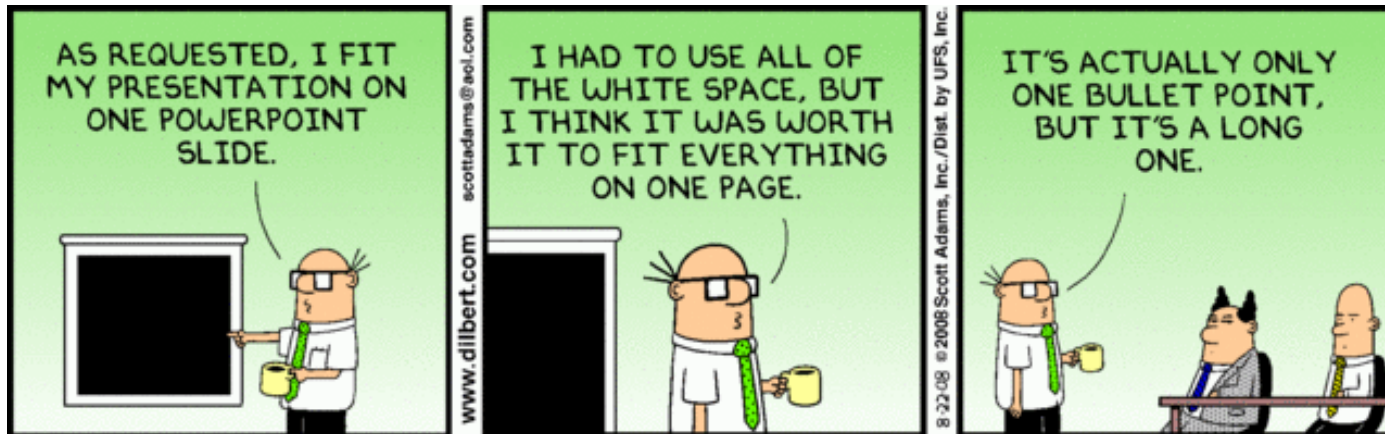
Pauline Shiach



eastbio
the East of Scotland Bioscience Doctoral Training Partnership

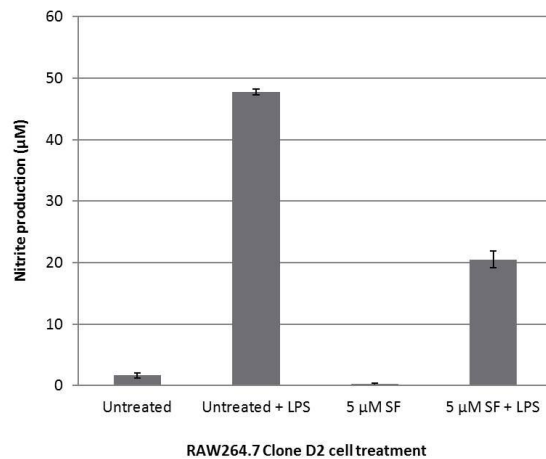


Thank you for listening

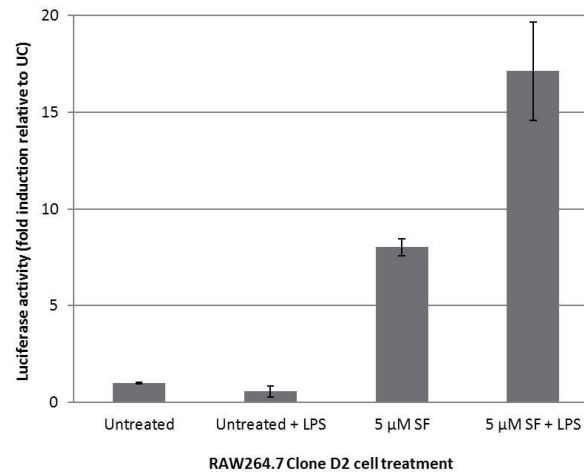


RAW264.7 NQO1-luc D2

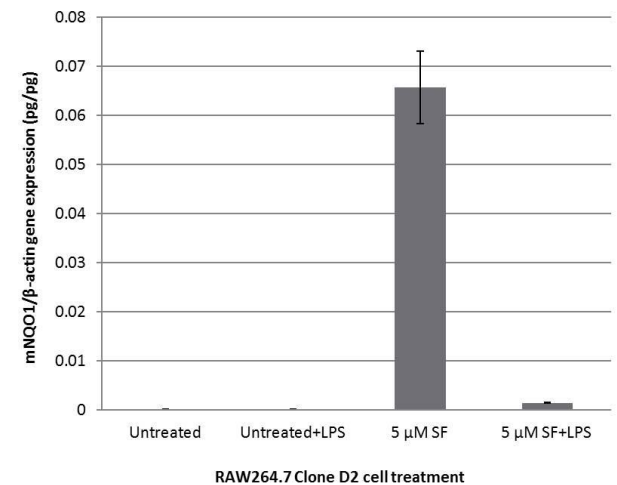
Nitrite production



NQO1-luc reporter induction



NQO1 gene expression



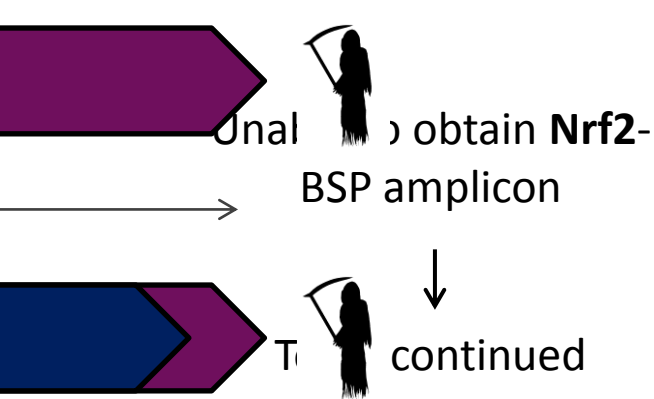
gDNA isolation from HepG2 C1 cells after 20 days treatment with SF (n=3)



Bisulfite conversion of gDNA



PCR with a selection of **Nrf2**-BSP and **NQO1**-BSP primers that span a CpG island within the respective gene promoters



Obtain 335 bp **NQO1**-BSP amplicon from nested PCR



Purify and clone into pGEMT vector



PCR & Sequence positive clones



Assess methylation status of 24 CpG dinucleotides (> 10 clones)

