



Towards Development of a 'Silence and Replace' Based Approach for the Treatment of Oculopharyngeal Muscular Dystrophy

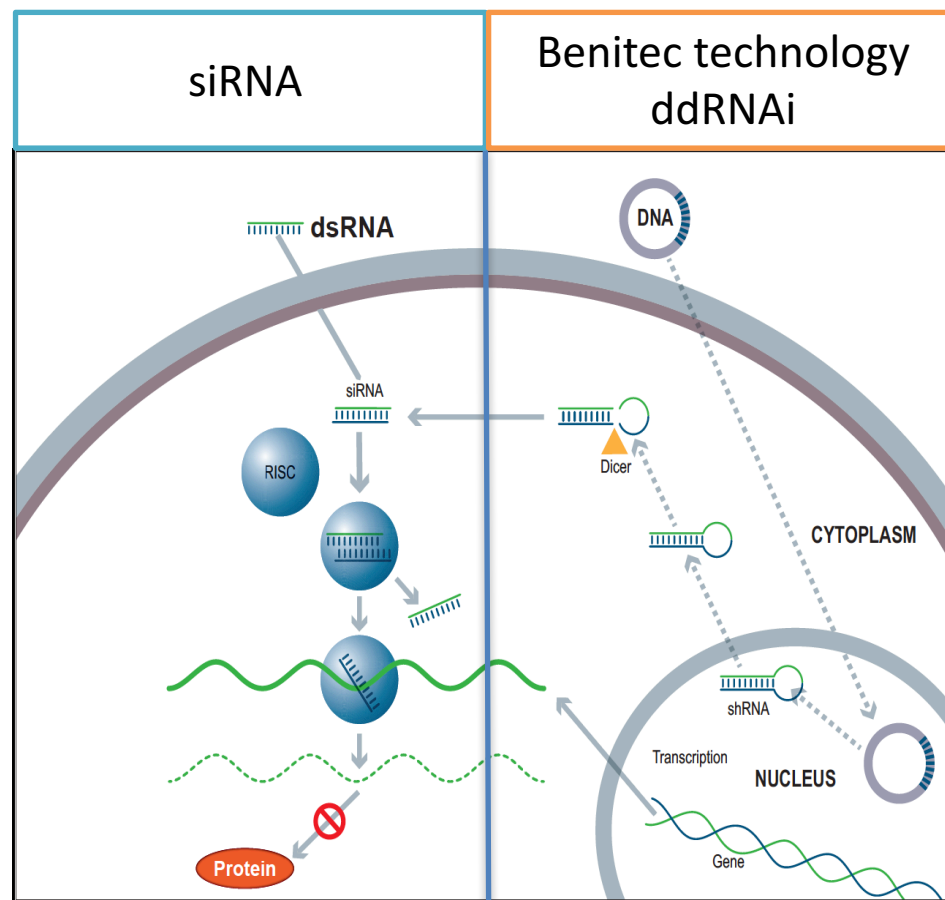
David Suhy / CSO

This presentation contains "forward-looking statements" within the meaning of section 27A of the US Securities Act of 1933 and section 21E of the US Securities Exchange Act of 1934. Benitec has tried to identify such forward-looking statements by use of such words as "expects," "intends," "hopes," "anticipates," "believes," "could," "may," "evidences" and "estimates," and other similar expressions, but these words are not the exclusive means of identifying such statements. Such statements include, but are not limited to, any statements relating to Benitec's pipeline of ddRNAi-based therapeutics, including the initiation, progress and outcomes of clinical trials and any other statements that are not historical facts. Such forward-looking statements involve risks and uncertainties, including, but not limited to, risks and uncertainties relating to the difficulties or delays in our plans to develop and potentially commercialize our product candidates, the timing of the initiation and completion of preclinical and clinical trials, the timing of patient enrolment and dosing in clinical trials, the timing of expected regulatory filings, the clinical utility and potential attributes and benefits of ddRNAi and our product candidates, potential future out-licenses and collaborations, our intellectual property position and duration of our patent portfolio, the ability to procure additional sources of financing and other risks detailed from time to time in filings that Benitec makes with US Securities and Exchange Commission, including our most recent annual report on Form 20-F and our reports on Form 6-K. Such statements are based on management's current expectations, but actual results may differ materially due to various factors, including those risks and uncertainties mentioned or referred to in this presentation. Accordingly, you should not rely on those forward-looking statements as a prediction of actual future results.

Pipeline Programs

Program	Delivery	Discovery	Preclinical	IND-Enabling	Phase I/II	Status
Oncology						
HNSCC - BB-401	Plasmid intratumoral	<div></div>				<ul style="list-style-type: none">Phase 1 clinical POC completePhase 2 FPE – 1Q18
HNSCC - BB-501	ddRNAi intratumoral	<div></div>				<ul style="list-style-type: none">Construct design complete<i>In vivo</i> proof of concept – 4Q17
Infectious Disease						
HBV - BB-103	AAV intravenous	<div></div>				<ul style="list-style-type: none">Pre-IND completedIND-enabling work ongoing
Ocular Disease						
AMD - BB-201	Novel AAV intravitreal	<div></div>				<ul style="list-style-type: none">Capsid biodistribution complete<i>In vivo</i> proof of concept – 4Q17
Orphan Disease						
OPMD - BB-301	AAV intramuscular	<div></div>				<ul style="list-style-type: none"><i>In vivo</i> proof of concept – 3Q17

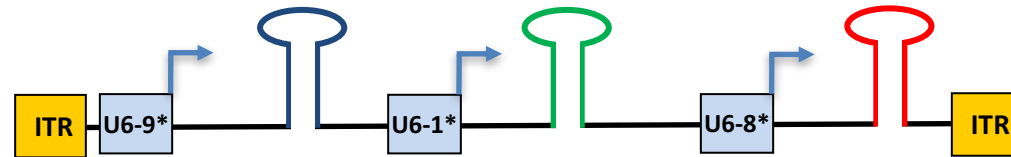
- Combines RNA interference with gene therapy delivery
- Long term therapeutic potential from a single administration
- Steady state levels of gene expression
- Silence a single gene or target multiple genes simultaneously
- Silence/replace strategies of mutant proteins



ddRNAi Platform: Flexibility in shRNA Expression Design to Fit Disease Indication

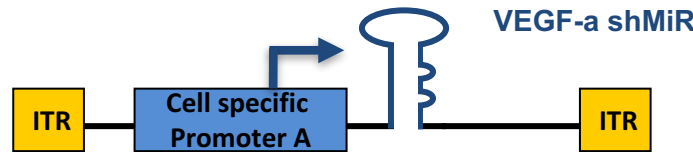
Characteristics:

- Multi-targeting
- Lowered shRNA expression through altered promoters
- Tissue restricted expression
- Defined shRNA loop processing
- Multi-targeting
- Defined shRNA loop processing
- More Robust shRNA expression
- Tissue restricted expression
- Silence disease causing allele
- Replace expression with wildtype protein

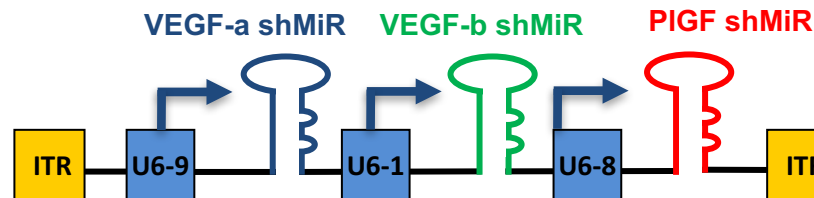


Drug:

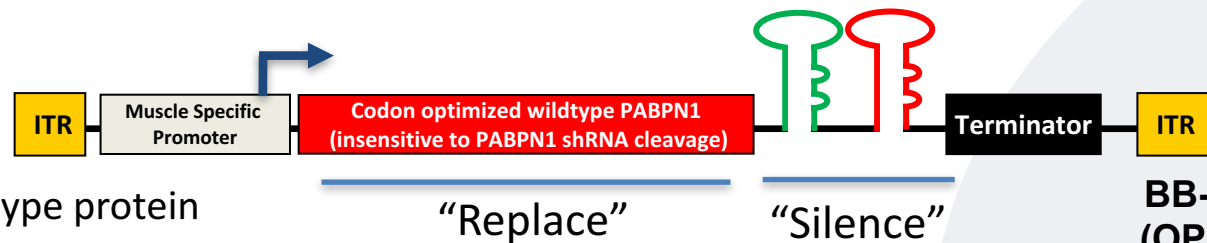
TT-034
(HCV)



BB-AMD-211
(Ocular)



BB-201
(Ocular)



BB-301
(OPMD)

OPMD

Oculopharyngeal Muscular Dystrophy

Rare autosomal dominant inheritance

- 1:100,000 (Europe)
- As high as 1:600 in specific populations
- Founder effect in Quebec, Canada

Typically onset occurs in the fifth to early sixth decade of life

Characterised by:

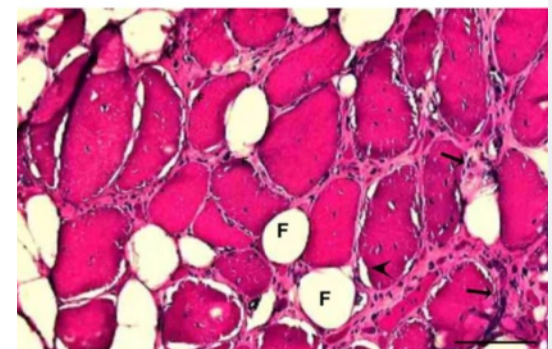
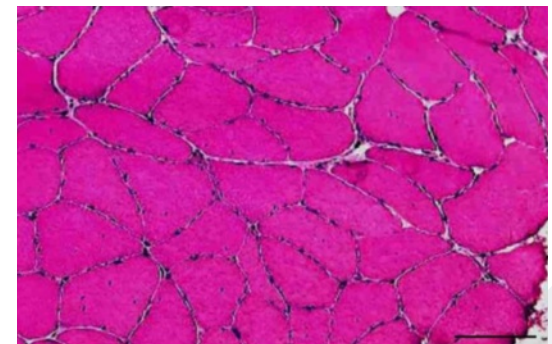
- eyelid drooping (ptosis)
- swallowing difficulty (dysphagia)
- proximal limb weakness
- death due to aspiration pneumonia & malnutrition



Histopathology

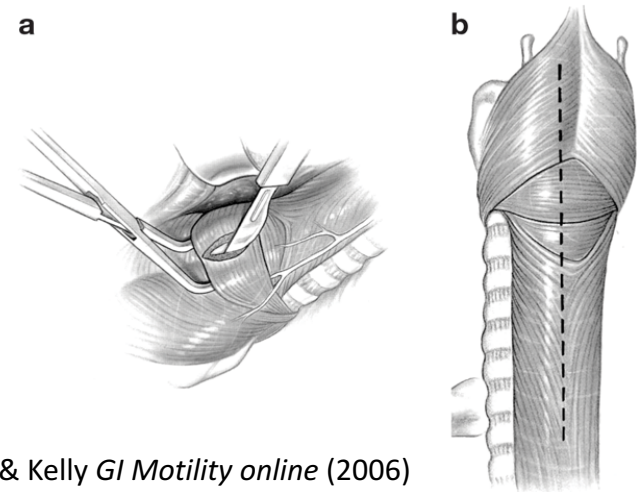
- Decrease of muscle fiber number
- Variation in the size of muscle fibers
- Fibrosis (connective tissue)

Raz et al., *BMC Neurology* 2013, **13**:70



Only Widely Used Treatment is a Surgical Procedure That is Ineffective Over Time

- **Cricopharyngeal myotomy** : a surgical intervention to improve swallowing, but does not correct the progression of the disease since it has a genetic basis.
- **Stem cell transplants**: grafting of autologous myoblasts isolated from unaffected quadriceps or sternocleidomastoid muscles into the esophagus of the patient. Some short term efficacy but transplanted cells still carry the genetic defect.



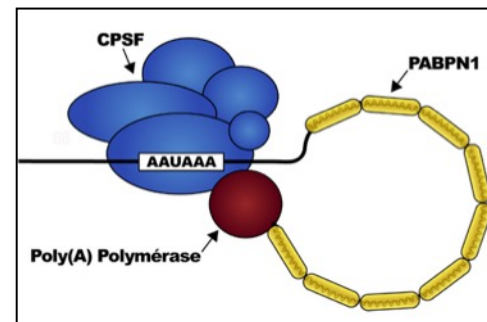
Chu & Kelly *GI Motility online* (2006)

Molecular Therapy (2014); **22** 1, 219–225

Genetic Basis of OPMD: Expansion of the Poly-alanine Tract Within PABPN1

PABPN1:

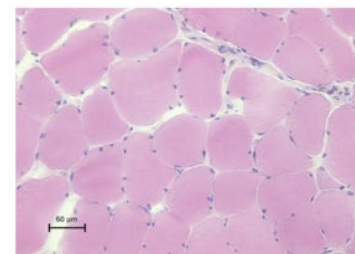
- a ubiquitous factor that promotes interaction between the poly(A) polymerase and CPSF (cleavage and polyadenylation specificity factor) and thus controls the length of mRNA poly(A) tails, mRNA export from the nucleus, and alternative poly(A) site usage.



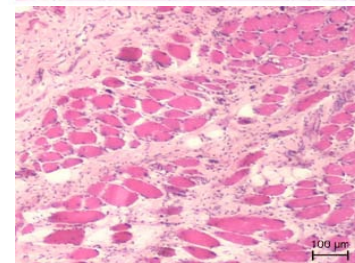
In OPMD:

- a genetic mutation results in trinucleotide repeat expansion within exon 1 of PABPN1 and results in an expanded poly-alanine tract at the N-terminal end of PABPN1.

WT	ATG (GCG) ₆	-----	(GCA) ₃	GCG GGG GCT GCG..
MUT	ATG (GCG) ₆	(GCG) ₁₋₇	(GCA) ₃	GCG GGG GCT GCG...--



Non-affected



Affected

Intranuclear Inclusions (INIs), the Hallmark of OPMD

Expansion of the short (GCG) trinucleotide repeat
in the coding sequence of PABPN1

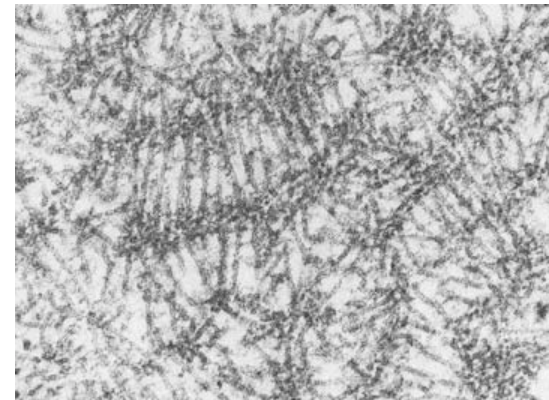


The mutated protein has 11-17 alanines in the N-Terminal
domain instead of 10



Protein aggregation forms intranuclear inclusions (INIs)

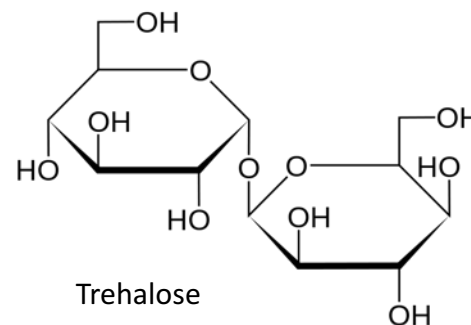
- Tubular filaments
- Resistant to degradation
- INIs found in the nuclei of skeletal muscle fibres (both affected and non-affected)



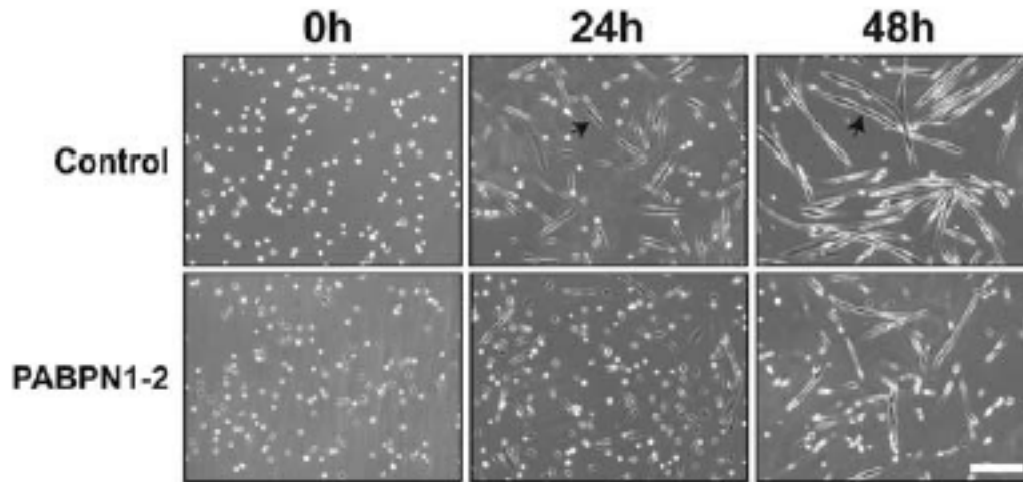
Tomé & Fardeau, 1980

Trehalose

- BioBlast is in Phase II clinical testing of Cabaletta, a chemical chaperone that prevents pathological aggregation of proteins within cells. The active ingredient Trehalose, a disaccharide of glucose, is thought to induce autophagy and stimulate intracellular clearance of the protein aggregates.
- The drug is administered weekly by intravenous infusion.



PABPN1 is Required to Maintain Muscle Function



Human Molecular Genetics, 2010, Vol. 19, No. 6 1058–1065
doi:10.1093/hmg/ddp569
Advance Access published on December 24, 2009

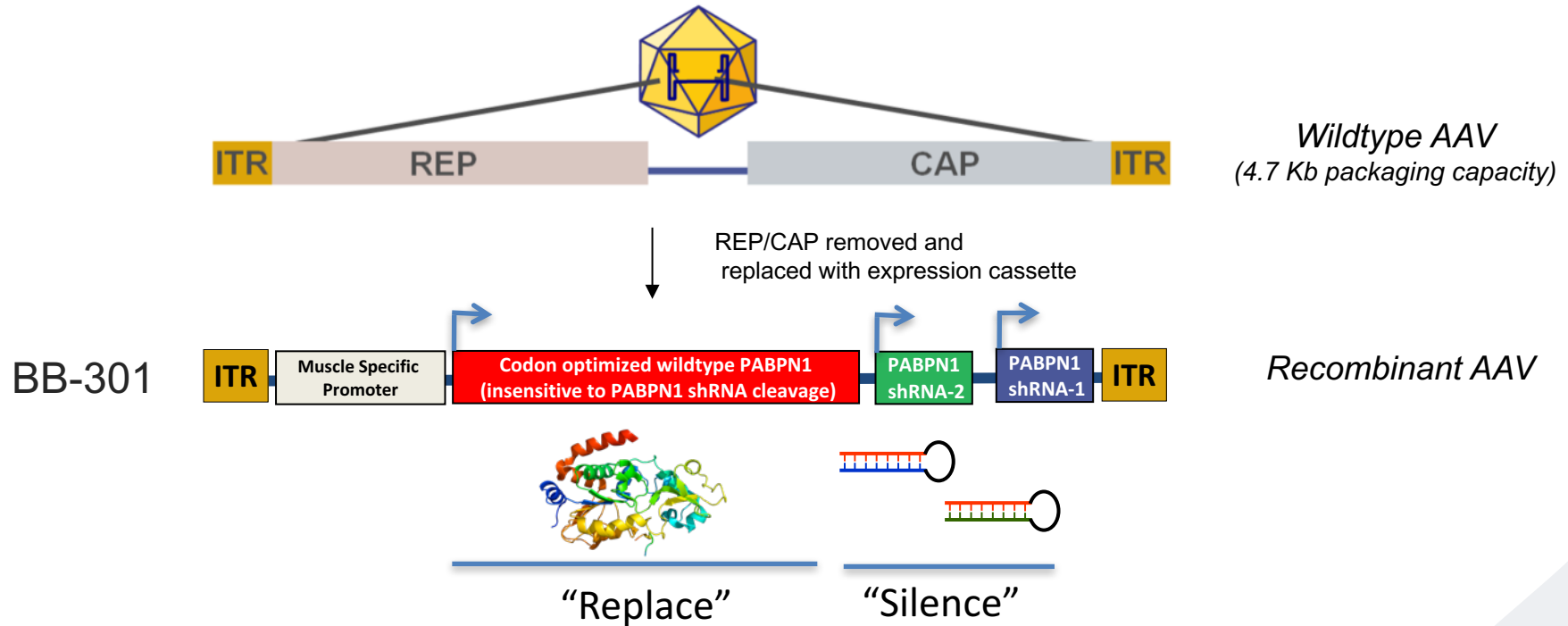
Loss of nuclear poly(A)-binding protein 1 causes defects in myogenesis and mRNA biogenesis

Luciano H. Apponi¹, Sara W. Leung², Kathryn R. Williams¹, Sandro R. Valentini³,
Anita H. Corbett^{2,*} and Grace K. Pavlath^{1,*}

- PABPN1 is required for normal myoblast proliferation and differentiation
- PABPN1 is required for proper polyadenylation in muscle cells
- PABPN1 is required for proper poly(A) RNA export from the nucleus

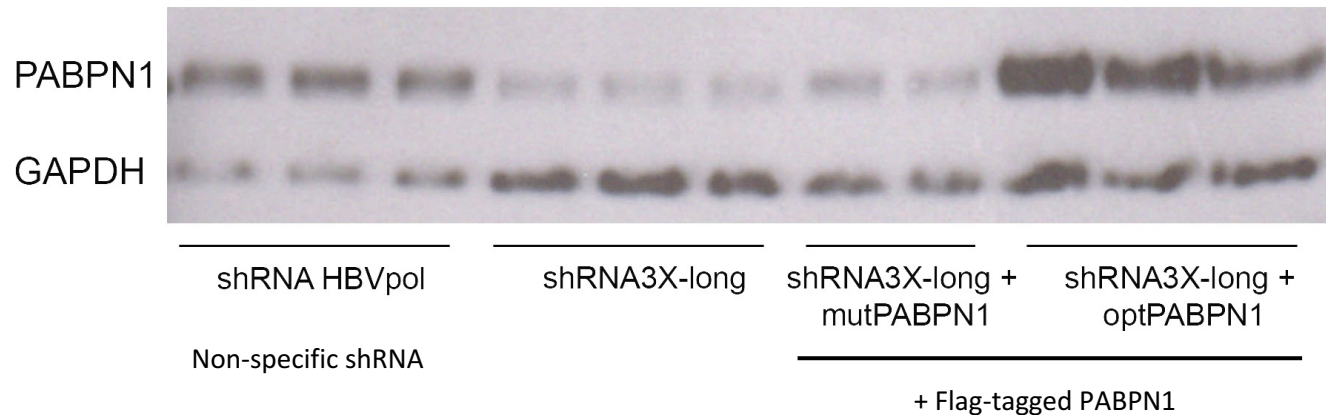
****Thus, an effective treatment likely requires maintaining endogenous function in addition to eliminating mutant protein aggregates**

BB-301: 'Silence and Replace' Approach



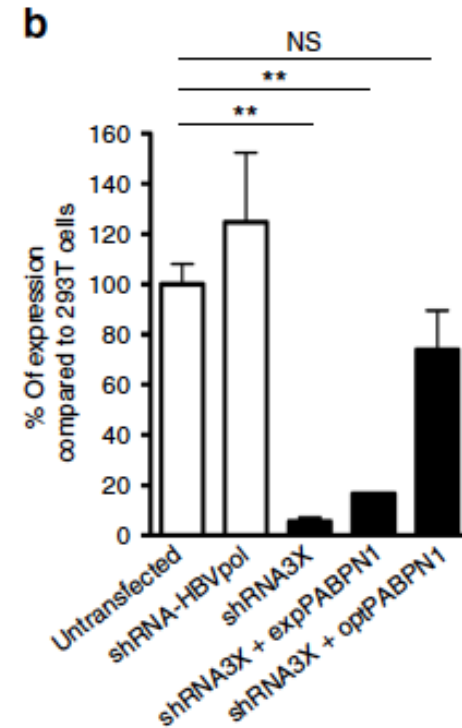
- Non-integrating, non-pathogenic viral delivery system
- To date, AAV has been used in over 173 clinical trials with excellent safety record
- Sustained expression (years) following single injection

Expression of “codon optimized” wildtype PABPN1 is not knocked down by shRNA



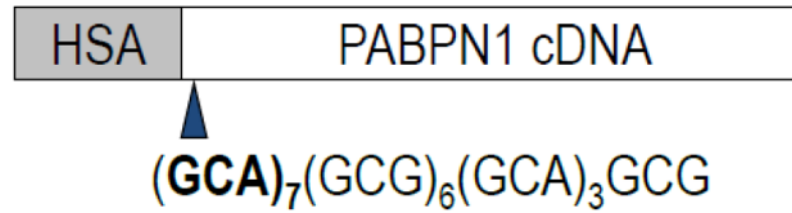
Initial sequence GGACATGGA GGAA GAA GC TGAGAAGCTAA AGGAG CTAC...

Codon-modified GGACATGGAAGA GGA GGC CGA AAAA CTAA C GGAG T TAC...



****Codon modified wildtype PABPN1 is resistant to knockdown with shRNA**

An Animal Model of OPMD: The 'A17' Mouse



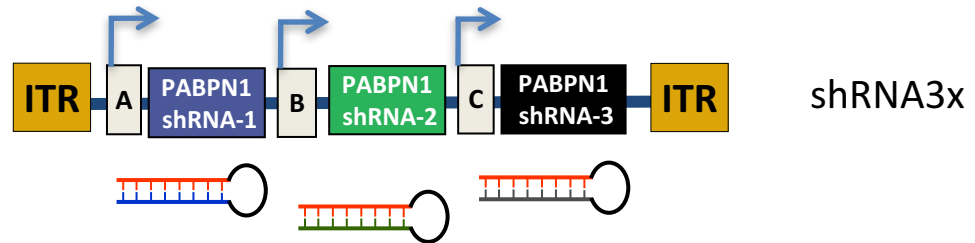
- Transgenic mouse: express a mutated bovine PABPN1 driven by the human skeletal actin promoter in addition to the endogenous PABPN1
- Recapitulates severe muscle atrophy
- Mimics many of the disease pathologies:
 - Progressive muscle weakness/ Atrophy
 - Fibrosis
 - Mitochondrial / Ubiquitin-Proteasome defects
 - Muscles contain intranuclear inclusions

Nature Medicine **11**, 672 - 677 (2005)
Published online: 1 May 2005 | doi:10.1038/nm1242

Doxycycline attenuates and delays toxicity of the oculopharyngeal muscular dystrophy mutation in transgenic mice

Janet E Davies¹, Lin Wang¹, Lourdes Garcia-Oroz¹, Lynnette J Cook¹,
Coralie Vacher¹, Dominic G O'Donovan² & David C Rubinstein¹

Assessment of Efficacy in the A17 Mouse Model



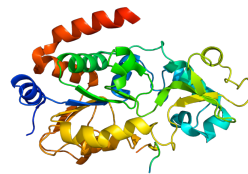
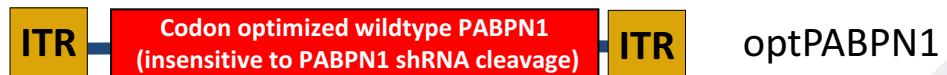
Mice TA Muscles injected with:

2.5e10 vg scAAV8-shRNA3X

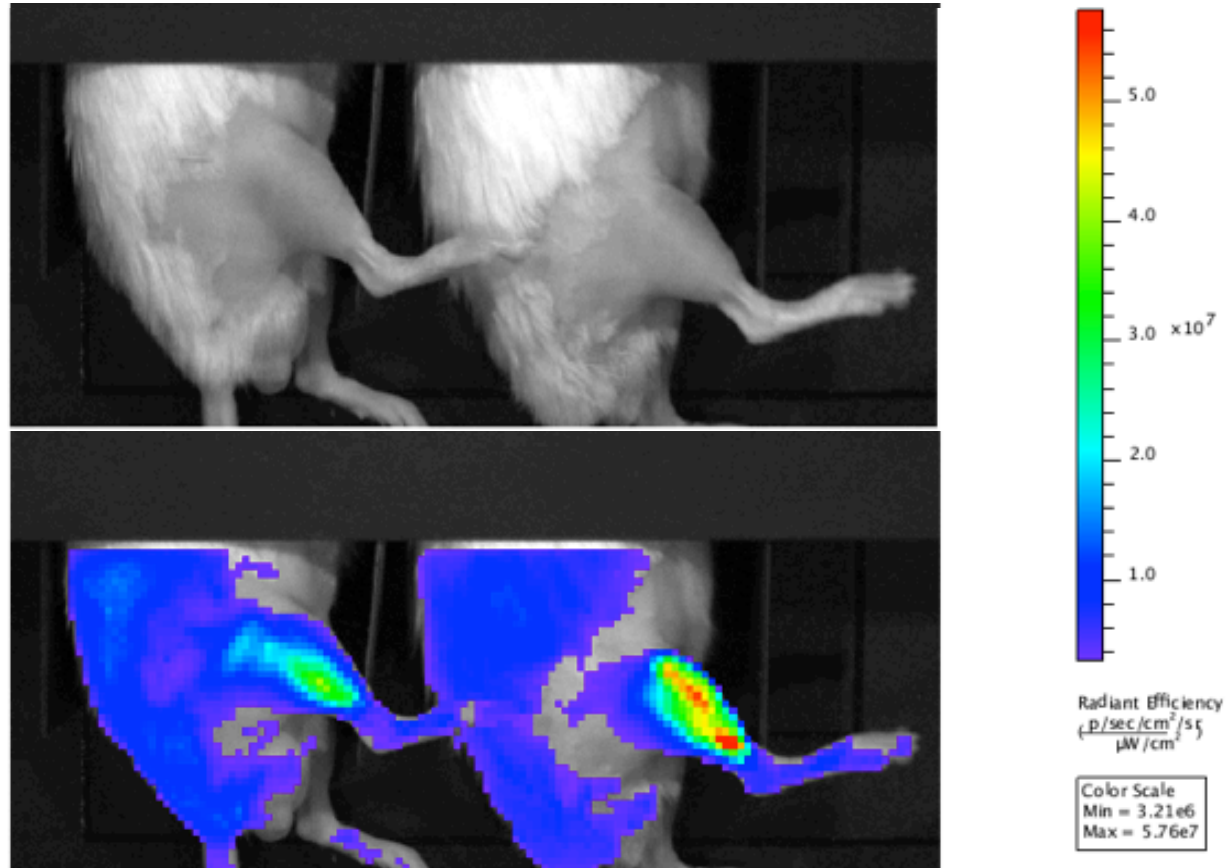
1.3e11 vg ssAAV9-optPABN1

Analyses at week 18 post injection

AND / OR

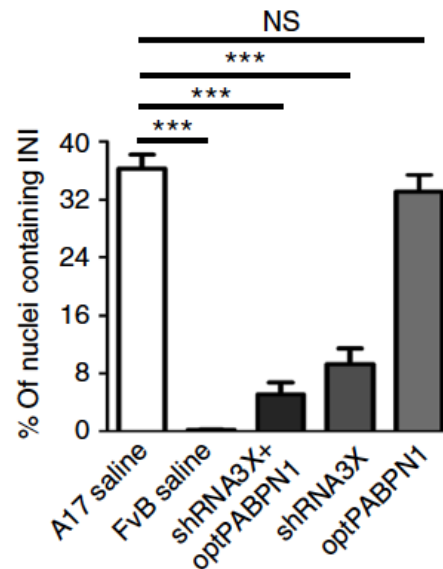
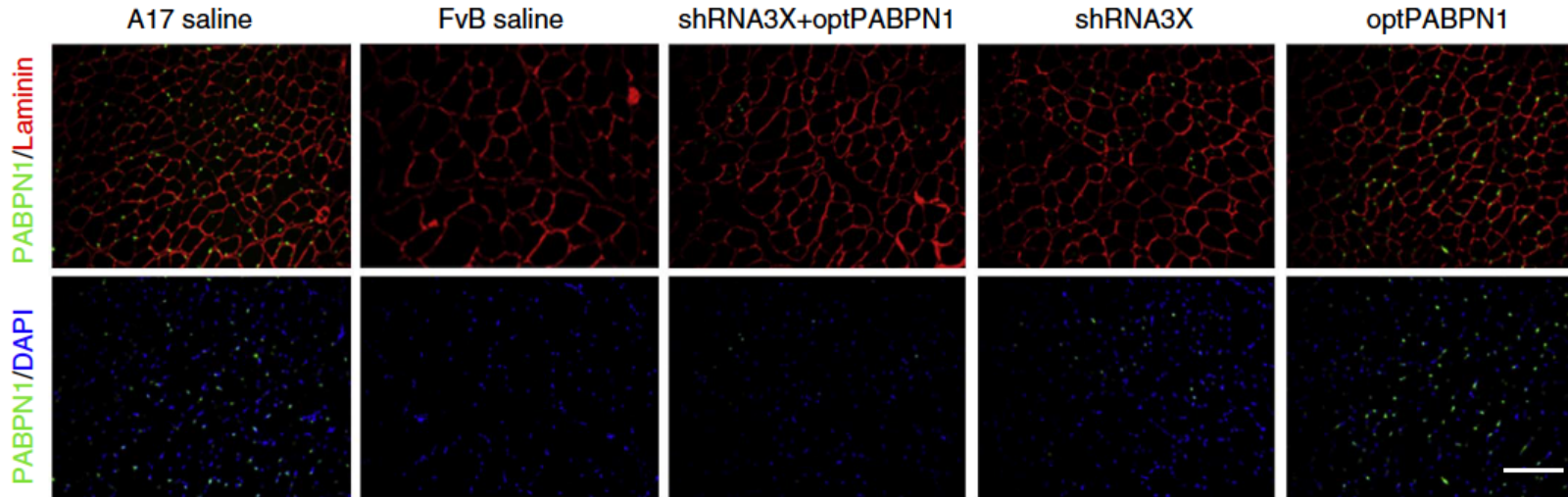


Use of AAV / Intramuscular Injections for Delivery into Muscles Tissues

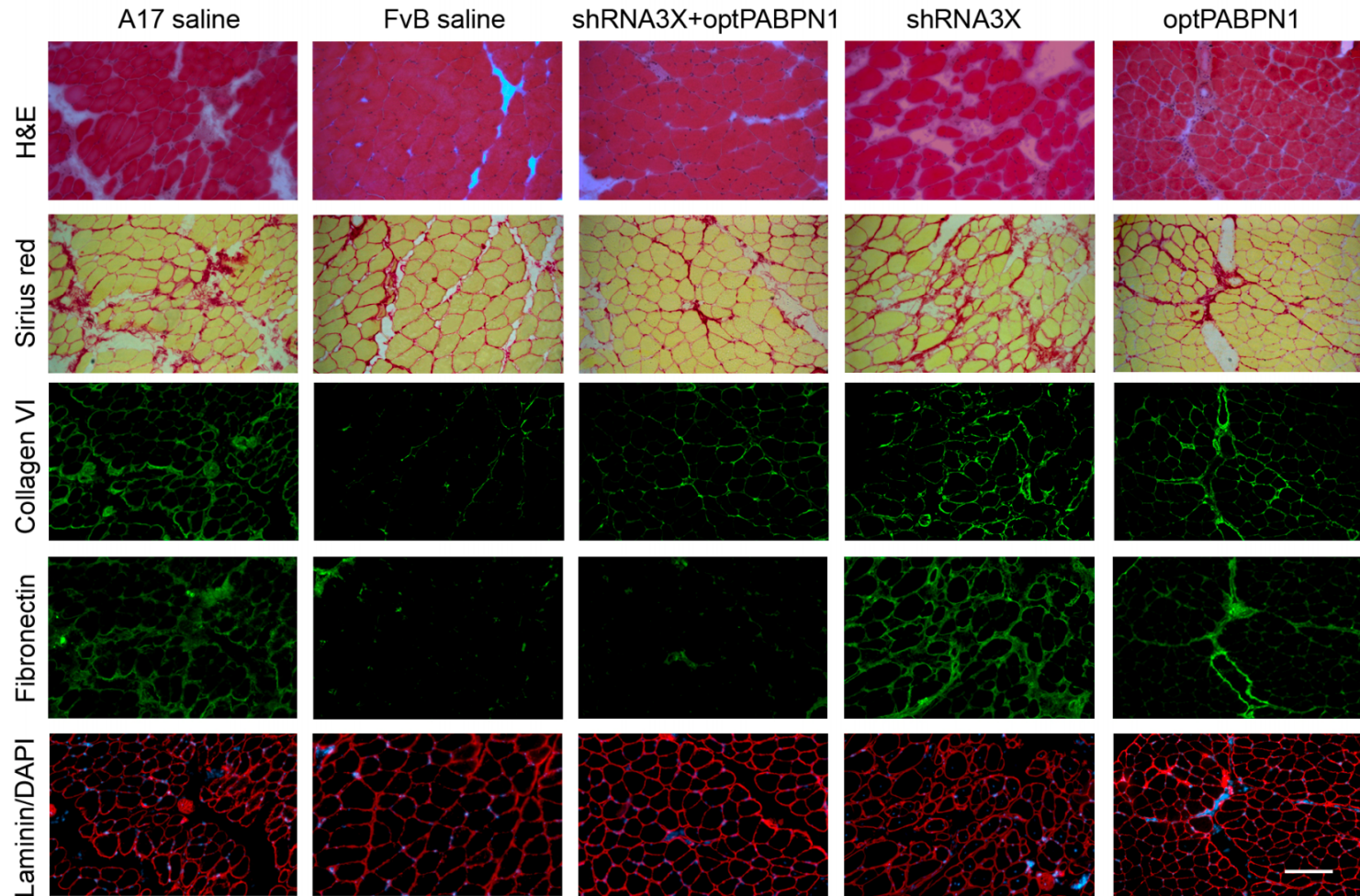


- **Intranuclear Inclusions**
- **Fibrosis**
- **Muscle Fiber Size**
- **Muscle Strength**
- **Gene Expression**

Silence and Replace Strategy Reduces Insoluble Aggregates in Muscle Sections of Treated A17 Mice

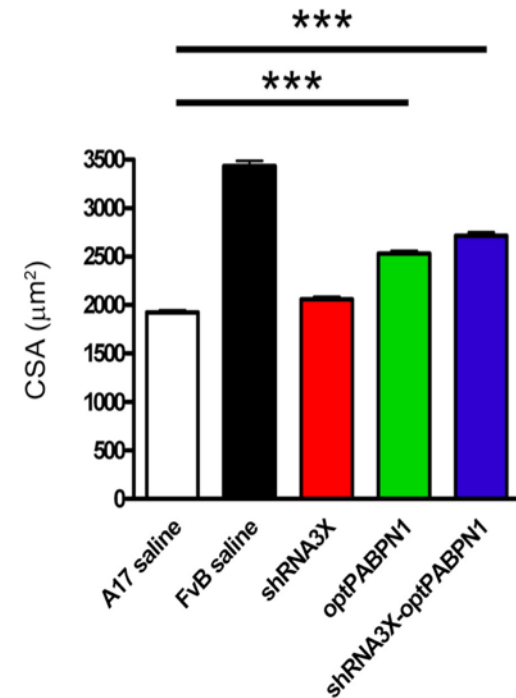
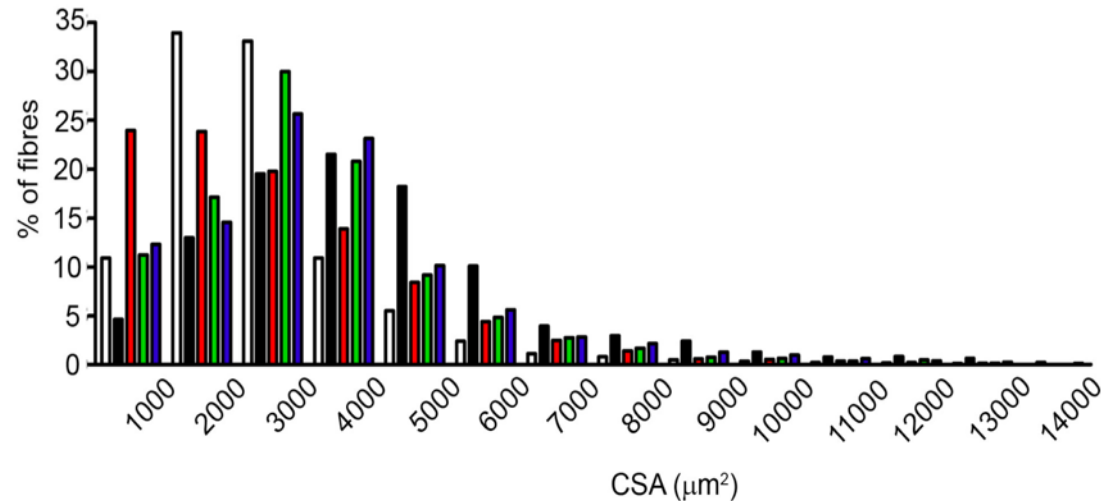


Silence and Replace Strategy Reduces Fibrosis in Transverse Muscle Sections of Treated A17 Mice

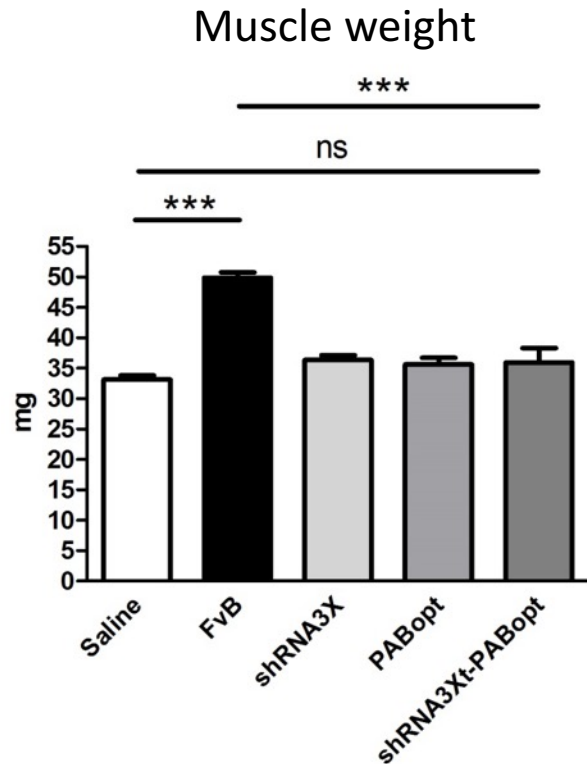


Partial Reversal of Atrophy: a Cross Section Analysis of Muscle Fibers in Treated A17 Mice

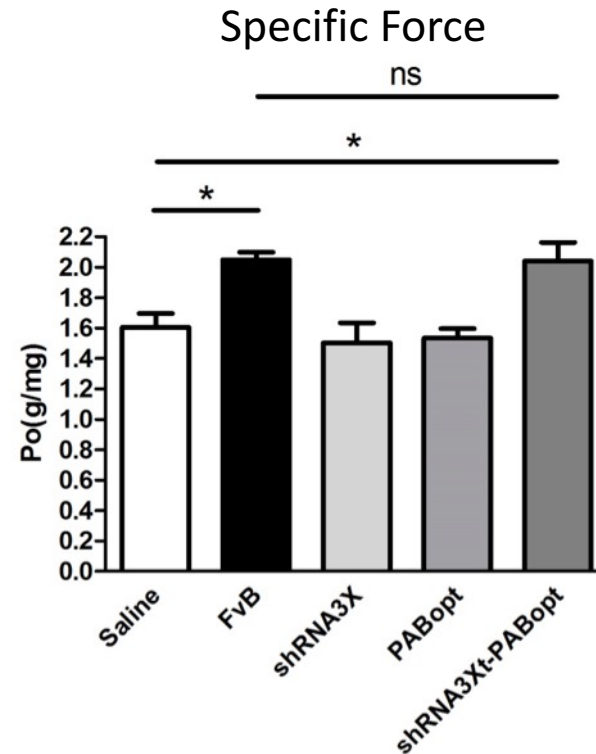
muscle fiber size assessed by quantifying fiber cross-sectional area (CSA)



Assessment of Muscle Atrophy and Restoration of Specific Force in Treated A17 Mice



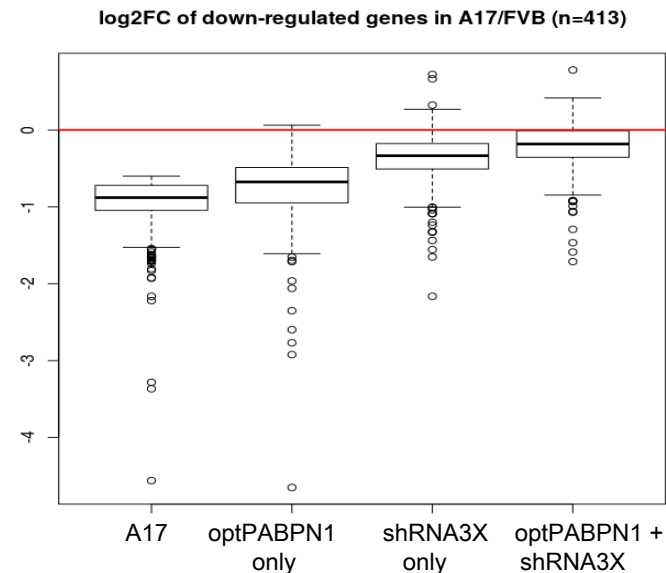
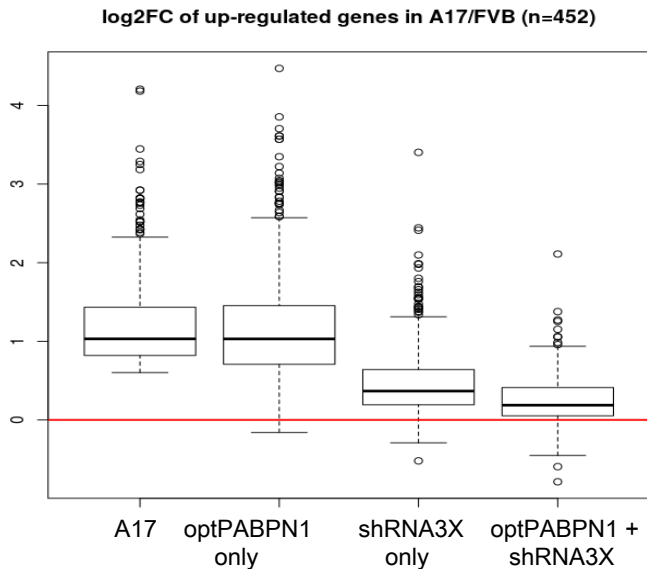
Reduced Fibrosis +
Partial Reversal of Atrophy



Specific force calculated by normalizing
maximal force for muscle weight

Impact of Treatment of A17 Mice: a Microarray Analyses

- expPABPN1 expression in A17 mice causes extensive remodelling of muscle transcriptome (Trollet et al. *HMG* 2010; Anvar et al. *Sk Muscle* 2011; Chartier et al. *Plos Genet* 2015)
- Transcriptome analyzed from current experiment



- In A17 mice vs wildtype, 865 transcripts were deregulated $FC > 2$; $p < 0.05$
- Dual treatment with shRNA3X + optPABPN1 results in only 12 genes deregulated, a 98% “correction”

- ddRNAI constructs design highlights unique 'silence and replace' therapeutic strategy for monogenic disease
- Neither overexpression of wildtype protein or knockdown on disease protein impacts disease. Only a combined approach efficiently restores function.
- A single vector approach simplifies the CMC approach for clinical materials
- Continuing to characterize *in vivo* efficacy of clinical candidates
- Orphan Drug Designation granted from the EU January 2017



Centre for Biomedical Sciences

George Dickson

Alberto Malerba

Houria Bachtarzi

Susan Jarmin



Vanessa Strings

Sonal Harbaran

Michael Graham



Myology Research Center, UMRS974 (Paris)

Capucine Trollet

Pierre Klein

Gillian Butler-Browne