

A New Genetic Test for Alzheimer's Disease and Dementia Risk – with transformational potential

A Polygenic Risk Score test for Alzheimer's Disease

variaTECTTM and SNPfitRTM provide an accurate genetic blood test for the assessment of Alzheimer's Disease risk



Best in class genotyping platform with one software solution offering two analysis options for amyloid PET positivity:-

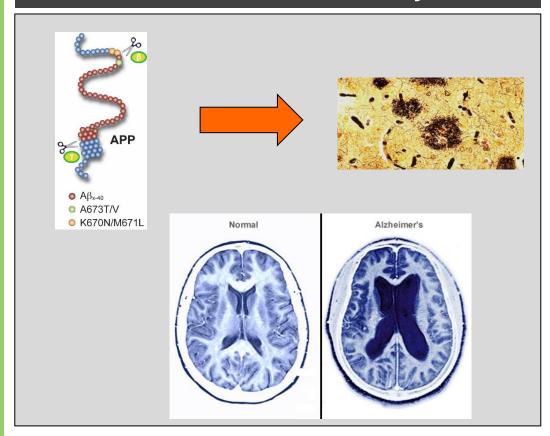
- 1. Assist Pharma in stratifying trial subjects
- 2. Support academic research
- 3. Clinical test for physicians



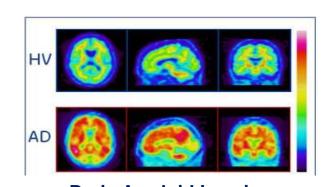


Biomarkers are crucial to the early identification of Alzheimer's Disease

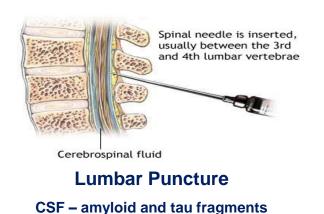
"Genetic analysis is the most effective way of deciding who should be assessed for early disease" – Professor John Hardy Detection of early changes is key:
Working at the molecular level



- PET Imaging: Very expensive and limited availability
- Lumbar Puncture: Invasive and high risk; limited reproducibility

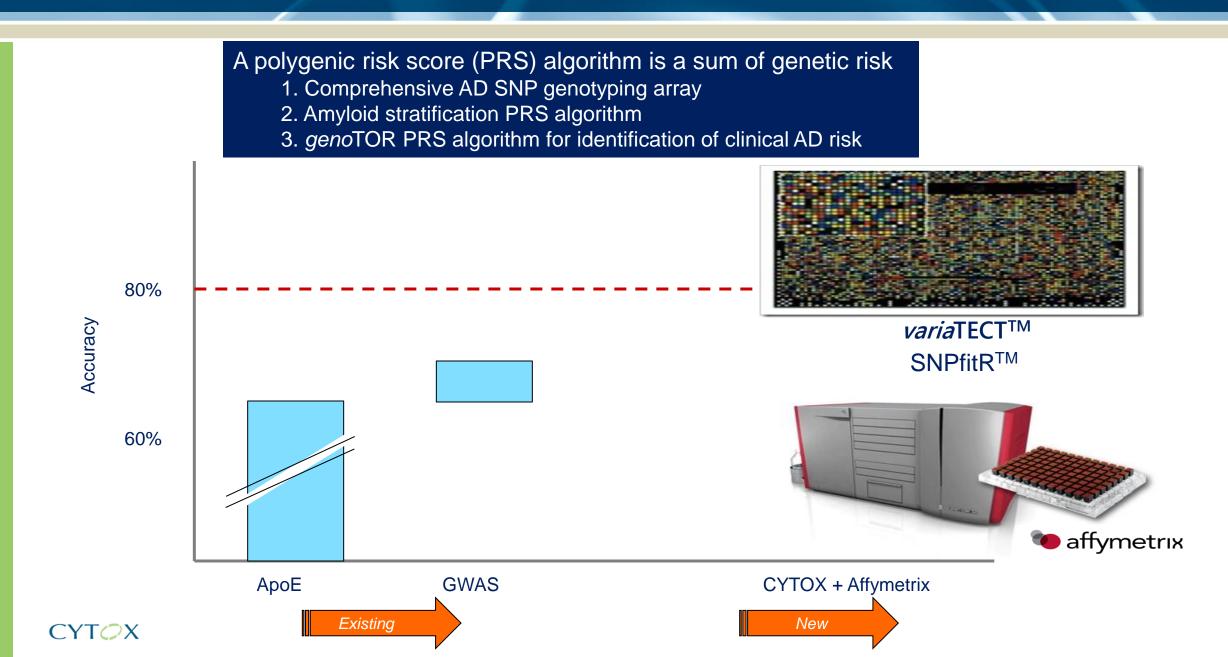


Brain Amyloid Imaging
PET – Positron Emission Tomography



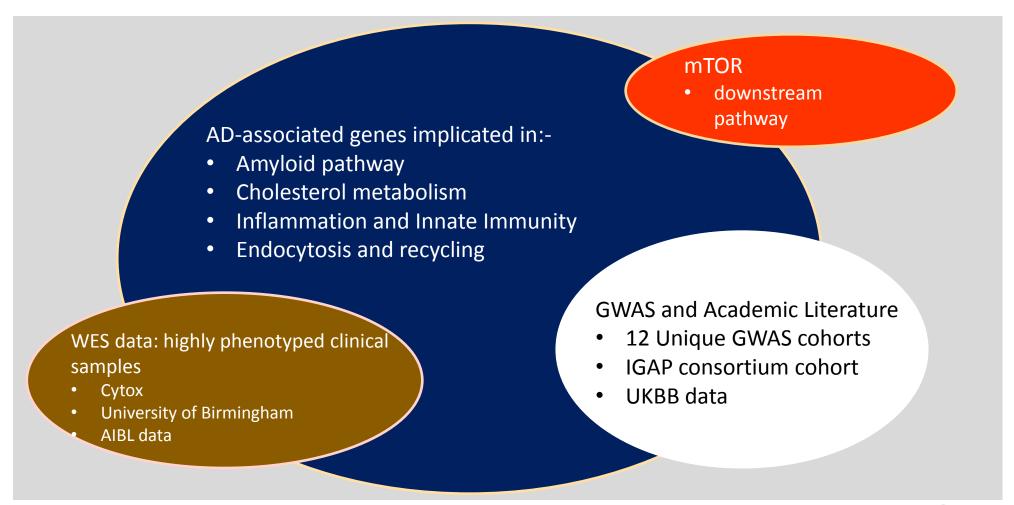


Genetic building block strategy: Polygenic Risk Score



The *varia*TECT™ SNP array: ~130,000 SNPs

The *varia*TECTTM panel is the most comprehensive panel available for the detection of Alzheimer's Disease informative SNPs, comprising novel and known variants in genes pertaining to pathways implicated in Alzheimer Disease aetiology







Polygenic Risk Score (PRS): modelling and validation

- Two basic models:-
 - Hypothesis-driven variant selection (Model 1)
 - Hypothesis-free variant selection (Model 2)
- All PRS models trained in PET-amyloid (or CSF) confirmed cases and controls
- •Blind validation of models in clinically assessed, post-mortem, pathology-confirmed cases and controls

Training and test clinical samples:-

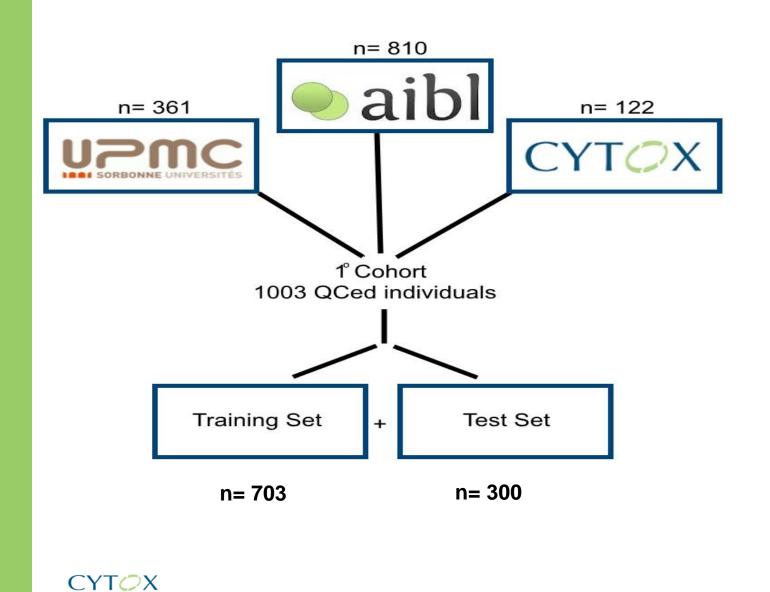
- AIBL Simon Laws, Colin Masters, Larry Ward
- INSIGHT Harald Hampel, Bruno Dubois, Simone Lista
- KU Leuven Rik Vandenberghe, Isabelle Cleynen

Validation Samples:-

UPenn (Brain Bank)
 John Trojanowski, Virginia Lee, Vivianna Van Deerlin, David Irwin



Training set description

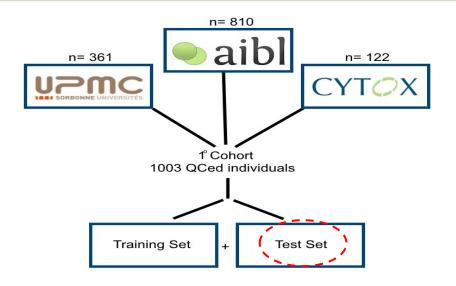


Age (mean)	72.82
Age (range)	55-93
Method of amyloid analysis	PET/CSF
Sample Size	703
Gender ratio M:F	296:407
Ethnicity	Caucasian

% Amyloid +ve	35%
ApoE4 carrier	31%
Clinical Diagnosis	
- Cognitively Normal	76%
- MCI	9%
- Alzheimer's Disease	15%

Test set results

Sample Size	300
% Amyloid +ve	34%
ApoE4 carrier	35%
Clinical Diagnosis	
- Cognitively Normal	71%
- MCI	12%
- Alzheimer's Disease	17%



Model name	Sensitivity	Specificity	AUC	PPV_33	NPV_33
1	72.55%	72.73%	78.93%	80.12%	63.62%`
2	86.29%	86.22%	94.35%	89.07%	82.85%



Validation test set results

Pathology Confirmed Samples from UPenn (blinded analysis results)

Sample Size	237
% Amyloid +ve	88%
ApoE4 carrier	50%
Clinical Diagnosis	
- Cognitively Normal	30%
- MCI	4%
- Alzheimer's Disease	53%
- Other dementia	13%

Model	Sensitivity	Specificity	AUC	PPV33	NPV33
1	0.80	0.80	85.55%	95.78%	41.22%
2	0.72	0.72	75.12%	91.51%	37.98%

Model name	False positive rate	False negative Rate	True Positive rate	True Negative Rate
1	52	4.808	95.192	48
2	24	33.173	66.827	76



Conclusions – presented at AD/PD Conference (Vienna, March 2017)

- Two basic models: Hypothesis-driven and Hypothesis-free variant selection
- Blind validation of models:-
 - True Positive Rate of 95% or greater
 - True Negative Rate around 76% is possible
- Models yield results significantly better than what is currently available in both ApoE4 negative cohorts and ApoE4 mixed cohorts

Level of performance consistent with potential utility in population stratification in clinical trials



Ongoing optimisation of PRS amyloid algorithms in MCI cohorts



PRS Algorithm locked for CLIA/LDT verification and validation:

- Research version available for collaboration today
- PRS Algorithm lock: 3Q 2017
- RUO Assay in CLIA-compliant lab: 4Q 2017



Collaborative opportunities

- Multiple PRS algorithms offering high overall accuracy for prediction of amyloid status
- Word-class scientific team
- Opportunities for true collaborative partnerships

- 1. Stratification for clinical trials savings in cost and time
- 2. Drug responder profiling clinical trial optimisation
- 3. Companion Diagnostic approaches

Next steps...



Genetics and Alzheimer's Disease Expertise

Experienced Management Team

Dr Andrew Carr (Chairman)

GE Healthcare; Thermo Fisher; Teraview; deltaDOT; Akubio

Dr Richard Pither (CEO)

GE Healthcare; Lorantis; UCB-Celltech – Alzheimer's

Dr Kevin Banks (Business Strategy and Marketing)

ADI; Affymetrix – Genetic assay platforms

Dr Alex Gibson (Business Development)

GE Healthcare Business Development – Alzheimer's

Dr Jonathan Wilde (Clinical Product Development)

Veracyte; Astra-Zeneca – Genetic products

Dr Greg Davidson (Software Engineering and Regulatory)

Quotient Diagnostics – regulated software

World-Class Scientific Partners

Prof John Hardy

 Chair of Molecular Biology of Neurological Disease at the UCL Institute of Neurology

Dr Maryam Shoai

Post-doctoral Expert in Genetic Statistics, UCL Institute of Neurology

Dr Zsuzsa Nagy

• Institute of Inflammation and Ageing, University of Birmingham

Profs Colin Masters, Simon Laws, Ian Cooke

• AIBL (The Australian Imaging, Biomarker & Lifestyle Flagship Study of Ageing)

Profs Harald Hampel and Bruno Dubois

Neurological Institute of the University Salpêtrière Hospital in Paris, UPMC

Profs Rik Vandenberghe and Isabelle Cleynen

Research Group Experimental Neurology, KU, Leuven

Profs John Trojanowski, Vivianna Van Deerlin and Virginia Lee

· University Pennsylvannia

