Myalgic Encephalomyelitis/Chronic Fatigue Syndrome (ME/CFS) -a complex disease that has come of age with the SARS-CoV-2 pandemic

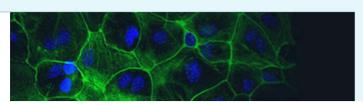
Emeritus Professor Warren Tate FRSNZ CNZM Department of Biochemistry University of Otago, Dunedin



Complex Chronic Illness Support 40th Anniversary Celebration July 3rd 2021







What was my stimulus to become involved in ME/CFS research?

Began with the dramatic onset in 1990 of a mystery disease with pervasive fatigue in my young daughter

- Medical practitioners in Dunedin in 1990 knew nothing of the disease and the default view was it was likely a psychological 'perception' disorder
- despite NZ having an outbreak 'Tapanui Flu' in 1984
- I put in a research proposal to the Health Research Council but reviewers questioned whether there was a real disease to study





The origins of ME/CFS

Documented Outbreaks of infectious diseases causing an ongoing post viral fatigue syndrome



Iceland 1946-49 (1090 cases) (Akureyri disease)

Royal Free Hospital London, 1955 (292 cases) (ME)

Incline Village Nevada 1984 (175 cases) (CFS)



A dominant global voice who firmly believed he had the answer to ME/CFS (1985-2005)



- Sir Simon Wesseley (Professor of Psychological Medicine Kings College London
- ME is "a perception disorder", an "opting out syndrome"
- "a general disorder of perception, perhaps of both symptoms and disability. At the heart of this misperception lies the sense of effort. CFS patients clearly experience increased effort in everyday physical and mental tasks"

In NZ
Medical, Social,
and Societal
attitudes towards
ME/CFS
my 30 years experience

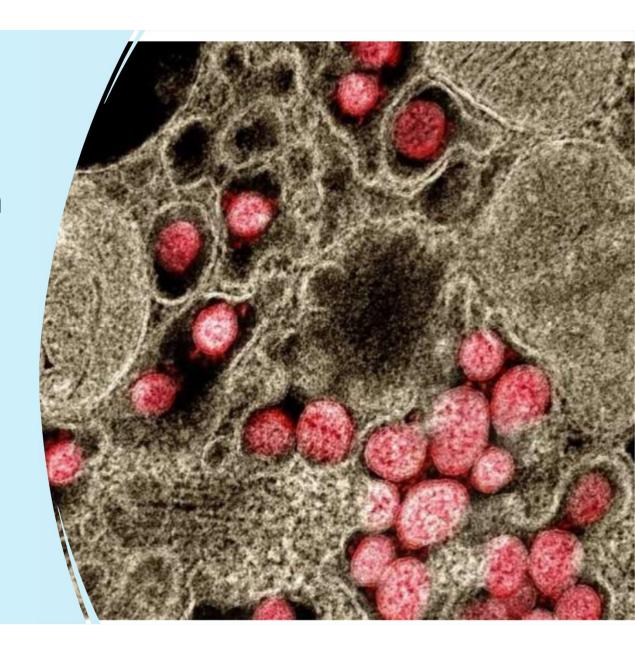
- Initially and even now among some a strong belief it is a somatic perception disorder
- Patients not having their illness affirmed by the health practitioners
- No medical school training and thereby knowledge
- Social services not accepting patients cannot work or need help
- Families not understanding –not a socially acceptable illness

Origins of Long Covid: arising from a specific viral infection

• SARS-CoV-2

(but now from 180 million infections! – it is not boutique!)

- a range of syndromes
 - Post viral fatigue
 - Ongoing lung pathology
 - Ongoing heart pathology
 - Post traumatic syndrome



Attitudes towards Long COVID?

- High degree of attention from the clinicians dealing with the pandemic - were surprised when patients did not recover as expected
- ME/CFS patients initially angry and now sympathetic but hope attention for their illness can coat tail.
- 'Long haulers' have had a strong voice through social media
- Special units set up in the UK to study ongoing the subgroups of 'long hauler' patients
- The NIH has promised over 1 billion dollars to support research into Long COVID
- In New Zealand slow response from MoH given there are debilitated patients

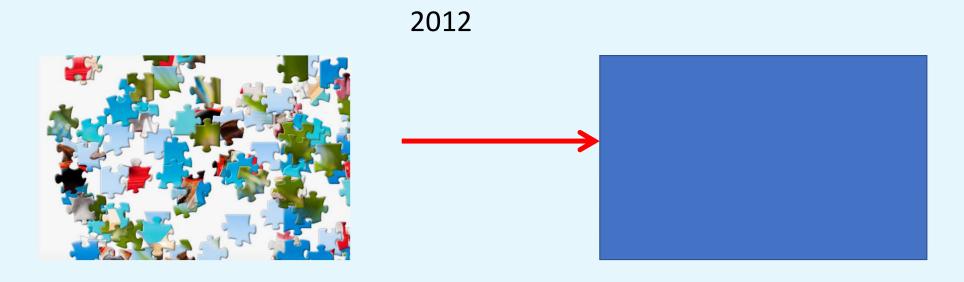
What is an ME/CFS patient like?



Changes in the USA in attitudes to ME/CFS in 2019 "just before the pandemic"

- The US senate passed a bill declaring ME/CFS was a serious illness
- Patients and caregivers had been poorly treated both medically and from social services
- Research support was promised from National Institute of Health – a few million dollars
- Research Centres on ME/CFS were set up at Harvard and Stanford

Solving the jigsaw puzzle of ME/CFS



Lots of interesting individual pieces of research

No integrated picture of the illness

As a parent in an ME/CFS family my research questions were:

Onset: What 'control centre' in the body is affected to cause such severe and diverse long-lasting symptoms?

Perpetuation: Why do the disease symptoms not resolve?

Relapse Cycles: What triggers the frequent relapses?

And then a new question was added on the 25th anniversary of my daughter's illness – aged 39

Physiological State: Why do the symptoms improve so markedly in **pregnancy** – at least for her?



Now a pre-teen on his 6th birthday



How do we get started on research with modest resources? Around 2012!

Approach:

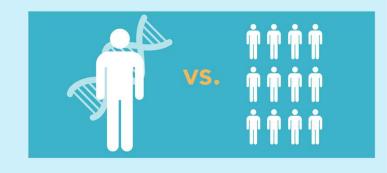
Inspired by 'Personal omics study' of Michael Snyder over 2 years of Snyder's own personal health – multi\$M -40 authors 2012 Cell paper

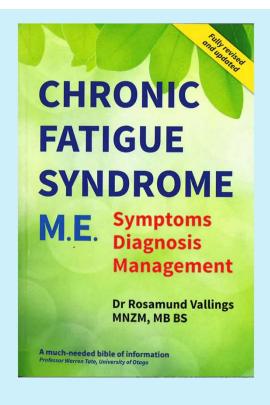
Size of patient studies

Inspired by my former student, Tony Manning, Momenta Pharmaceuticals -study of rare diseases with very few patients

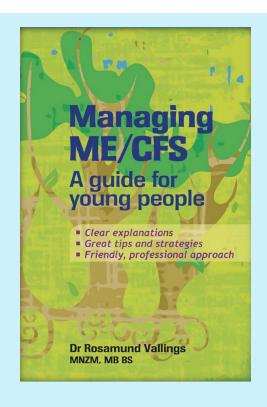
Used precision medicine approach





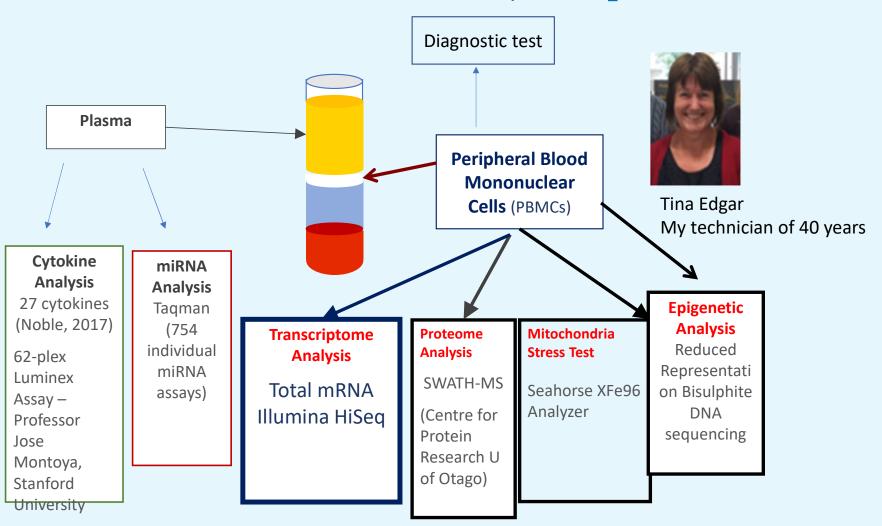






NZ 's expert GP for ME/CFS: Ros Vallings

Our molecular studies with ME/CFS patients



Active genes in immune cells -13000 transcripts

Top 3 changes in ME/CFS involved in inflammation

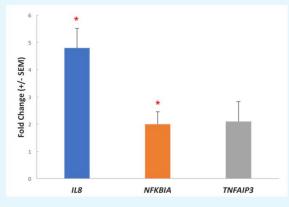
• 27 gene transcripts increased and 6 decreased (P<0.01).

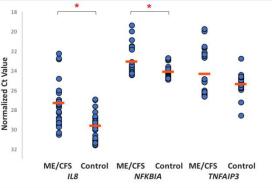
Gene	Name	P -value	Fold Change
IL8	Interleukin-8	2.5x10 ⁻⁶	5.6
NFKBIA	NF-κB Inhibitor Alpha*	2.0x10 ⁻⁵	2.4
TNFAIP3	Tumour Necrosis Factor Alpha-Induced	1.3x10 ⁻⁴	3.6
	Protein 3*		

Eiren Sweetman



Anti-inflammatory responders to TNF- induced NF-κB activation





As a PhD student

Functional Pathways

- Functional Network Analysis (n=33, P<0.01) and Ingenuity Pathway Analysis (n=165, P<0.05) detected effects on:
 - Mitochondrial function (energy production)
 - Cellular stress response/oxidative stress
 - Lowered Metabolism
 - Immune/inflammatory pathways
 - Circadian clock function

Sweetman, E., Ryan, M., Edgar, C., MacKay, A., Vallings, R., & Tate, W. (2019). Changes in the transcriptome of circulating immune cells of a New Zealand cohort with myalgic encephalomyelitis/chronic fatigue syndrome. *International Journal of Immunopathology & Pharmacology, 33,* 1-8

Cellular proteins – immune cell proteome

Mass spectrometry Analysis

- 1. Digested isolated proteins into tiny pieces
- Built a spectral library with small samples from each of all the patients and controls
 (~3000 proteins)

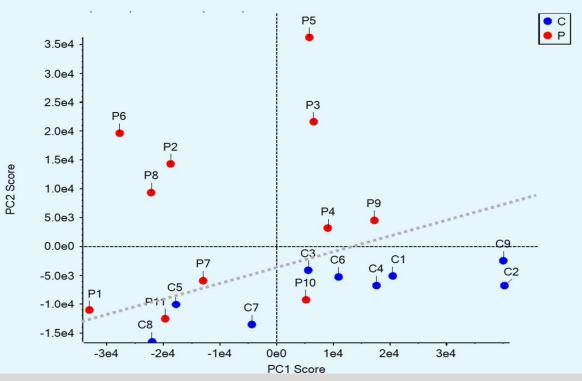
Eiren Sweetman



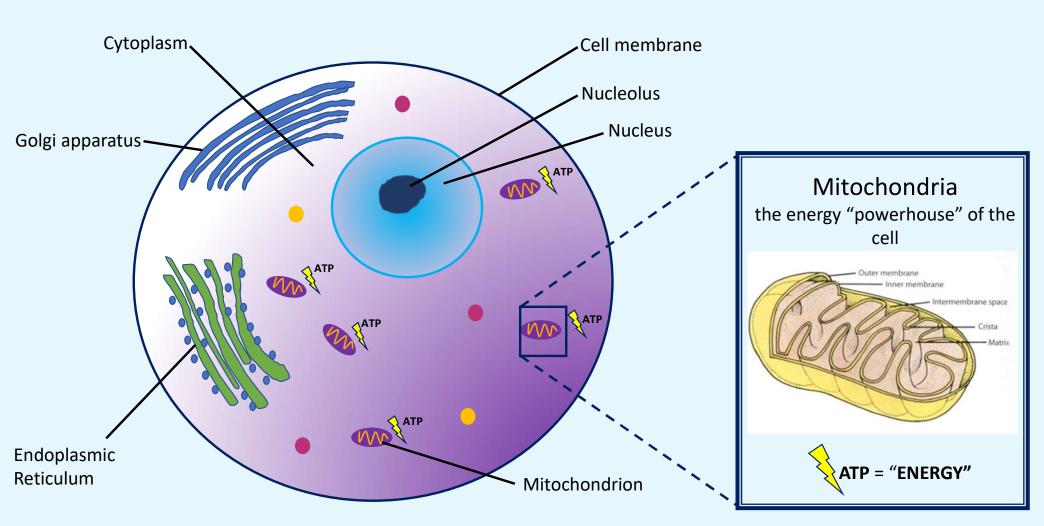
3. Quantify proteins in all samples :Sequential Acquisition of all theoretical Fragment Ion Spectra mass Spectrometry (SWATH-MS)

Principal Component Analysis (PCA)

A principal component analysis (PCA) was used to stratify groups based on protein abundance patterns, which segregated the majority of the ME/CFS patients from the controls.

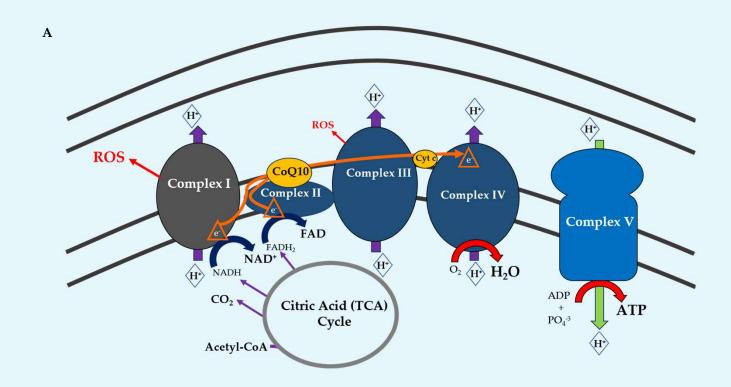


9/11 ME/CFS patients separated from healthy controls (linear regression analysis)



Basic structure of a human cell

Key finding: Many mitochondrial proteins up or down regulated



Sweetman ES, Kleffmann T, Edgar CD, DeLange, M,, Vallings R, &Tate WP (2020) A Swath-MS analysis of Myalgic Encephalomyelitis /Chronic Fatigue Syndrome peripheral blood mononuclear cell proteomes reveals mitochondrial dysfunction *J Translat Med* 18:365

Mitochondrial proteins affected –structural, metabolic pathways, electron chain complexes, ROS response

Mitochondrion

FUN14 domain-containing protein 2 (FUNDC2)

Putative transferase CAF17 (IBA57) Endophilin-B1 isoform 1 (SH3GLB1) BRI3-binding protein (BRIB3P) Biogenesis of lysosome-related organelles complex 1 subunit 1 (BLOC1S1)

Mitochondrial fission 1 protein (FIS1) Glycine--tRNA ligase (GARS)

39S ribosomal protein L12 (MRPL12) 39S ribosomal protein L43 (MRPL43) Peptidyl-prolyl cis-trans isomerase

NIMA-interacting 4 (PIN4) Prohibitin (PHB)

Single-stranded DNA-binding protein (SSBP1)

Dynamin-1-like protein (DNML1) Choline transporter-like protein 1 (SLC44A)

Stimulator of interferon genes protein (TMEM173)

Thiosulfate sulfurtransferase (TST) CCA tRNA nucleotidyltransferase 1

ATP-binding cassette sub-family B member 7 (ABCB7)

39S ribosomal protein L23 (MRPL23)

Citric Acid (TCA) Cycle

D-beta-hydroxybutyrate dehydrogenase (BDH1)

2-oxoglutarate/malate carrier protein (SLC25A11)

Glycerol-3-phosphate dehydrogenase (GPD2) Pyruvate dehydrogenase E1 component subunit beta (PDHB) Isocitrate dehydrogenase [NAD] subunit beta (IDH3B)

Aconitate hydratase(ACO2) 2-oxoglutarate dehydrogenase (OGDH)

Isocitrate dehydrogenase (NADP+) (IDH2)

Mitochondrial import inner membrane translocase subunit Tim8 A isoform 1 (TIMM8A) Glycerol kinase (GK)

Beta-oxidation of fatty acids

Short-chain specific acyl-CoA dehydrogenase (ACADS) Trifunctional enzyme subunit beta, mitochondrial (HADHB)

Complex I

Beta subcomplex subunit 3, accessory subunit (NDUFB3) Alpha subcomplex subunit 3,

accessory subunit (NDUFA3) Beta subcomplex subunit 11, accessory subunit (NDUFB11) Alpha subcomplex subunit 12, accessory subunit (NDUFA12) Alpha subcomplex subunit 5,

ROS stress response

accessory subunit (NDUFA5)

Glyceraldehyde-3-phosphate dehydrogenase, testis-specific (GAPDHS)

Thioredoxin reductase 2 (TXNRD2) Thioredoxin domain-containing protein 17 (TXNDC17)

Peroxiredoxin-1 (PRDX1)

Peroxiredoxin-2 (PRDX2)

Selenoprotein H [Homo sapiens] (SELENOH)

Peptidyl-prolyl cis-trans isomerase F, mitochondrial precursor (PPIF)

Oxidation resistance protein 1 (OXR1)

Peroxiredoxin 6 (PRDX6)

Complex III

Ubiquinone biosynthesis protein COO9 (COO9) Cytochrome C, Somatic (CYCS)

Complex IV

Cytochrome c oxidase assembly factor 6 homolog (COA6)

Complex V

ATP synthase F1 subunit epsilon (ATP5E)

Phosphate carrier protein, mitochondrial isoform b precursor (SLC25A3)

ATP synthase F1 subunit delta (ATP5D)

ATP synthase F1 subunit beta (ATP5F1)

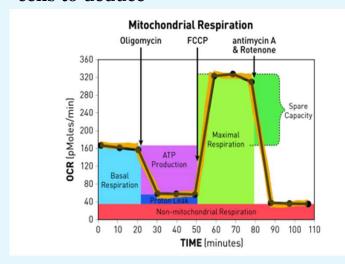
Up-regulated during skeletal muscle growth protein 5 (USMG5)

ATP synthase mitochondrial F1 complex assembly factor 1 (ATPAF1)

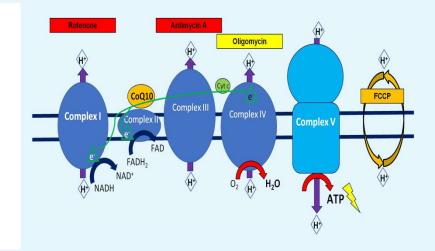
Seahorse analysis of Mitochondria

Mito Stress Test

The test measures mitochondrial function by directly measuring the oxygen consumption rate (OCR) of live cells to deduce

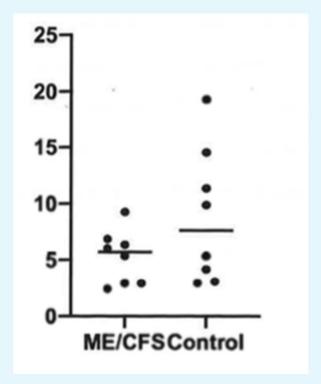


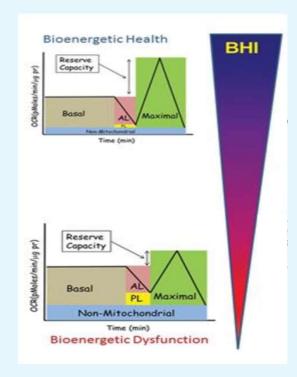
ATP production, Maximal respiration, Non-mitochondrial respiration, Proton leak and spare respiratory capacity

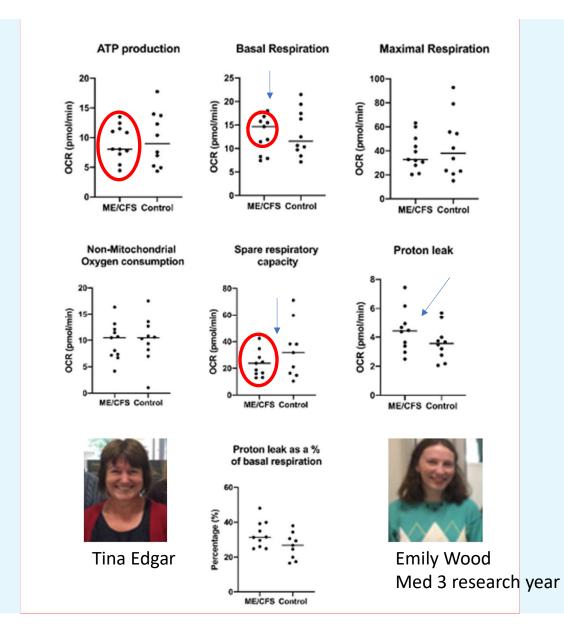


Bioenergetic Health Index

Results suggest a lower BHI in ME/CFS







My big questions as a parent in an ME/CFS family

Control centre for the disease? symptoms, non recovery, relapses

brain

Why not recovery?

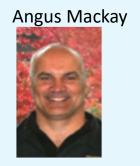
Ongoing stressor

Explanation

neuroinflammation?

A neuroinflammatory paradigm to explain symptoms, lack of recovery, and relapses in ME/CFS

Focused on the hypothalamus and the 'stress centre of the brain' – a cluster of neurons within the paraventricular nucleus



Paradigm: "Cycles of fluctuating chronic neuro-inflammation can drive stress related relapses in ME/CFS and support the failure to recover the 'normal' homeostatic set point that would allow recovery"

We think damaged mitochondria have a role as the danger signal

Mackay, A & Tate WP (2018) A compromised paraventricular nucleus within a dysfunctional hypothalamus: A novel neuroinflammatory paradigm for ME/CFS Internat J Immunopath and Pharmacol 2: 1–8

Changes in epigenetic code of the DNA associated with ME/CFS

Changes during relapse recovery cycle

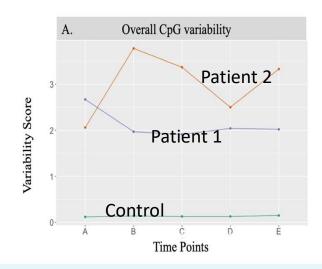
Amber Helliwell

A. Summary of self reported health scores (A-E)

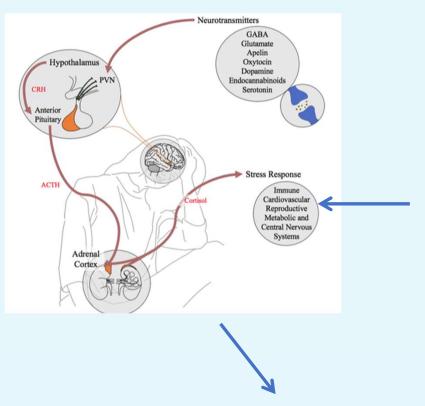
	A	В	C	D	E
Patient One	"Well"	"Fragile"	"Fragile"	"Satisfactory"	"Satisfactory"
	7	-3	-2	7	7
Patient Two	"Fragile" 5	"Fragile" 6	"Relapsed" 2	"Satisfactory" 6	"Fragile" 4
Control	"Well"	"Well"	"Well"	"Well"	"Well"
	10	10	10	10	10

Generated reduced genome DNA methylation maps By reduced representation bisulphite sequencing Aniruddha Chatterjee (Pathology)

Patient epigenetic methylome code was 20 fold more dynamic than the control



Hypothalamus/Pituitary/adrenal (HPA) axis with neurotransmitter pathways identified as affected in ME/CFS



Effects on neurotransmitters identified by five hypomethylated gene regions:

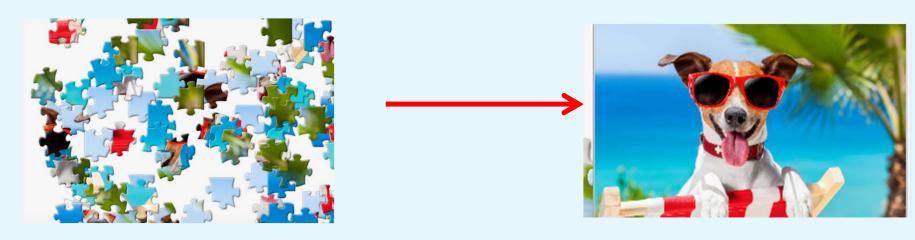
RYR1, GNAS, GNG7, GABRB3 and APBA2

Low cortisol associated with fatigue and Post Exertional Malaise



Solving the jigsaw puzzle of ME/CFS

2021



Lots of interesting individual pieces of research

An integrated picture of the disturbed physiology In ME/CFS

What now in 2021 & 2022?

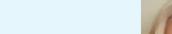
- A. The most extensive study yet of **Post Exertional Malaise** a core symptom of ME/CFS
- (i) Cardiac Physiology (ii) DNA methylome (iii) mitochondrial function
- (iv) Oxidative stress biomarkers in Protein and DNA, glutathione oxidation status





B. Long COVID – a comparative study with ME/CFS does it have the same molecular signatures?
(i) Immune cell proteome (ii) DNA methylome

(iii) Oxidative stress



Tina Edgar Anna Blair

r Planned

C. Long COVID immune cellular and molecular signatures before and after vaccination (with Anna Brooks - specialist immunologist UoA)

(i) Serology (ii) Specific immune cell dysfunctions (iii) molecular signatures



Jemma Elley

Dr Anna Brooks

Acknowledgements

Collaborators

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Michael de Lange — Biostatistics
Margaret Ryan — Dept of Anatomy
Peter Stockwell — Dept of Pathology
Lynette Hodges , School of health and exercise Massey U
Anna Brooks — School of Biological Sciences UoA

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We also wish to express our sincere gratitude towards all of our ME/CFS study participants.

Collegial Discussion

Paul Fischer – Monash U, Vic Jose Montoya – Stanford, Calif Paul Naviaux – Stanford, Calif Sonya Marshall-Gradnisek Griffiths U, Que







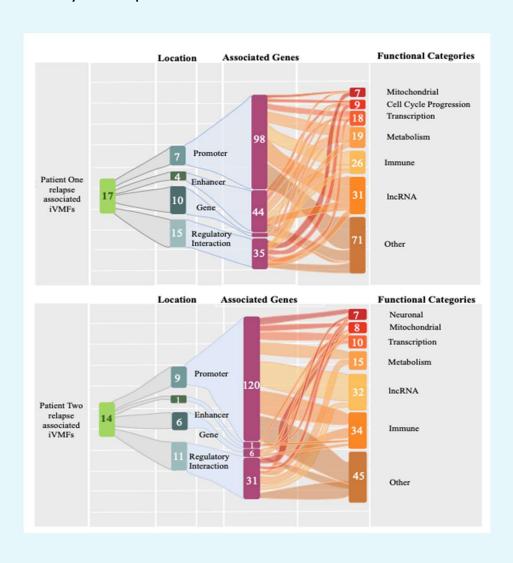


DNA methylation changes at specific sites

chr13:114918456-114918525 chr19:33885306-33885381 (PEPD) chr19:55464080-55464189 (NLRP7) A. 100 75 50 Wethylation % 100 chr8:145003618-145003684 (PLEC) chrX:149106531-149106576 (CXorf40B 75 50 25 Patient One B. Time Point chr19:4543716-4543762 chr22:18027985-18028072 chr1:155098923-155098964 100 75 Methylation % 25 chrX:15353393-15353393 100 75 50 25

Time Point

Summary of relapse associated features and functions



First Study Group

Clinical characteristics	ME/CFS participants	Control participants	
Number	11	9	
Median age (years)	43.2	38.0	
Age range (years)	11.3-69	12.5-60	
Gender	F = 7	F=6	
	M = 4	M=3	
Median BMI	23.7	b	
Nationality	NZ/European	NZ/European	
Median illness duration (years)	11	N/A	
Stage of illness ^a	Acute = 2 Chronic = 9	N/A	
Potential initial ME/CFS 'trigger'	Acute infection = 7 Surgery = 1 Stress = 1 Unsure = 2	N/A	

All of our ME/CFS participants were diagnosed by Dr Vallings using the 2003 Canadian Consensus Criteria

b Data not collected