

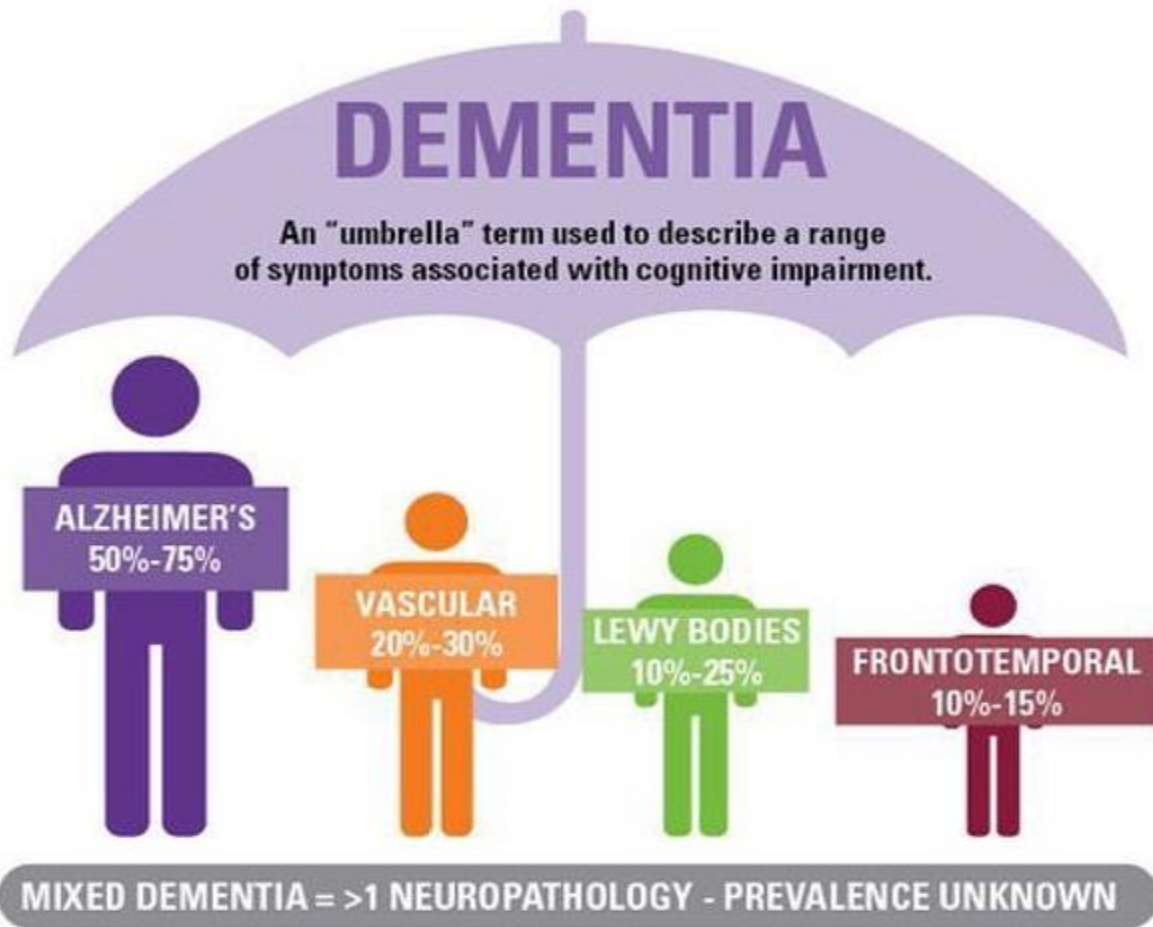
A close-up photograph of a person wearing blue nitrile gloves holding a clear 96-well microplate. The plate is filled with a blue liquid. The background is dark and out of focus, showing some laboratory equipment.

AmplideX[®] PCR/CE *TOMM40* Kit (RUO)

2500-0665

For Research Use Only. Not for use in diagnostic procedures

Types of Dementia



Alzheimer's

Deposits of the protein fragment beta-amyloid (plaques) and twisted strands of the protein tau (tangles)

Vascular

Occurs most commonly from blood vessel blockage

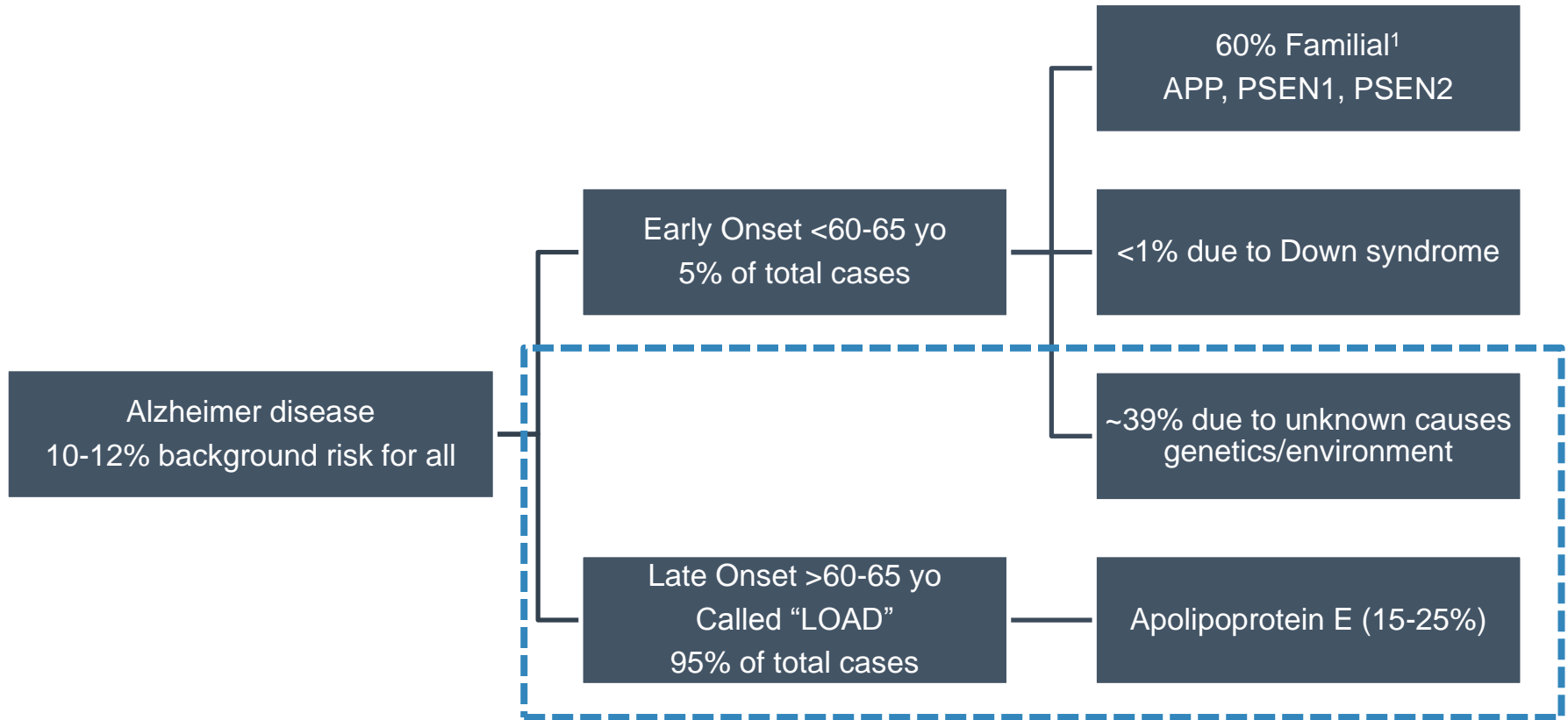
Lewy Bodies

Lewy bodies are abnormal aggregations (or clumps) of the protein alpha-synuclein

Frontotemporal

A mild to severe decrease in overall brain weight and atrophy of the frontal and temporal lobe

Alzheimer's Disease Genetics



¹Campion D. *et al.* Am J Hum Genet. 1999;65:664–70; Brickell KL, *et al.* Arch Neurol. 2006;63:1307–11.

Apolipoprotein E (*APOE*) Identified as AD Risk Factor

Allele	ε2	ε3	ε4
General Frequency	8.4%	77.9%	13.7%
AD Frequency	3.9%	59.4%	36.7%

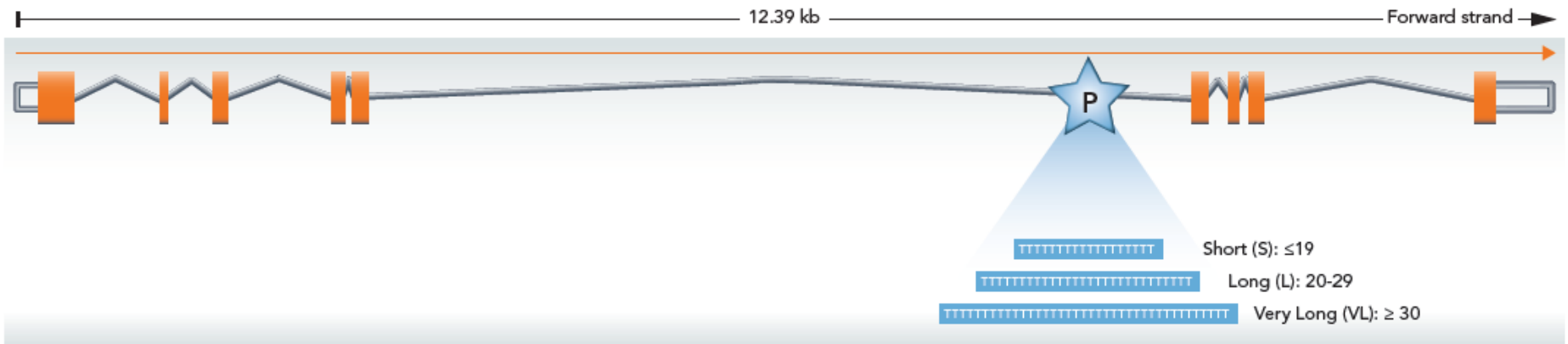
- Principal cholesterol carrier in the brain
- Three alleles: ε2, ε3, and ε4
 - ε2 considered protective for AD
 - ε4 associated with increased risk of AD

APOE Alone Unable to Predict AD Risk

Overall Risk:

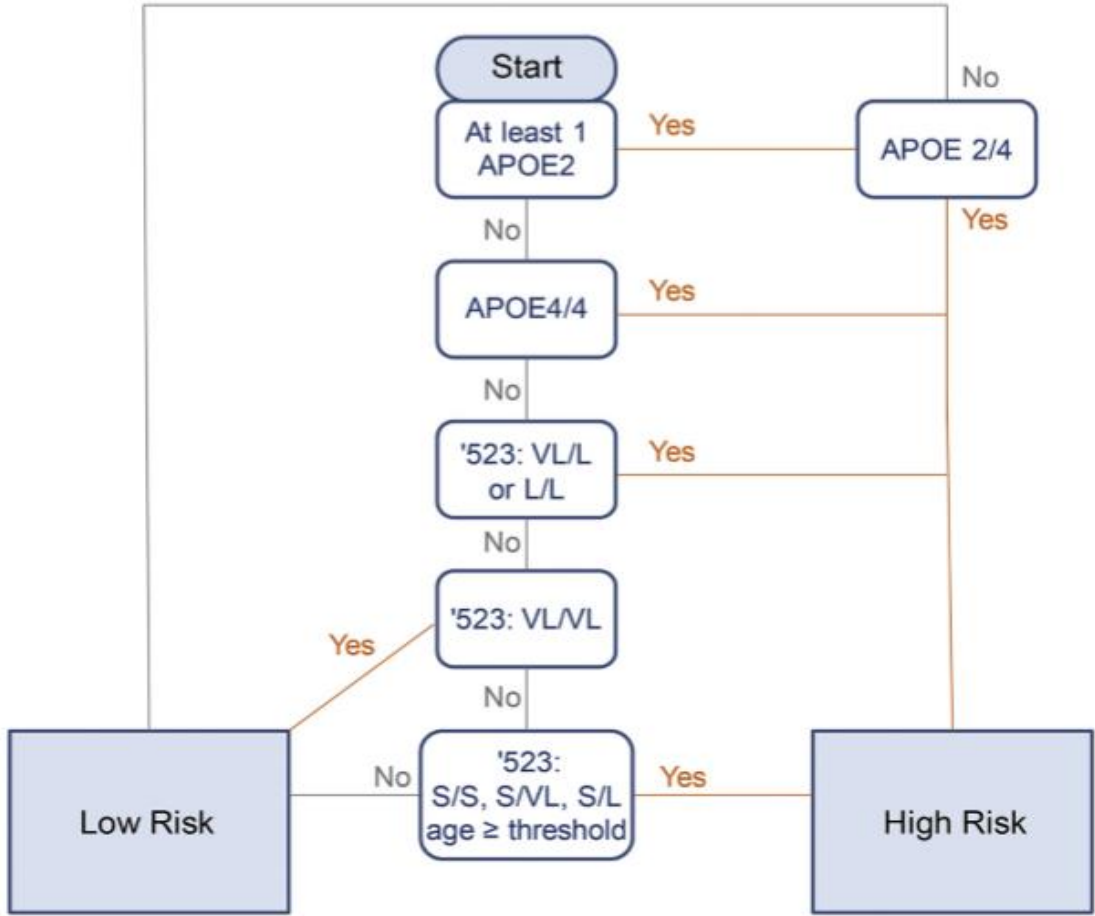
- Depends on combination of risk factors
- Presence of one or more $\epsilon 4$ alleles does NOT mean AD
- Lifetime risk still associated with family history
 - $\epsilon 4$ negative does not remove 2-3x increased risk
- Varies based on age, gender, and ethnicity

TOMM40: Missing Piece For AD Risk



- Encodes Tom40, the translocase of the outer mitochondrial membrane pore subunit
- Mitochondrial Dysfunction: Early defect in LOAD pathogenesis and other age-related diseases
- Poly-T length repeats linked with LOAD

APOE + TOMM40 Poly T Improves the PPV of AD Risk Prediction?

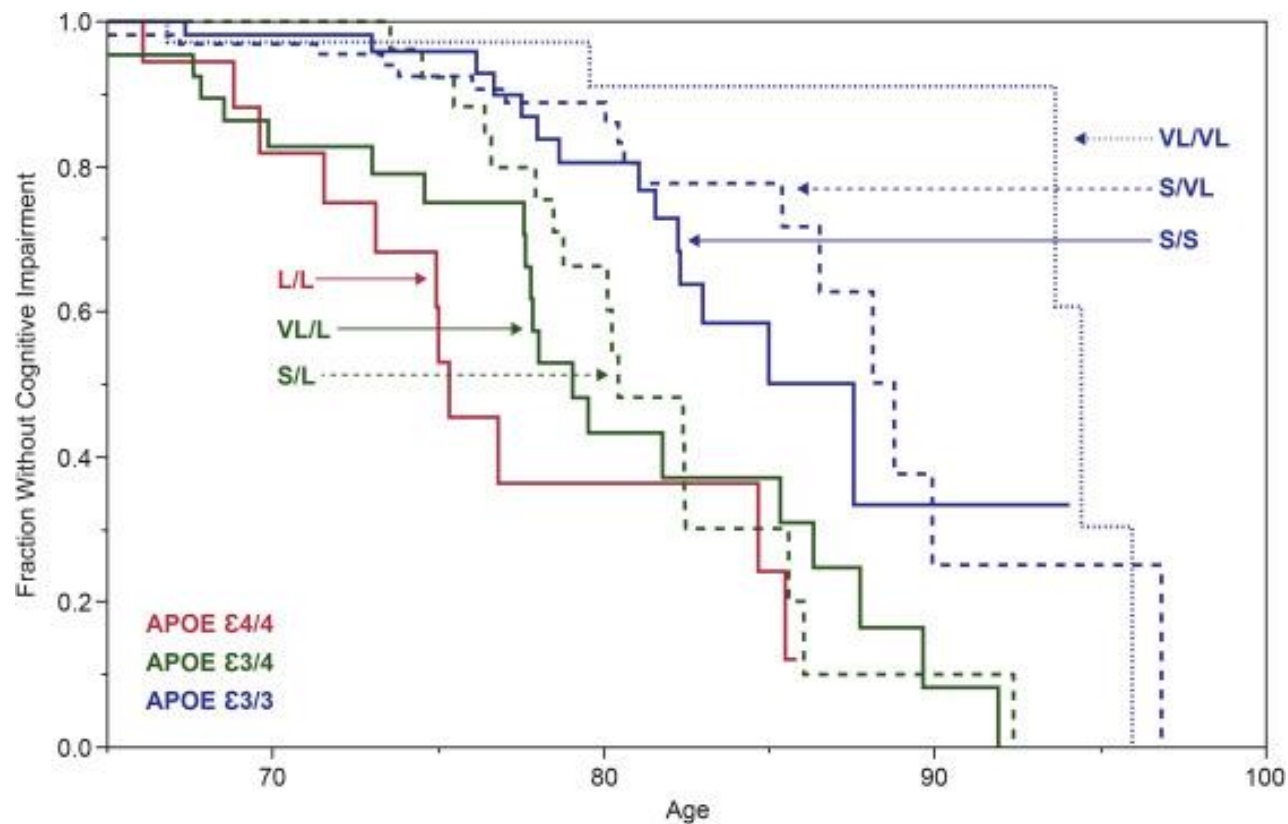


Short (S; ≤19Ts)
 Long (L; 20–29 Ts)
 Very Long (VL; ≥30Ts)



M.W. Lutz et al. / Alzheimer's & Dementia: Translational Research & Clinical Interventions 2 (2016) 30-44

Association of *APOE/TOMM40* with AD Development



Challenges of *TOMM40* '523 Genotyping

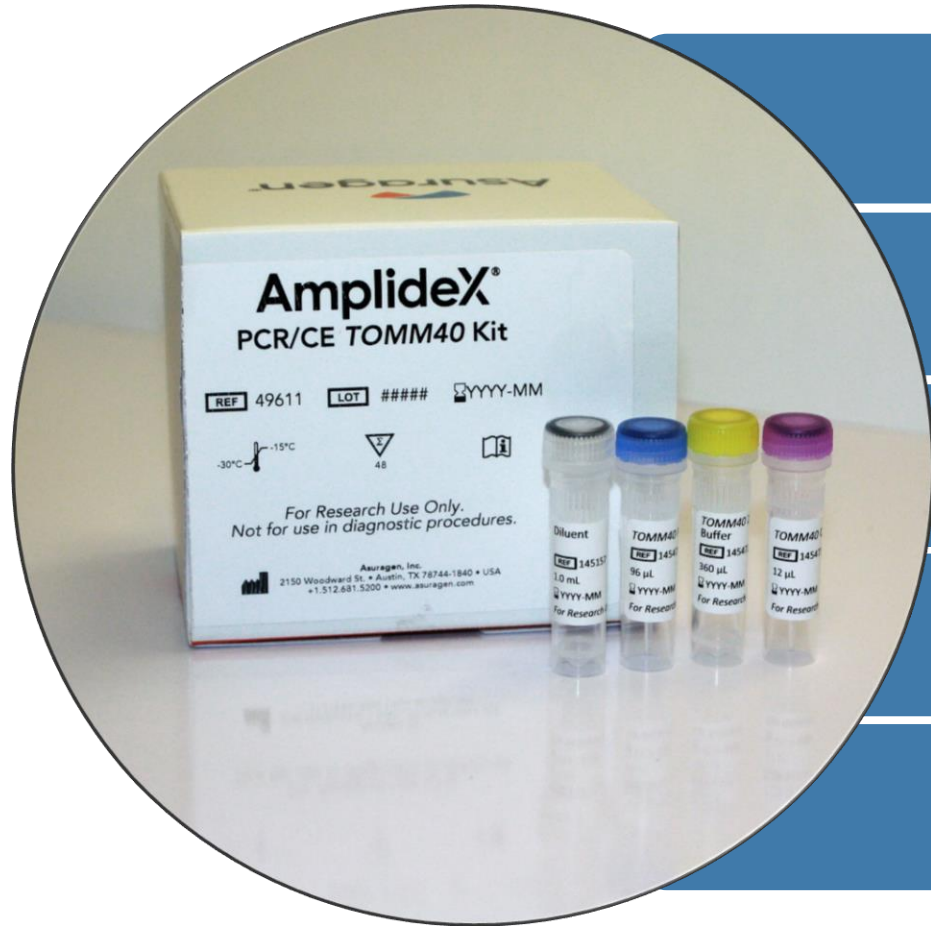
AT Rich Region

- Challenging to amplify
- Hard to resolve (polymerase stutter and slippage)

Lack of Robust Standardized Assays

- Inaccurate results
- Inability to perform high throughput studies

Simple and Robust Test for Detection of *TOMM40* Poly T Repeats



Scalable run sizes

All-inclusive Reagents

Optimized TAT with Minimized HOT

PCR Calibrator for accurate sizing

Sizing accuracy down to 1 bp

AmplideX[®] PCR/CE TOMM40 Kit

Enables Simple, Reliable TOMM40 Gene Research

Technology

Workflow

Performance

- Leverages proven AmplideX[®] chemistry for high homology sequences

- < 3 hr DNA to result
- 30 min Hands-on-Time
- Easy to interpret

- Repeatable
- Reproducible
- Accurate sizing w Single-Base Resolution

Simple workflow minimizes hands on time



Includes all PCR Reagents

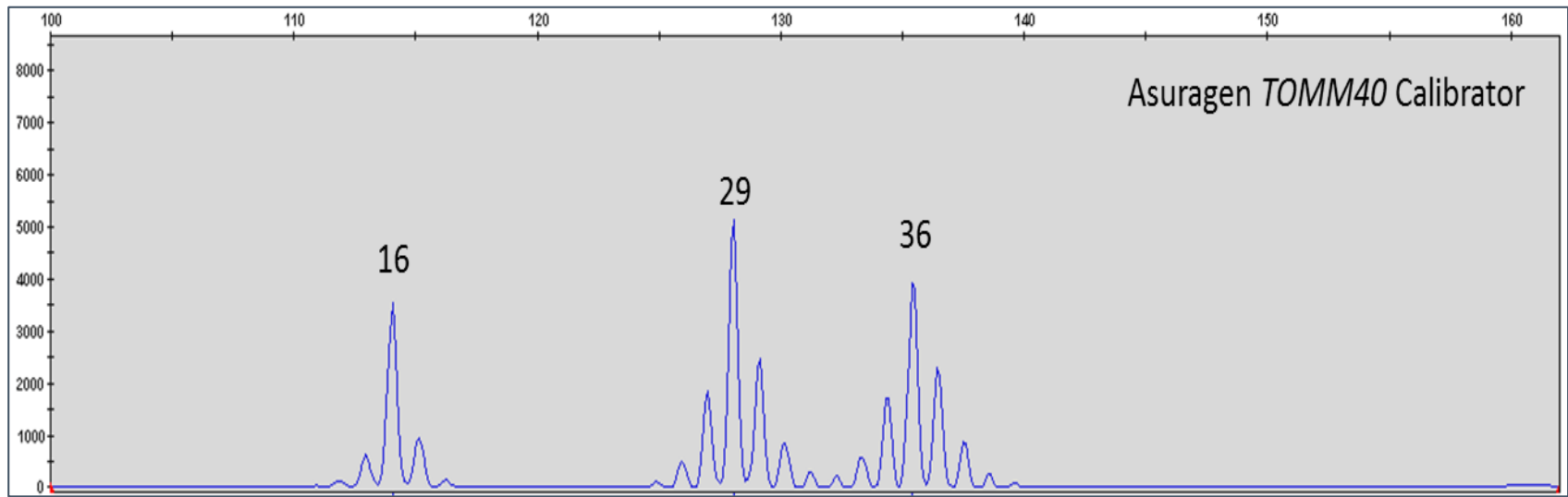
- Calibrator for accurate sizing
- Primer mix
- PCR enzyme and buffers

Detailed protocol guide

- Master mix composition
- PCR cycling conditions

Calibrator Enables Accurate Sizing

- Calculate mobility and correction factor (run or instrument specific)
- Use provided equation from protocol guide to calculate Poly T repeats

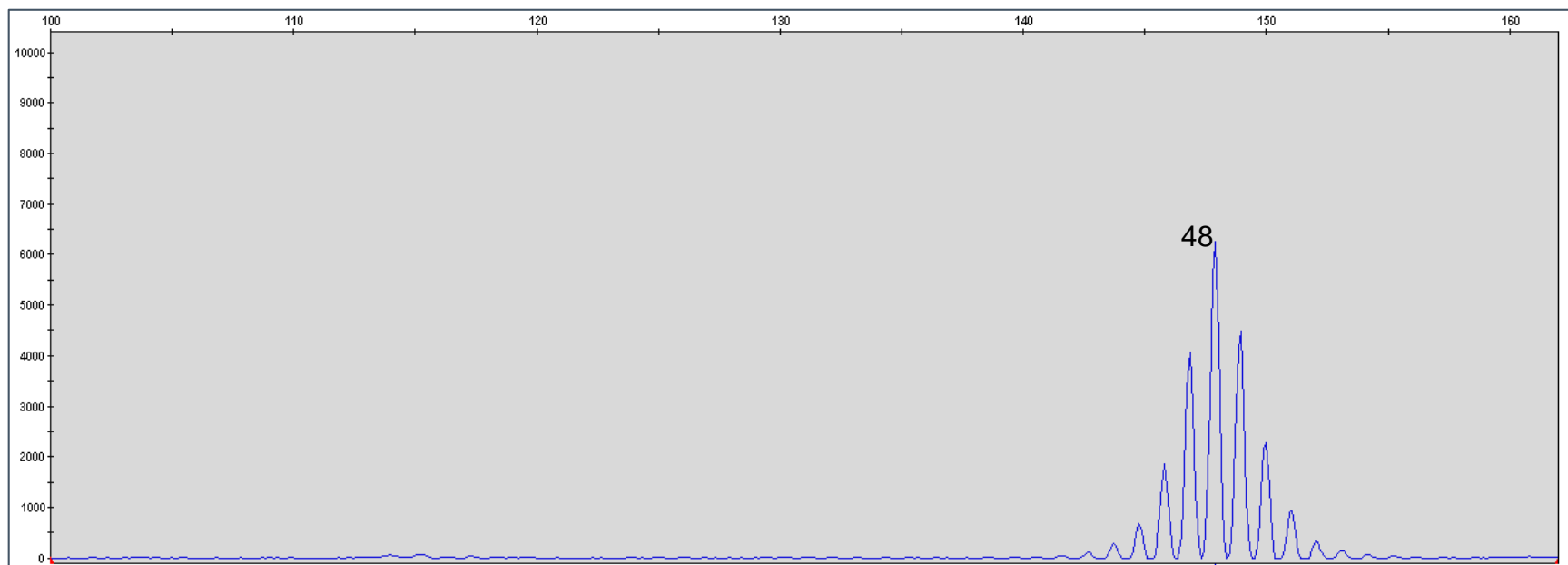


$$T_n = \frac{(CE \text{ Sizing} - Intercept)}{Slope}$$

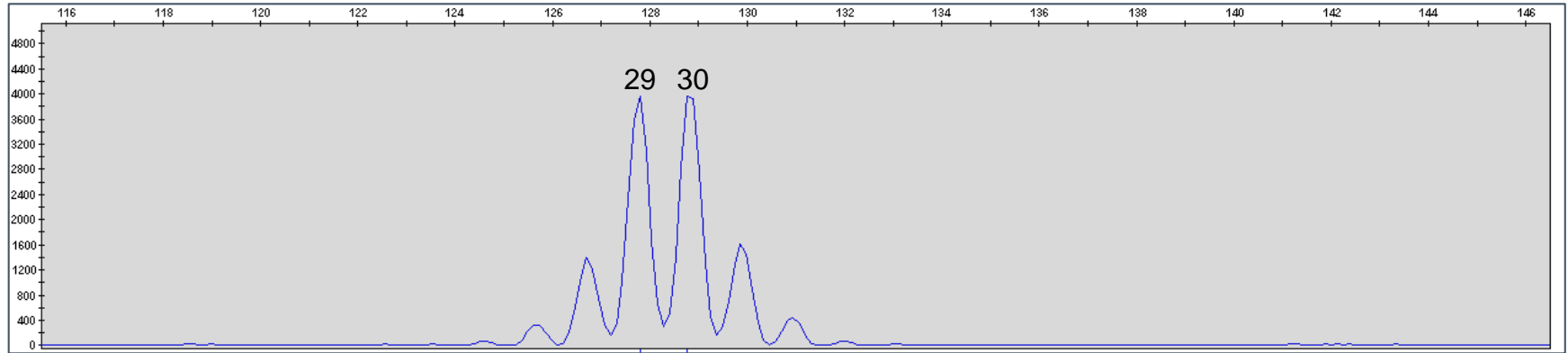
Configuration		C_0	m_0
3730, 3730xL	36 cm	96.92	1.07
3500, 3500xL	50 cm	96.96	1.07

Accurate Sizing Enables Accurate Sample Categorization

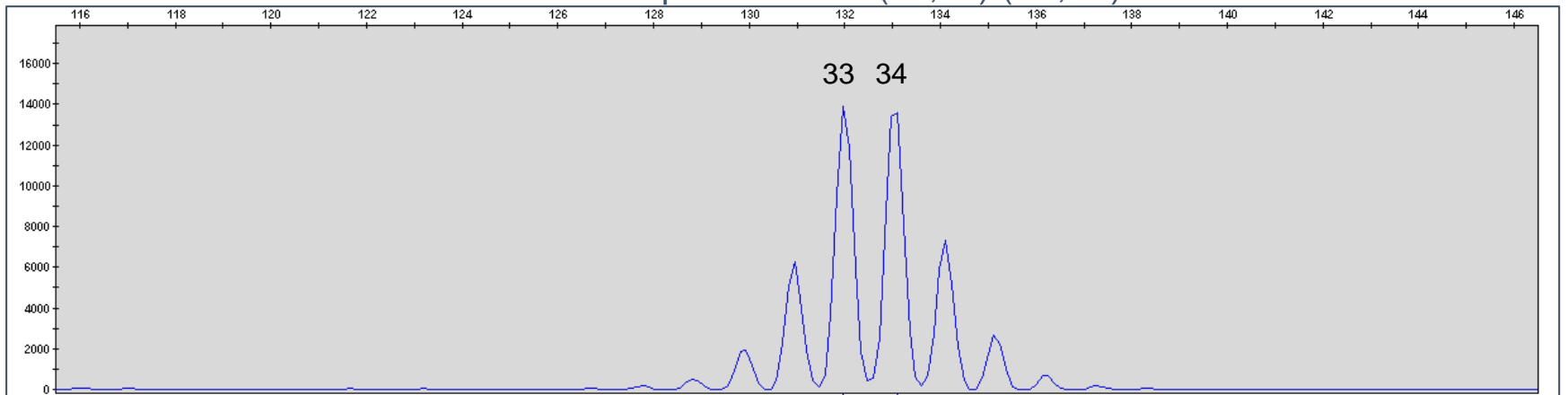
A synthetic 48 mer was accurately sized



Single bp resolution allows ability to distinguish n,n+1 alleles



Coriell sample NA11325 (33,34) (VL,VL)

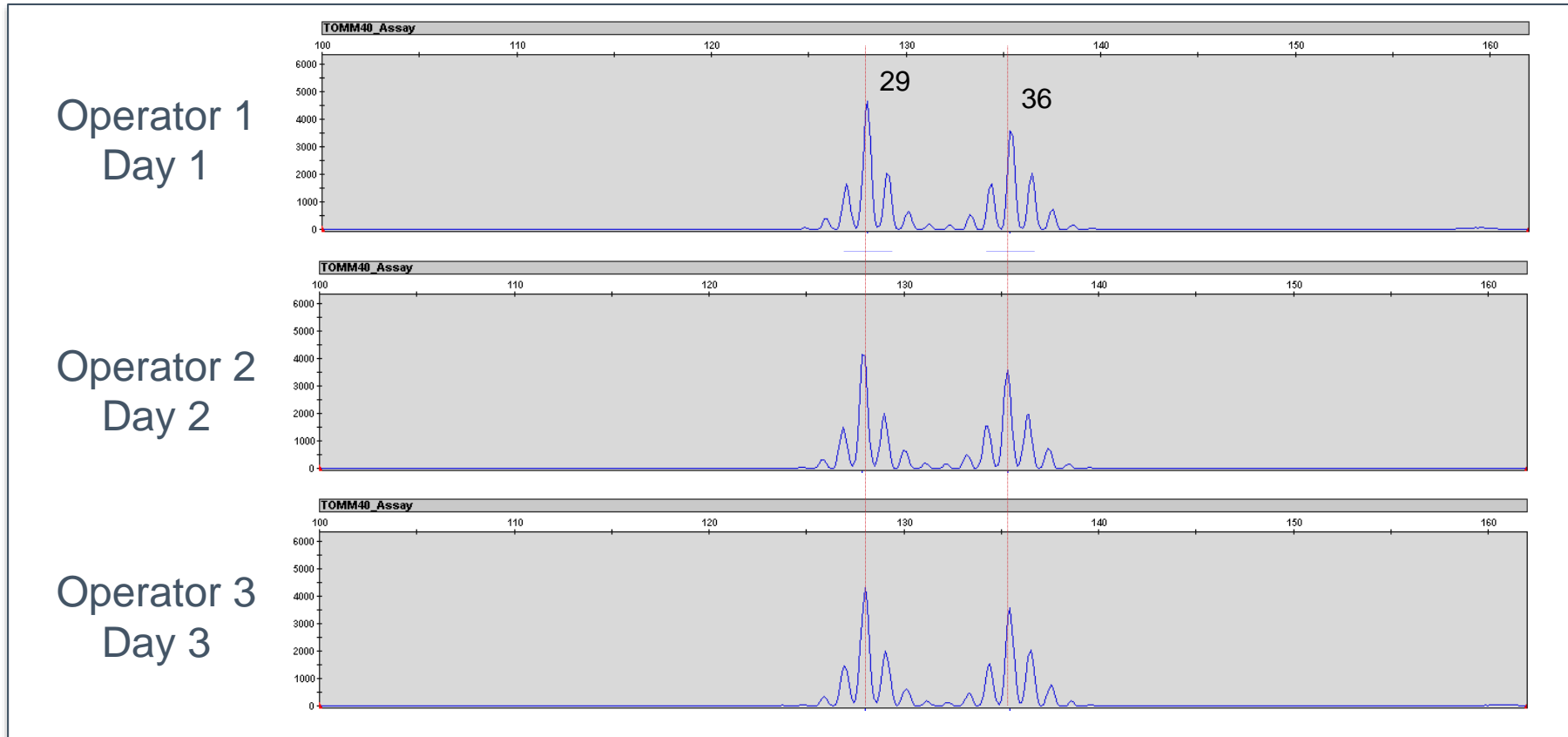


Compatibility with Different Sample Types

Sample Name	Blood	Buccal
F01	16/16	16/16
M02	16/34	16/34
M03	16/16	16/16
F04	16/22	16/22
F05	16/37	16/37
F06	16/29	16/29
M07	16/36	16/36
F08	16/16	16/16
M09	22/36	22/36
F10	16/38	16/38
M11	35/35	35/35
M14	16/36	16/36
M15	16/34	16/34
M18	16/34	16/34

*6 females and 8 males with matched blood and buccal samples

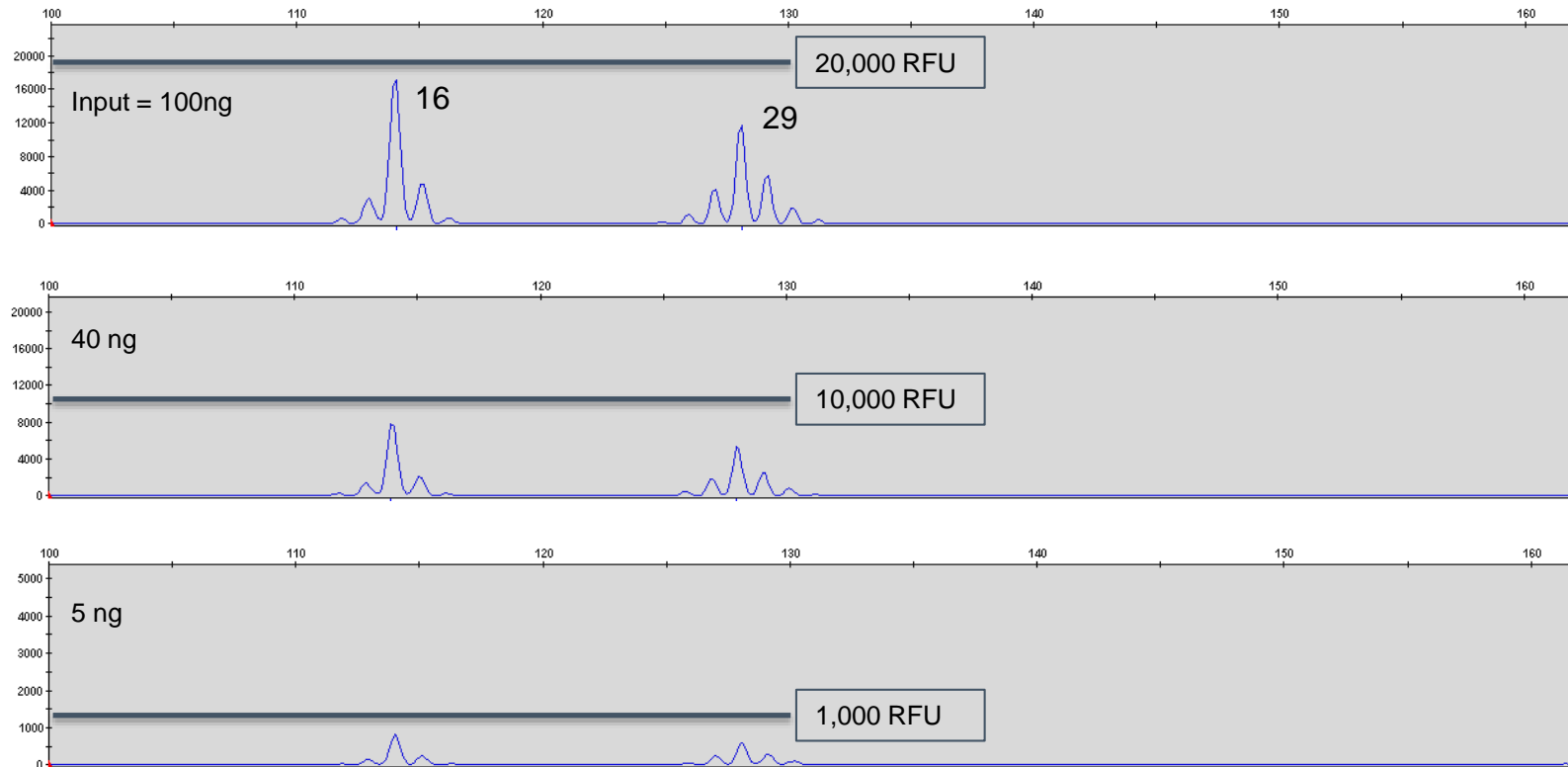
Inter-Operator and Multi-Day Runs Demonstrate Assay Reproducibility



Sample: L/VL, 29/36 alleles

Assay Tolerates Wide Range of DNA Input

Recommended assay input range = 5-100 ng



Simplified Workflow, Reliable Performance

Reducing Complexity

Reagents and protocol enables easy assay set up

Clear data interpretation

Optimizing Workflow

<30 mins of hands on time

<3 hrs of TAT

Quality Performance

1 bp Resolution

Reliable and accurate results