

Bezisterim Effects on Biological Age, Alzheimer's Epigenetics, and Neurologic Assessments

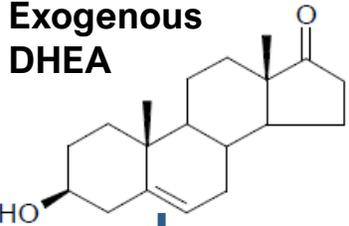
Christopher Reading, PhD
BioVie Inc.
ARDD Copenhagen, 2024

In 2000, we asked

“Why does DHEA show

- anti-inflammatory
- insulin-sensitizing
- neuroprotective
- anti-aging effects

in mice, and not in humans?”



INFLAMMATION

MICE

MEN

✓ **ACTIVE**

✗ **INACTIVE**

- NFκβ
- cytokines
- chemokines
- IRS1/2 p-Serine
- insulin resistance

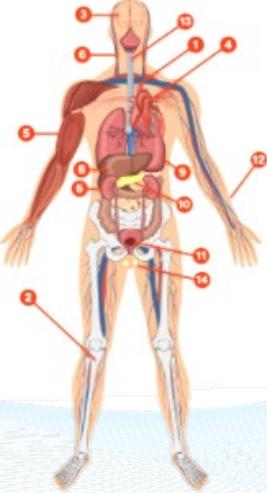


- amyloid b
- cognitive decline
- Alzheimer’s
- neuron death

- hypertriglyceridemia
- adipogenesis
- obesity
- diabetes
- longevity



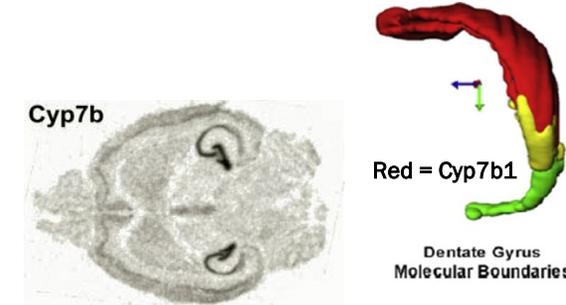
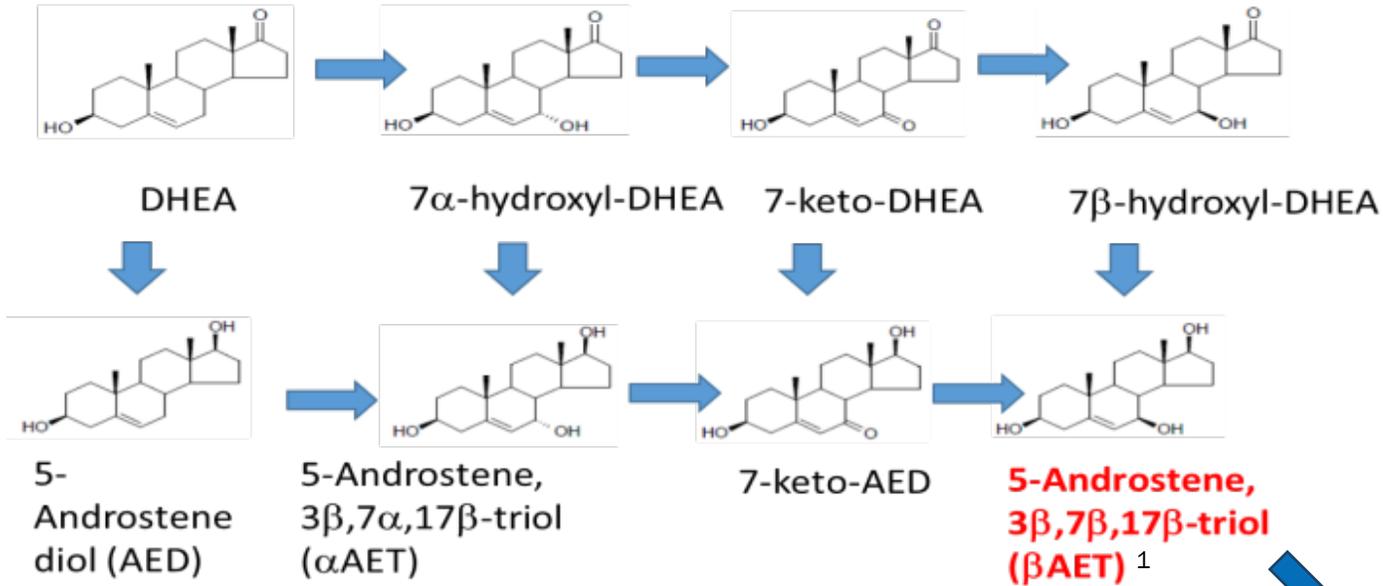
Harvard Health Publishing
HARVARD MEDICAL SCHOOL



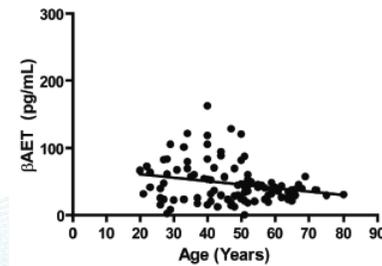
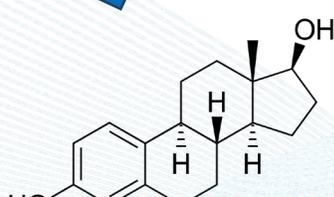
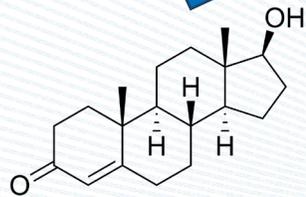
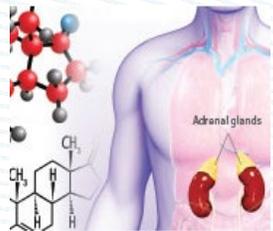
- insulin resistance
- hyperglycemia
- obesity
- dyslipidemia
- osteoporosis
- cognitive decline
- Alzheimer’s
- decreased quality of life

Thousands of publications

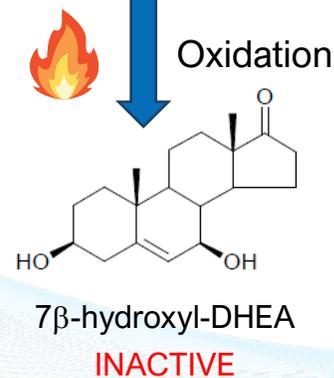
Bezisterim Developed by Understanding the Differential Metabolism of Androst-5-ene-3,17-diol Between Rats, Canines, Monkeys and Humans¹



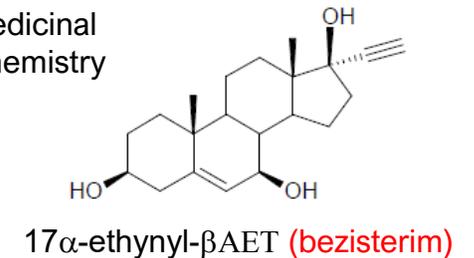
Neurosteroid: Cyp7B defines hippocampal dentate gyrus^{3,4}



β AET decreases with age²



Medicinal Chemistry



Metabolic stability
Oral bioavailability
Blood-brain permeable

Bezisterim

In vitro, in vivo and clinical findings align with expected pharmacology

- Binds extracellular-signal regulated kinase⁵, (ERK), and inhibits inflammatory^{6,7}, but not homeostatic⁸ ERK, NFκB and TNF signaling. It is more potent than βAET for inhibition of NFκB transcription
- These activities have been demonstrated in primary **macrophages**, and in **monocyte/macrophage cell lines**, as well as in vivo animal models
- Decreased inflammatory mediators and insulin resistance in animal models and human subjects with obese impaired glucose tolerance or T2D^{9,10}
- Improved motor activity and decreased neurodegeneration in Parkinson's disease models¹¹, and improved motor activity in a phase 2a PD study¹²
- Has a well-tolerated safety profile to date¹³
- Positive signals of clinical activity in dementia in AD clinical studies

Positive Signals of Bezisterim Clinical Activity in Dementia

Open label 14-week study in MCI and Mild AD

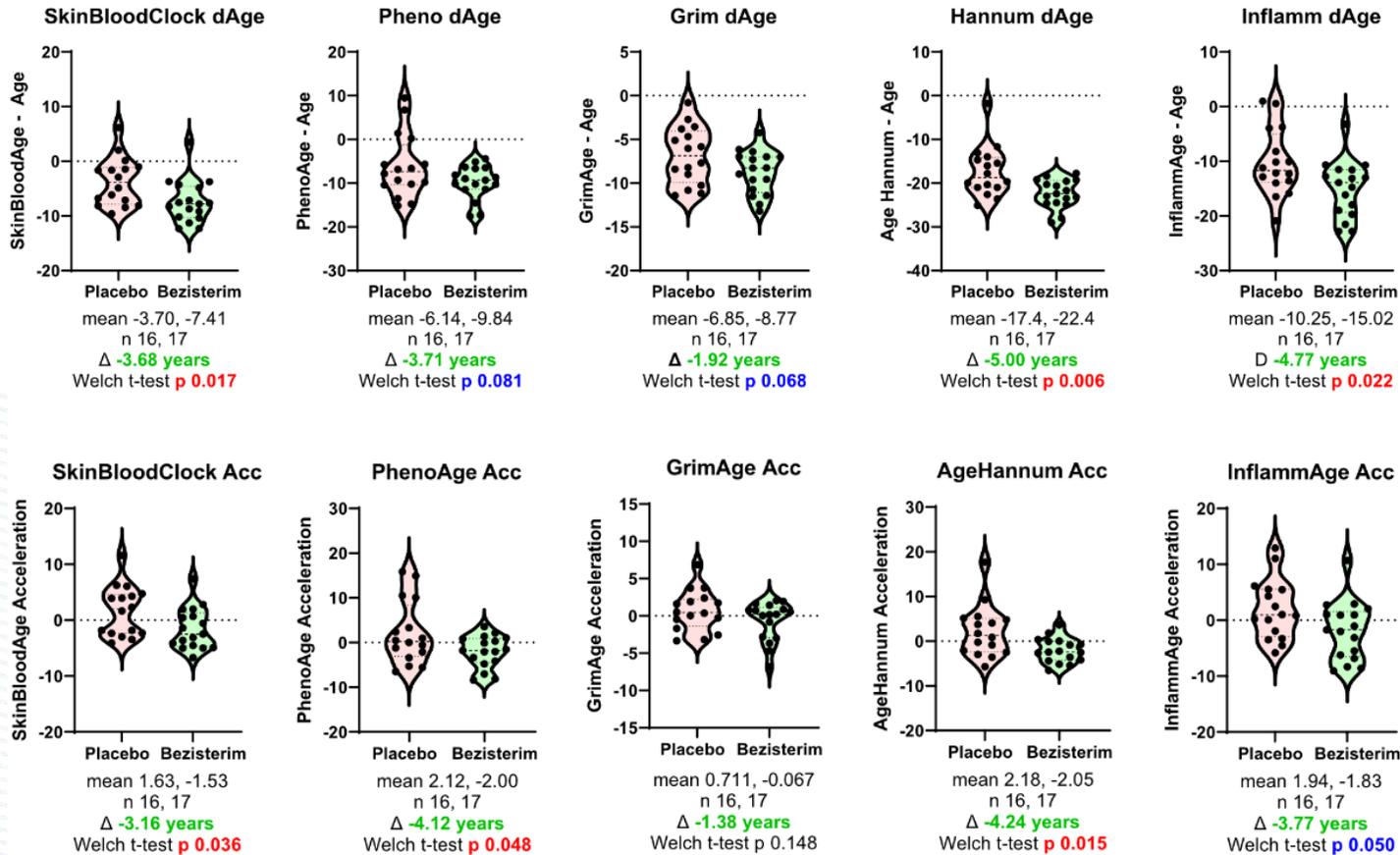
- Improved neurological, neuroimaging, and dementia and inflammation biomarkers in an open-label 14-week study in MCI and mild AD¹⁴
- Observed 3.4-year decrease from baseline in biological age (Horvath SkinBlood Clock), and altered monocyte methylome¹⁵, but array batch effects confounded the interpretation

Placebo-controlled 30-week study in Mild/Mod AD^{16,17}

- Positive signals found in -
 - Neurological and biomarker measures in 50 per-protocol subjects (26 placebo and 24 bezisterim)
 - DNA methylation data for 33 subjects at completion (16 placebo and 17 bezisterim)

Focus of this presentation

Bezisterim Subjects Showed a Lower Biological Age Than Placebo Subjects at 30 Weeks



Difference between biologic and chronological age (dAge) as calculated by different epigenetic aging clocks¹⁸⁻²²
 (SkinBlood Clock difference consistent with previous study¹⁵)

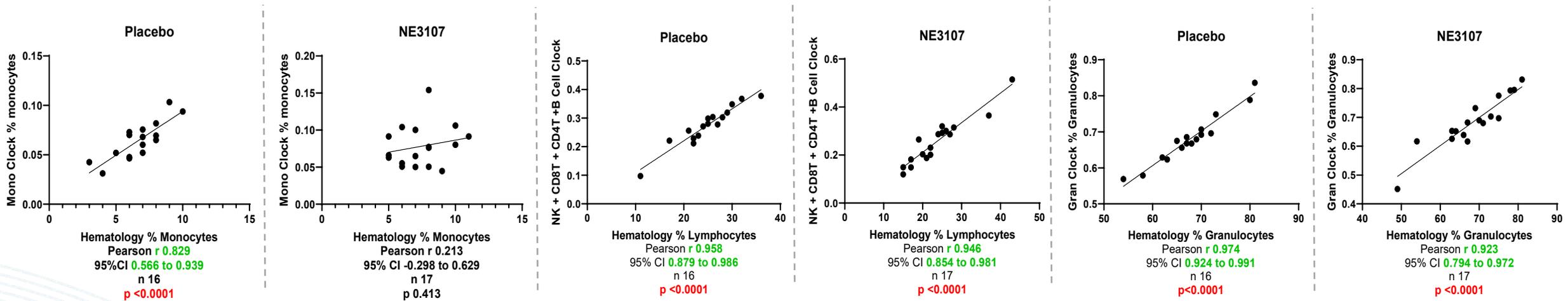
A consistent trend was observed when using age acceleration (similar to dAge but not confounded by chronological age)
 Note consistent placebo acceleration, Bezisterim deceleration

Raw data from Epigenetic Clock Development Foundation; Analyses of diverse epigenetic aging clocks performed by Chronomics/Hurdle Group, including their new InflammAge Clock²²

18. Horvath 2018 *Aging* 10 1758; 19. Levine 2018 *Aging* 2018 10 573; 20. Lu 2019 *Aging*. 2019 11 303; 21. Hannum 2013 *Mol Cell* 49 359; 22. Schunk *bioRxiv*. 2023. doi.org/10.1101/2023.12.21.572866

Bezisterim Appeared to Modify Week 30 Monocyte Methylome

Consistent with hypothesis of a transition from pro- to anti-inflammatory



Cell Type	Placebo Pearson r, P	Bezisterim Pearson r, P	Fisher transformation Z test statistic, P
Monocytes	0.829, <0.0001	0.213, 0.413	2.51, 0.012
Lymphocytes	0.958, <0.0001	0.946, <0.0001	0.334, 0.738
Granulocytes	0.974, <0.0001	0.923, <0.0001	1.44, 0.149

- Hematology and clock data are correlated for lymphocytes and granulocytes
- For placebo, but not bezisterim subjects, monocytes are not correlated
- Consistent with the previous open-label study result and hypothesis of a transition from **pro-inflammatory (M1) to anti-inflammatory (M2) myeloid cells**¹⁵

Analysis of Promoter Methylation

- Increased promoter methylation has been associated with decreased gene expression, however gene expression was not measured in this study
- Monocytes represent less than 10% of leukocytes; if significant promoter methylation differences were predominantly represented in monocytes²³, the magnitude of changes is much higher in monocytes.
- Significant (FDR $p < 0.05$) differences in promoter CpG methylation β values (bezisterim – placebo) identified were almost exclusively increased methylation by bezisterim

Bezisterim Increases in Promoter Methylation of Genes Associated with Disease Progression, Inflammation and Metabolic Regulation

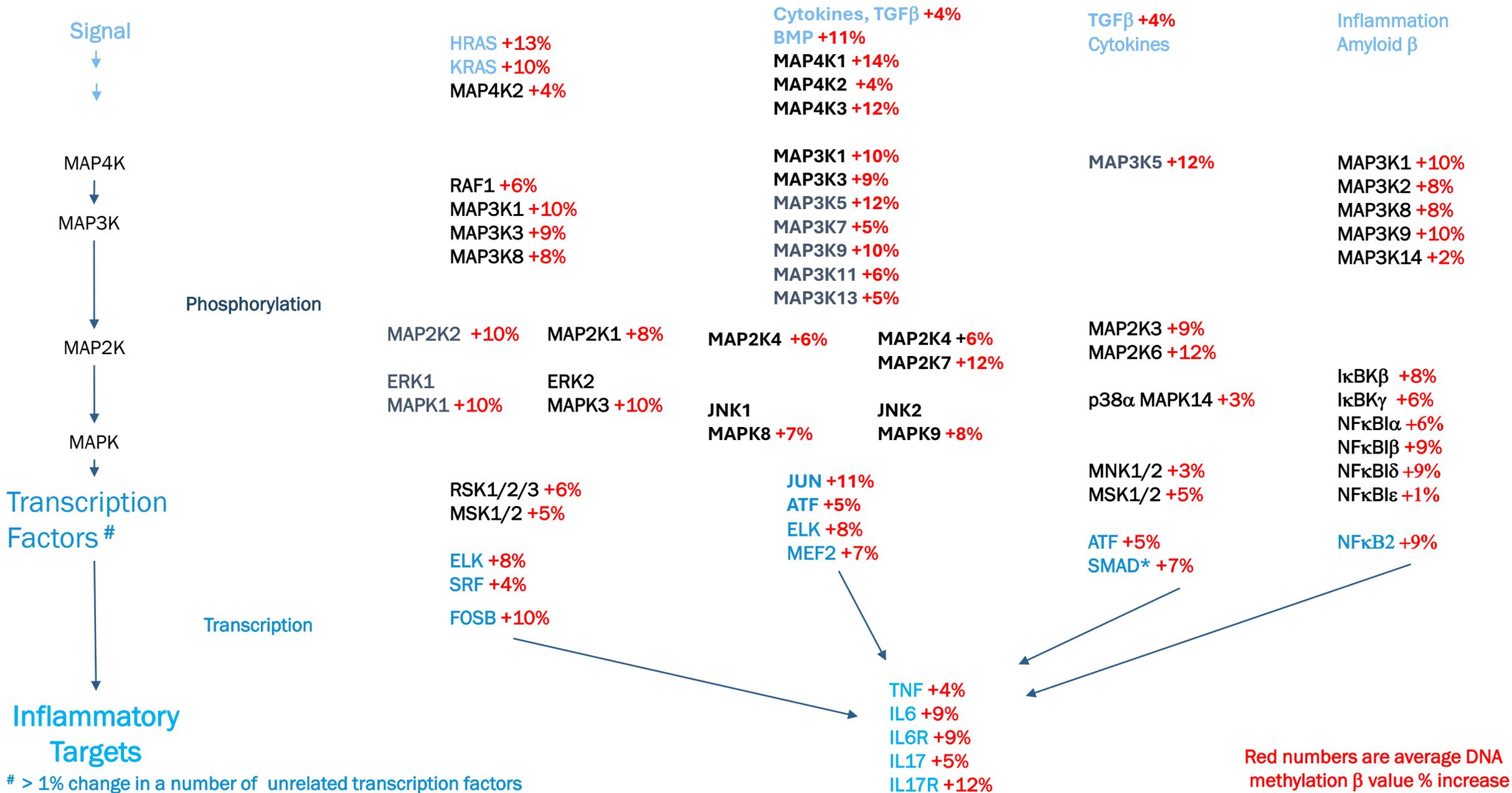
- Compared to placebo, **bezisterim** subjects had significantly (FDR $p < 0.05$) increased methylation in promoter CpGs of 4934 genes
- The top 50 genes (FDR $p < 0.005$) genes identified **associations with disease progression** for the following:
 - Cognitive decline, AD, PD and other CNS diseases
 - Inflammatory response, neurodegeneration, cell death
 - Obesity, insulin resistance, and T2D
 - Neurodegeneration biomarkers
 - Age acceleration
- Compared to placebo, **bezisterim** subjects had significantly (FDR $p < 0.05$) **decreased methylation** in promoter CpGs in 21 genes
- We identified these 6 of the 21 genes were associated with the following **improved homeostasis**:
 - Autophagy, mitochondrial energy
 - Positive regulation of transcription
 - Repression of inflammatory gene expression, decreased ROS
 - Promotion of beta cell regeneration

Key Genes in the AD Pathway for Promoter Methylation Increased in Bezisterim Compared to Placebo Subjects

Activity may reduce gene expression

- ADAM10
 - α -Secretase, decreases A β
- APP
 - Decreases A β
- Bid
 - Apoptosis death domain agonist
- ATF6
 - ER stress response decreased in Alzheimer's disease
- CASP7
 - Caspase-cleaved **tau** catalyzes filament formation, adopts a conformation found in early-stage tangles, and can be hyperphosphorylated
- TNF
 - Pro-inflammatory cytokine, NF κ B activation, decreases LTP

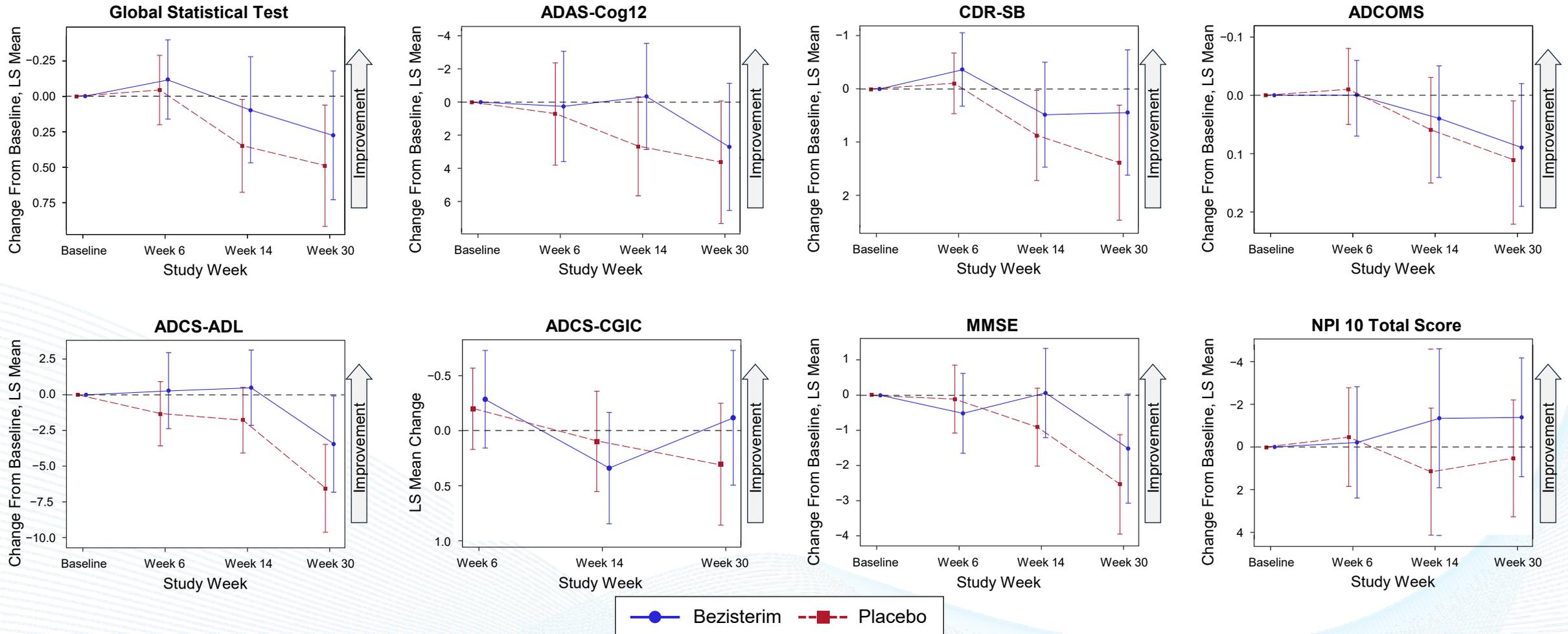
Inflammatory Gene Expression Decreased by Compounded Sequential Kinase Increased Promoter Methylation with Bezisterim vs Placebo



Bezisterim KEGG Pathways Enrichment

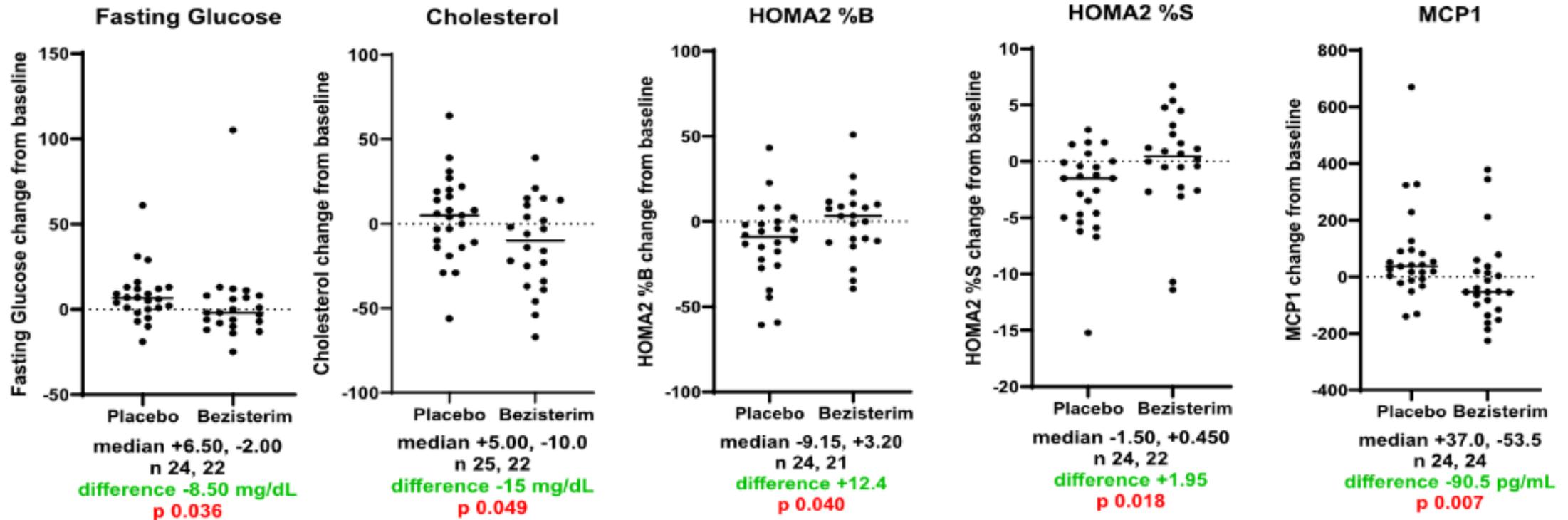
Pathway	Fold Enrichment	FDR Adjusted P	Genes/pathway Genes
Alzheimer's disease	1.6	6.2e ⁻⁶	110/384
Amyotrophic lateral sclerosis	1.7	1.8e ⁻⁷	112/364
Neurodegeneration multiple diseases	1.6	1.7e ⁻⁶	134/476
Parkinson's disease	1.7	7.0e ⁻⁶	82/266
Huntington's disease	1.6	1.4e ⁻⁴	86/306
Cellular senescence	1.8	1.3e ⁻⁴	51/156
Apoptosis	1.8	2.5e ⁻⁴	45/136
Proteasome	2.3	1.8e ⁻³	19/46
TNF signaling	1.5	2.9e ⁻²	31/112
NFκB signaling	1.6	3.4e ⁻²	29/104

Bezisterim Showed Improvements Over Time in Primary and Secondary Endpoints



- **Primary and secondary efficacy endpoints.** Scores for bezisterim and placebo from baseline through week 30. Magnitude of responses was comparable to results reported for approved medications lecanemab²⁴ and aducanumab²⁵

Directional Improvements From Baseline for Metabolic and Inflammatory Biomarkers with Bezisterim



There were non-significant directional improvements in fasting insulin, HOMA2 insulin resistance, triglycerides, fructosamine, adiponectin, leptin, systolic and diastolic blood pressure, and weight (data not shown) with bezisterim. There were no significant differences in pTau, GFAP, NfL or A β 42/40 ratio.

Biomarker Correlations With Neurological Assessments

- Changes from baseline in biomarkers in placebo, but not bezisterim subjects correlated with neurological assessment decline
 - For example, GFAP, pTau 217 and CRP
- Directions of improvement in scales
 - Increased MMSE and ADL = improvement
 - Decreased GST, CDR-SB, CGIC, & ADCOMS = improvement

Measure	Biomarker	Placebo		Bezisterim	
		Pearson R	P<0.1	Pearson R	P<0.1
GST <small>Global Statistical Test</small>	GFAP	0.671	1.7e ⁻⁴	0.253	
	pTau	0.444	0.023	0.174	
	Systolic BP	0.431	0.028	0.313	
	Cholesterol	-0.060		0.495	0.014
	NfL	0.169		0.480	0.018
CDR-SB <small>Clinical Demential Rating</small>	RANTES	0.165		-0.567	0.004
	SBC dAge	0.209		0.473	0.055
	GFAP	0.511	0.008	0.272	
	RANTES	0.398	0.044	-0.445	0.004
	Systolic BP	0.435	0.026	0.388	
Cog12 <small>ADAS-Cognition</small>	GFAP	0.589	0.002	0.272	
	pTau	0.410	0.037	0.133	
	Cholesterol	-0.082		0.550	0.005
	NfL	0.083		0.458	0.024
	RANTES	-0.033		-0.637	0.001
MMSE <small>Mini-Mental State Exam</small>	SBC dAge	0.167		0.455	0.067
	Cholesterol	0.020		-0.468	0.021
	RANTES	0.003		0.516	0.010
	SBC dAge	-0.011		-0.580	0.015
	C1q	-0.395	0.046	0.294	
CGIC <small>Clinician Global Impression of change</small>	pTau	0.414	0.036	0.208	
	SBC dAge	0.017		0.467	0.059
	CRP	-0.458	0.019	0.322	
ADL <small>Activities of Daily Living</small>	Fructosamine	0.391	0.048	-0.459	0.024
	HOMA2 %B	-0.395	0.046	-0.014	
	NfL	-0.039		-0.642	0.001
	GFAP	0.497	0.010	0.294	
ADCOMS <small>AD Composite Score</small>	pTau 217	0.393	0.047	0.169	
	Systolic BP	0.525	0.006	0.370	
	NfL	0.120		0.454	0.026
	RANTES	0.227		-0.575	0.003
	SBC dAge	0.134		0.469	0.058

Biomarker Correlations With Neurological Assessments

Measure	Biomarker	Placebo		Bezisterim		
		Pearson R	P<0.1	Pearson R	P<0.1	
GST	GFAP	0.671	1.7e ⁻⁴	0.253		
	pTau	0.444	0.023	0.174		
	Systolic BP	0.431	0.028	0.313		
	Cholesterol	-0.060		0.495	0.014	
	NfL	0.169		0.480	0.018	
	RANTES	0.165		-0.567	0.004	
CDR-SB	SBC dAge	0.209		0.473	0.055	
	GFAP	0.511	0.008	0.272		
	RANTES	0.398	0.044	-0.445	0.004	
	Systolic BP	0.435	0.026	0.388		
	Cog12	GFAP	0.589	0.002	0.272	
		pTau	0.410	0.037	0.133	
Cholesterol		-0.082		0.550	0.005	
NfL		0.083		0.458	0.024	
RANTES		-0.033		-0.637	0.001	
SBC dAge		0.167		0.455	0.067	
MMSE	Cholesterol	0.020		-0.468	0.021	
	RANTES	0.003		0.516	0.010	
	SBC dAge	-0.011		-0.580	0.015	
CGIC	C1q	-0.395	0.046	0.294		
	pTau	0.414	0.036	0.208		
	SBC dAge	0.017		0.467	0.059	
ADL	CRP	-0.458	0.019	0.322		
	Fructosamine	0.391	0.048	-0.459	0.024	
	HOMA2 %B	-0.395	0.046	-0.014		
	NfL	-0.039		-0.642	0.001	
ADCOMS	GFAP	0.497	0.010	0.294		
	pTau 217	0.393	0.047	0.169		
	Systolic BP	0.525	0.006	0.370		
	NfL	0.120		0.454	0.026	
	RANTES	0.227		-0.575	0.003	
	SBC dAge	0.134		0.469	0.058	

- Changes from baseline in biomarkers in bezisterim but not placebo subjects correlated with neurological assessment improvement
 - For example, decreases in neurofilament light chain (NfL) and cholesterol
- Directions of improvement in scales
 - Increased MMSE and ADL = improvement
 - Decreased GST, CDR-SB, CGIC, & ADCOMS = improvement

Biomarker Correlations With Neurological Assessments

Measure	Biomarker	Placebo		Bezisterim	
		Pearson R	P<0.1	Pearson R	P<0.1
GST	GFAP	0.671	1.7e ⁻⁴	0.253	
	pTau	0.444	0.023	0.174	
	Systolic BP	0.431	0.028	0.313	
	Cholesterol	-0.060		0.495	0.014
	NfL	0.169		0.480	0.018
	RANTES	0.165		-0.567	0.004
	SBC dAge	0.209		0.473	0.055
CDR-SB	GFAP	0.511	0.008	0.272	
	RANTES	0.398	0.044	-0.445	0.004
	Systolic BP	0.435	0.026	0.388	
Cog12	GFAP	0.589	0.002	0.272	
	pTau	0.410	0.037	0.133	
	Cholesterol	-0.082		0.550	0.005
	NfL	0.083		0.458	0.024
	RANTES	-0.033		-0.637	0.001
	SBC dAge	0.167		0.455	0.067
MMSE	Cholesterol	0.020		-0.468	0.021
	RANTES	0.003		0.516	0.010
	SBC dAge	-0.011		-0.580	0.015
CGIC	C1q	-0.395	0.046	0.294	
	pTau	0.414	0.036	0.208	
	SBC dAge	0.017		0.467	0.059
ADL	CRP	-0.458	0.019	0.322	
	Fructosamine	0.391	0.048	-0.459	0.024
	HOMA2 %B	-0.395	0.046	-0.014	
	NfL	-0.039		-0.642	0.001
ADCOMS	GFAP	0.497	0.010	0.294	
	pTau 217	0.393	0.047	0.169	
	Systolic BP	0.525	0.006	0.370	
	NfL	0.120		0.454	0.026
	RANTES	0.227		-0.575	0.003
	SBC dAge	0.134		0.469	0.058

- The chemokine RANTES can alter astrocyte driven microglial M1>M2 transition
 - For placebo subjects, increased RANTES change from baseline was associated with decline in Clinical Dementia Rating
 - For bezisterim subjects, increased RANTES was associated with improvements in neurological assessments
- Directions of improvement in scales
 - Increased MMSE and ADL = improvement
 - Decreased GST, CDR-SB, CGIC, & ADCOMS = improvement

Biomarker Correlations With Neurological Assessments

- SkinBloodClock (SBC) dAge at completion correlated with neurological improvements in bezisterim subjects. There were no correlations in placebo subjects.
- Directions of improvement in scales
 - Increased MMSE and ADL = improvement
 - Decreased GST, CDR-SB, CGIC, & ADCOMS = improvement

Measure	Biomarker	Placebo		Bezisterim	
		Pearson R	P<0.1	Pearson R	P<0.1
GST	GFAP	0.671	1.7e-4	0.253	
	pTau	0.444	0.023	0.174	
	Systolic BP	0.431	0.028	0.313	
	Cholesterol	-0.060		0.495	0.014
	NfL	0.169		0.480	0.018
	RANTES	0.165		-0.567	0.004
	SBC dAge	0.209		0.473	0.055
CDR-SB	GFAP	0.511	0.008	0.272	
	RANTES	0.398	0.044	-0.445	0.004
	Systolic BP	0.435	0.026	0.388	
Cog12	GFAP	0.589	0.002	0.272	
	pTau	0.410	0.037	0.133	
	Cholesterol	-0.082		0.550	0.005
	NfL	0.083		0.458	0.024
	RANTES	-0.033		-0.637	0.001
	SBC dAge	0.167		0.455	0.067
MMSE	Cholesterol	0.020		-0.468	0.021
	RANTES	0.003		0.516	0.010
	SBC dAge	-0.011		-0.580	0.015
CGIC	C1q	-0.395	0.046	0.294	
	pTau	0.414	0.036	0.208	
	SBC dAge	0.017		0.467	0.059
ADL	CRP	-0.458	0.019	0.322	
	Fructosamine	0.391	0.048	-0.459	0.024
	HOMA2 %B	-0.395	0.046	-0.014	
	NfL	-0.039		-0.642	0.001
ADCOMS	GFAP	0.497	0.010	0.294	
	pTau 217	0.393	0.047	0.169	
	Systolic BP	0.525	0.006	0.370	
	NfL	0.120		0.454	0.026
	RANTES	0.227		-0.575	0.003
	SBC dAge	0.134		0.469	0.058

Bezisterim Increased Methylation of Gene Promoters may be Associated with Clinical Improvements

- We identified 18 significant (FDR $p < 0.05$) **bezisterim** subject clinical measure changes from baseline correlations with individual CpGs of genes that may be **beneficial** for AD subjects
- There were 6 significant (FDR $p < 0.05$) **placebo** subject clinical measure changes from baseline correlations with CpGs of metabolic, inflammatory and dementia biomarker genes that may be **detrimental** for AD subjects
- The data suggest that bezisterim might be altering methylation of AD related genes in concert with **improvements in cognition and function**

Bezisterim Summary

- Analog of the neurosteroid β AET and may recapitulate, in humans, the beneficial effects on diseases of aging seen with DHEA in rodents
- Decreases inflammatory signal transduction and restores insulin sensitivity
- Well-tolerated safety profile to date
- Decreases biological age acceleration
- Bezisterim may act predominantly in myeloid innate immune cells (monocytes, macrophages, astrocytes, microglia)
- May act through RANTES transition of macrophage and microglia (M1) inflammatory to (M2) restorative programs through modification of the monocyte DNA methylome
- Bezisterim subjects had higher DNAm of genes enriched in neurodegeneration pathways
- Alters correlations with inflammatory metabolic disease, and biologic age biomarkers
- May improve Parkinson's disease motoric activity and Alzheimer's disease metabolic inflammation, cognition, function, imaging and biomarkers in clinical studies
- Bezisterim may improve systems dysregulation to restore homeostasis through expression changes of key neurodegeneration gene products
- This may be the first study to demonstrate improvement in longevity and clinical benefit in a disease of aging

Collaborators



Jiayan Yan
Clarence Ahlem
Penelope Markham
Jeffrey Zhang
Harvey Yuan
Lixia Wang
Stephen O'Quinn
Joseph Palumbo



Juozas Gordevicius
Bobby Brooke



HARVARD MEDICAL SCHOOL AND
BRIGHAM AND WOMEN'S HOSPITAL

Donald Simonson, MD, MPH, ScD



Marcia Testa, MPH, PhD

HURDLE

Lisa Schmunk
Hira Javaid
Dani Martin-Herranz