#### University of Massachusetts Medical School

#### eScholarship@UMMS

UMass Center for Clinical and Translational Science Research Retreat 2013 UMass Center for Clinical and Translational Science Research Retreat

May 8th, 3:30 PM - 5:00 PM

#### Molecular Mechanisms of FSH Muscular Dystrophy Pathogenesis

Peter L. Jones University of Massachusetts Medical School Worcester Et al.

#### Let us know how access to this document benefits you.

Follow this and additional works at: https://escholarship.umassmed.edu/cts\_retreat

Part of the Cancer Biology Commons, Cell Biology Commons, Molecular Biology Commons, Molecular Genetics Commons, Musculoskeletal Diseases Commons, Nervous System Diseases Commons, Neurology Commons, and the Translational Medical Research Commons

Jones PL, Jones TI. (2013). Molecular Mechanisms of FSH Muscular Dystrophy Pathogenesis. UMass Center for Clinical and Translational Science Research Retreat. Retrieved from https://escholarship.umassmed.edu/cts\_retreat/2013/presentations/15

Creative Commons License



This work is licensed under a Creative Commons Attribution-Noncommercial-Share Alike 3.0 License.

This material is brought to you by eScholarship@UMMS. It has been accepted for inclusion in UMass Center for Clinical and Translational Science Research Retreat by an authorized administrator of eScholarship@UMMS. For more information, please contact Lisa.Palmer@umassmed.edu.







# Molecular mechanisms of FSH muscular dystrophy pathogenesis

Peter L. Jones, Ph.D. and Takako I. Jones, Ph.D. Principal Investigators

Departments of Cell and Developmental Biology & Neurology

# Facioscapulohumeral Muscular Dystrophy (FSHD)

Most prevalent muscular dystrophy afflicting children and adults (~1:7,000-15,000)

Autosomal dominant

Facio: refers to face

Scapulo: refers to shoulders

**Humeral:** refers to humerus (upper arm bone)



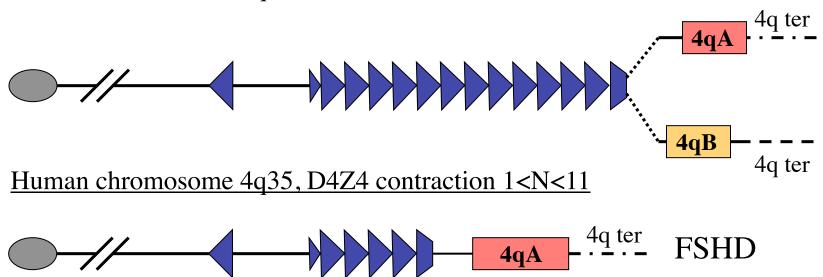
©2003 shoulderdoc.co.uk

Winging on both sides in a patient with FSHD due to weakness of all the scapula stabilizing muscles

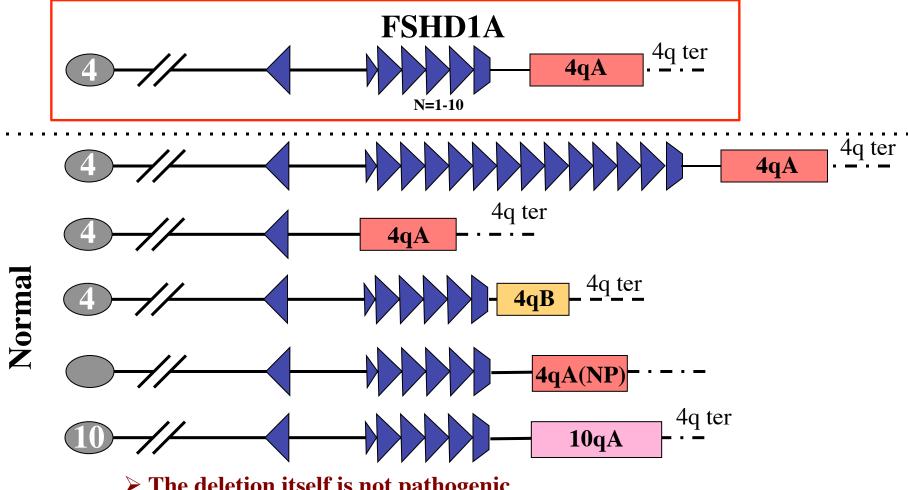
Great genetic and clinical heterogeneity
Each patient may differ in severity
Most patients exhibit symptoms by age 20
>50% of patients retain ability to walk

## The FSHD1 genetic lesion is a deletion in a tandem repeat array at 4q35

Human chromosome 4q35, Normal D4Z4 n=11-150

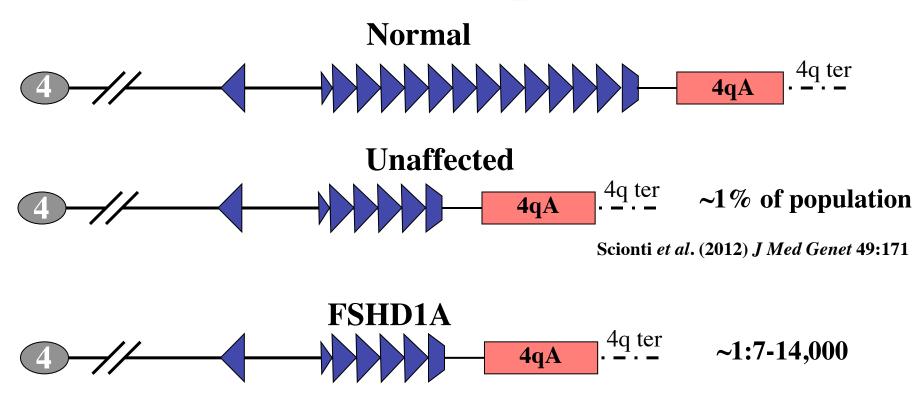


### A pathogenic FSHD1 deletion is complex



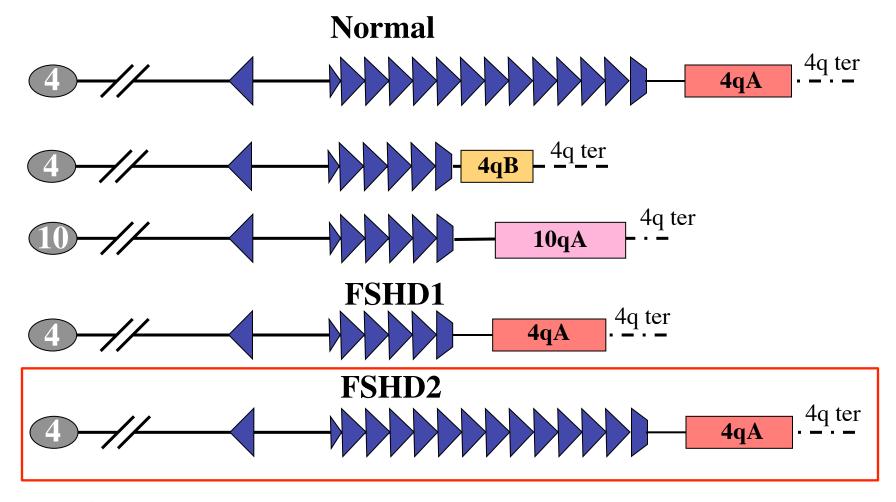
- **➤** The deletion itself is not pathogenic
- ➤ The 4qA sub-telomere is permissive, not pathogenic
- **➤** The 4qB sub-telomere is not permissive
- > Requires at least 1 D4Z4 repeat unit for FSHD
- > Chromosome 10 arrays (devoid of chr. 4 D4Z4) are not linked to FSHD

## A putatively pathogenic FSHD1 deletion shows very low penetrance



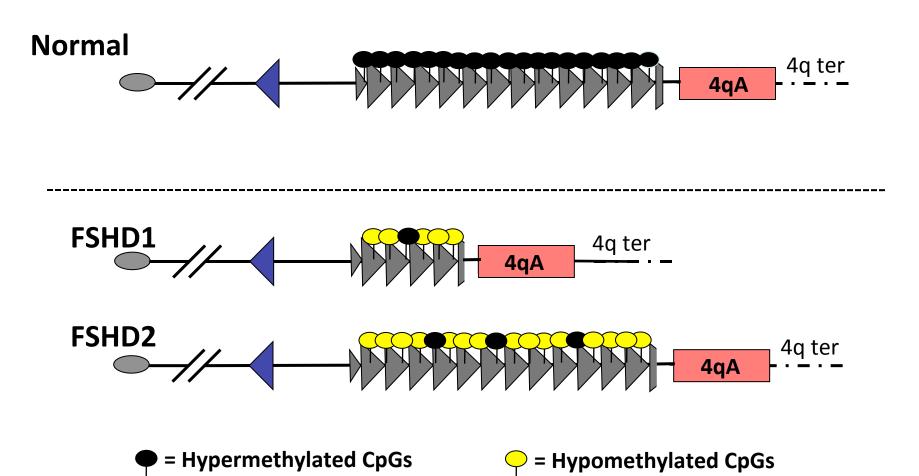
The deletion itself is not pathogenic The 4qA sub-telomere is permissive, not pathogenic

### FSHD2 is independent of the contraction



- > The 4qA sub-telomere is required for FSHD1 and 2
- ➤ At least 1 D4Z4 is required for FSHD1 and 2

## FSHD is linked to D4Z4, the A type subtelomere and the epigenetic status of the 4q35 D4Z4 repeat



more euchromatic

more heterochromatic

# The 3 types of FSHD are linked by epigenetic dysregulation

FSHD1: Dominant deletions at 4q35 D4Z4 array Apparently low penetrance <a href="DNA Hypomethylation">DNA Hypomethylation</a> of shortened 4q35

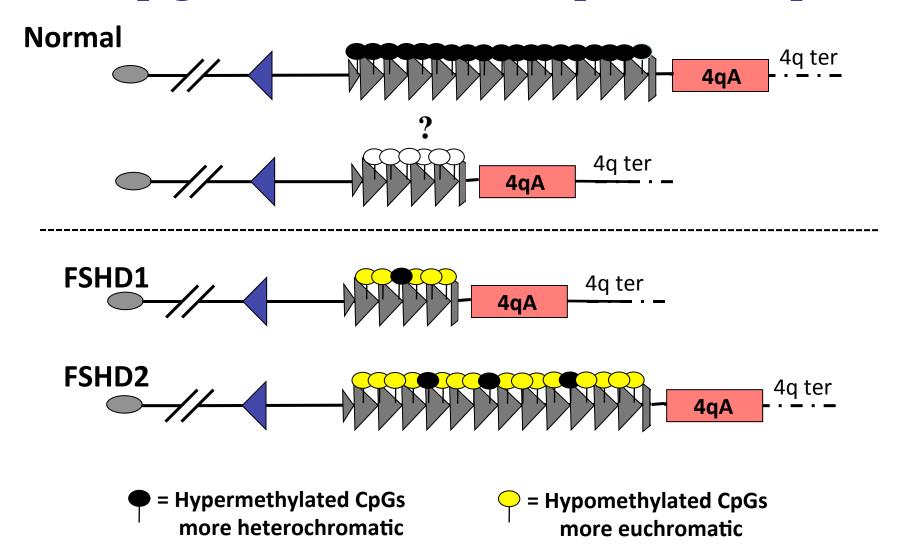
FSHD2: SMCHD1 inactivating mutations
--ATPase chromatin remodeling protein
--Modifier of metastable epialleles

<u>DNA Hypomethylation</u> of 4q35 and 10q26 arrays

IFSHD: Infantile form of FSHD1 or FSHD2, much more severe <u>DNA Hypomethylation</u> of FSHD1 or 2



## FSHD is linked to the A type subtelomere <u>and</u> the epigenetic status of the 4q35 D4Z4 repeat



# FSHD results from an epigenetic-mediated dysregulation of gene repression

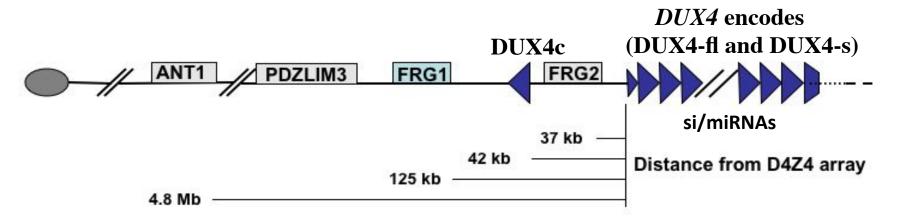
- > DNA methylation
- > Histone modifications
- > Chromatin structure
- **Long non-coding RNAs**
- > Nuclear organization
- > High variability within the clinical population

**Severity** 

Age of onset

**Gene expression** 

# Which gene(s) is responsible for FSHD pathology?

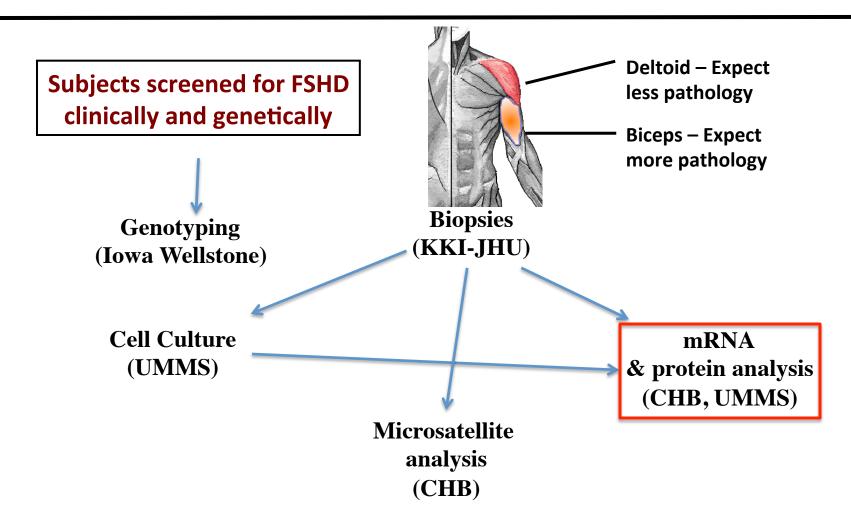


#### The FSHD causal gene:

- 1. Should be misexpressed in FSHD (Up or down)
  - → mRNA, protein, cell type, developmental timing
  - → adversely affect skeletal muscle and potentially vasculature
- 2. Explain the 4qA linkage and under epigenetic repression

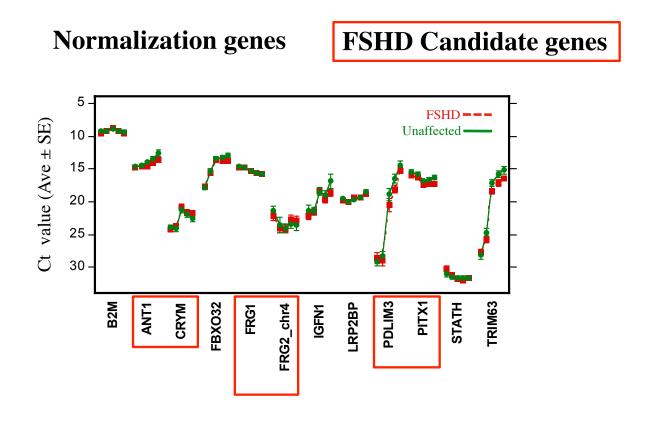
Both FRG1 and DUX4 produce phenotypes consistent with FSHD when overexpressed in animal models

## Wellstone family cohorts of muscle biopsies and myogenic cell cultures



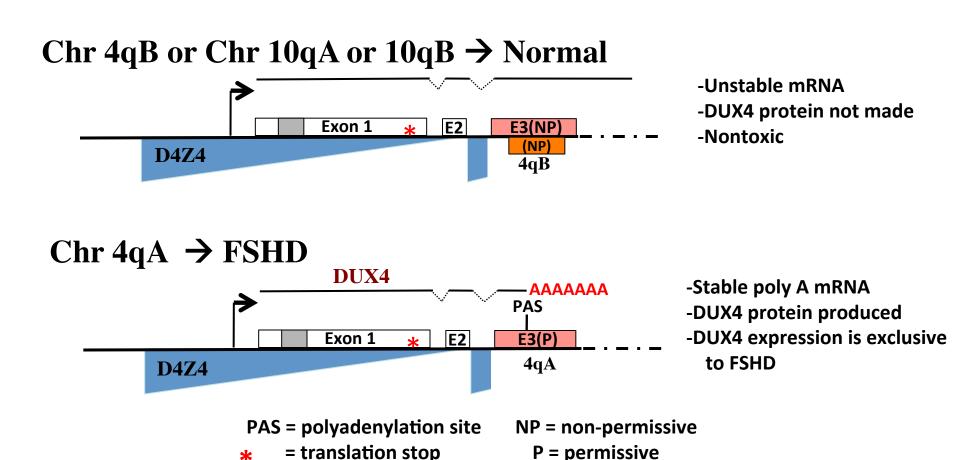
FSHD1 affected and genetically unaffected 1st degree relatives

### Myogenic cultures from FSHD and unaffected firstdegree relatives have similar patterns of gene expression during proliferation and myogenic differentiation



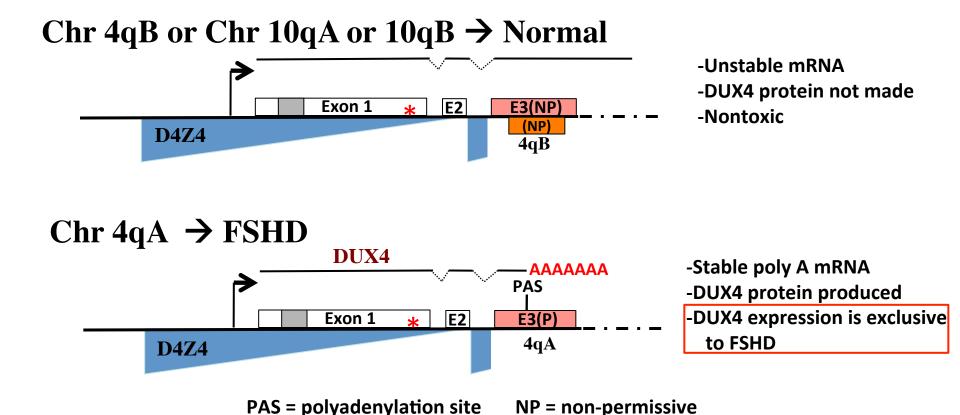
#### "A Unifying Model for FSHD" based on DUX4

FSHD permissive 4qA subtelomeres encode a third exon containing a polyadenylation site that stabilizes the DUX4 mRNA



#### "A Unifying Model for FSHD" based on DUX4

FSHD permissive 4qA subtelomeres encode a third exon containing a polyadenylation site that stabilizes the DUX4 mRNA

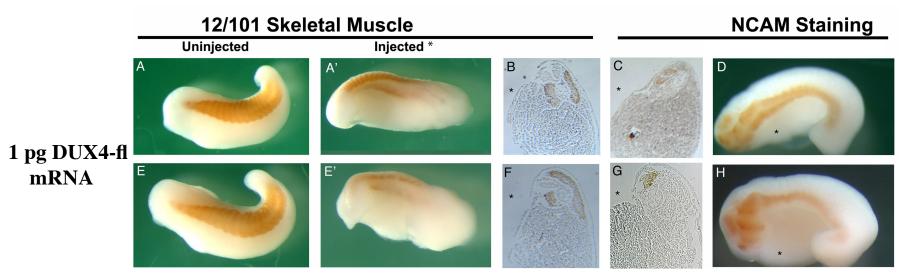


P = permissive

= translation stop

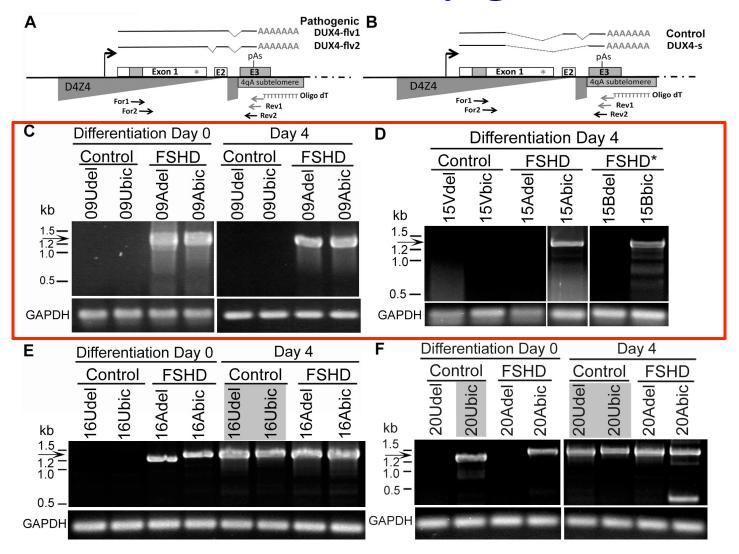
## DUX4-FL expression leads to massive loss of developing myogenic cells

#### Xenopus embryos

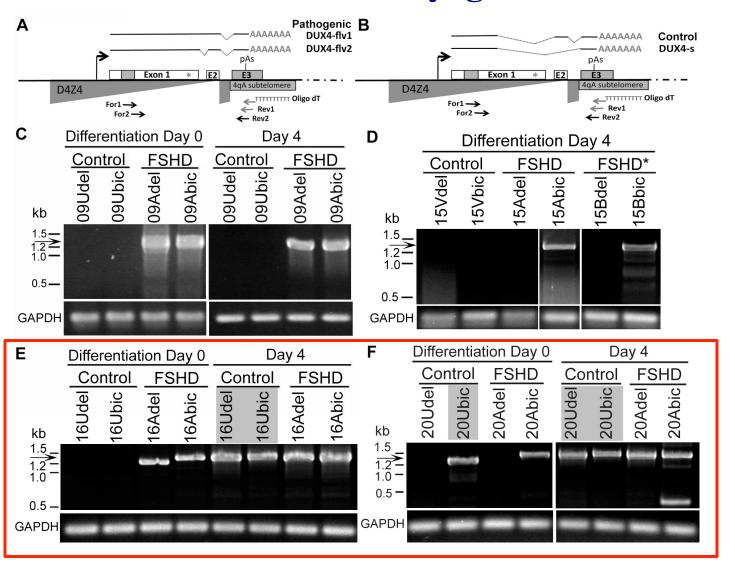


Immunostaining for developing skeletal muscle

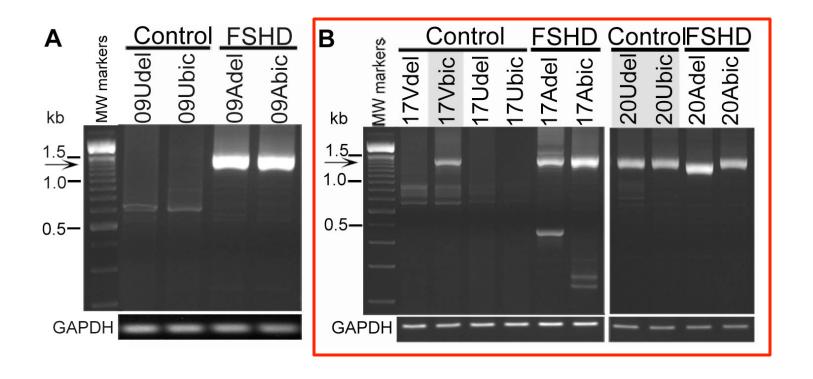
## DUX4-fl mRNA is expressed in FSHD1-derived myogenic cells



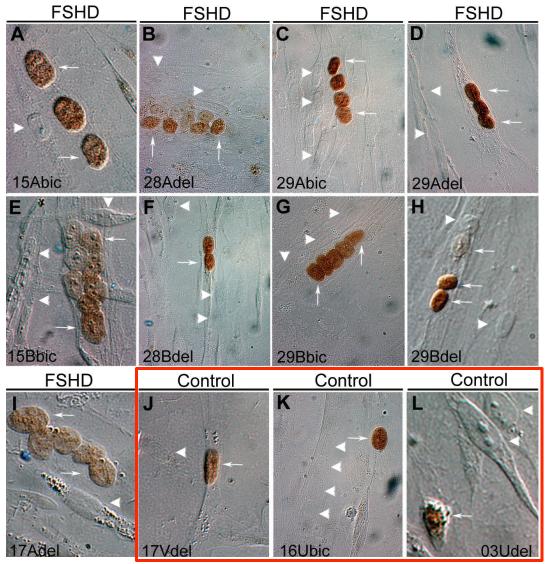
## DUX4-fl mRNA expression is not exclusive to FSHD1-derived myogenic cells



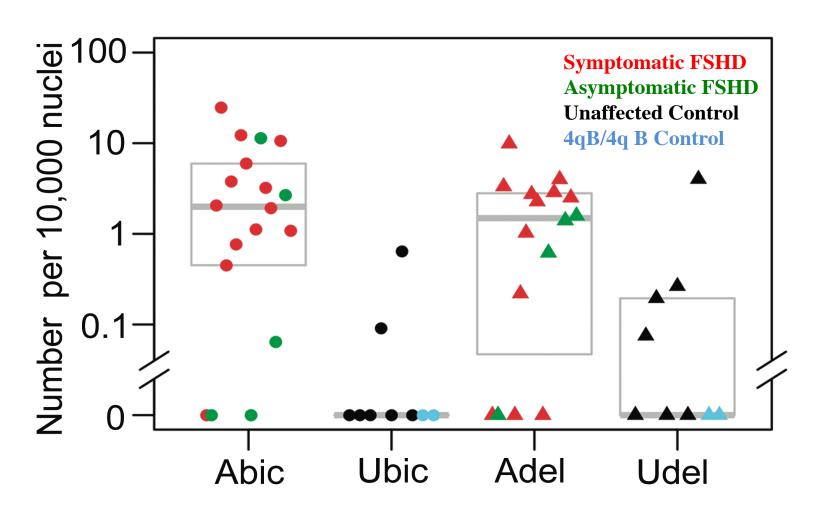
### DUX4-fl mRNA expression is expressed in both FSHD1-derived and unaffected muscle biopsies



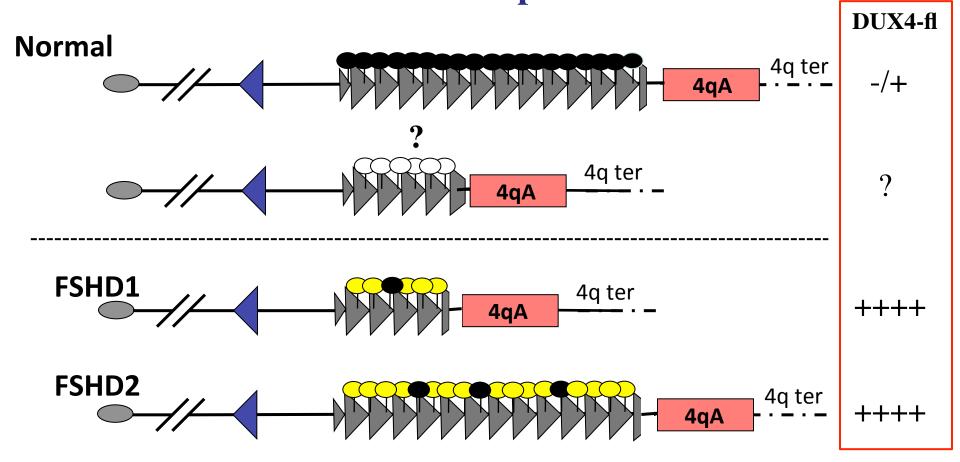
## Differentiated myogenic cells from genetically FSHD1 and control subjects express DUX4-FL protein



# DUX4-FL expression in myogenic cells from FSHD affected unaffected subjects suggests a quantitative model of pathogenesis

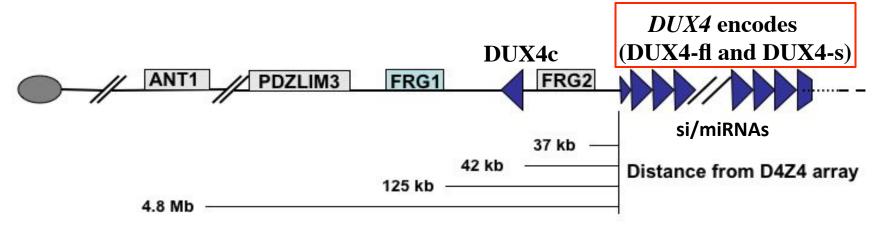


FSHD is linked to the permissive 4qA subtelomere, the epigenetic status of the 4q35 D4Z4 repeat and DUX4-fl expression



= Hypermethylated CpGs more heterochromatic = Hypomethylated CpGs more euchromatic

# Which gene(s) is responsible for FSHD pathology?

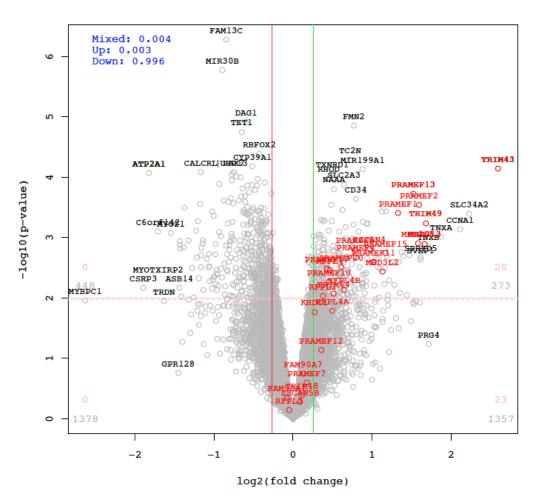


#### The FSHD causal gene:

- 1. Should be misexpressed in FSHD (Up or down)
  - → mRNA, protein, cell type, developmental timing
  - → adversely affect skeletal muscle and potentially vasculature
- 2. Explain the 4qA linkage and under epigenetic repression

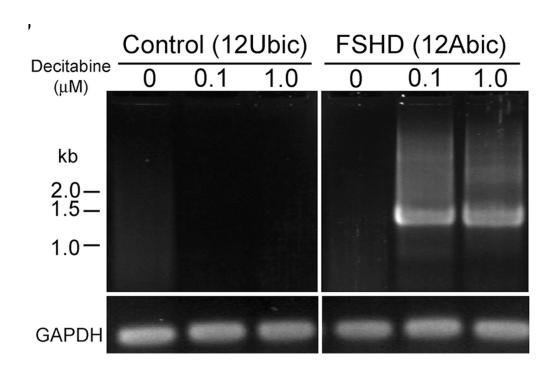
**DUX4** fulfills these criteria

# Many DUX4-FL responsive genes are upregulated in FSHD myotubes (biomarkers)



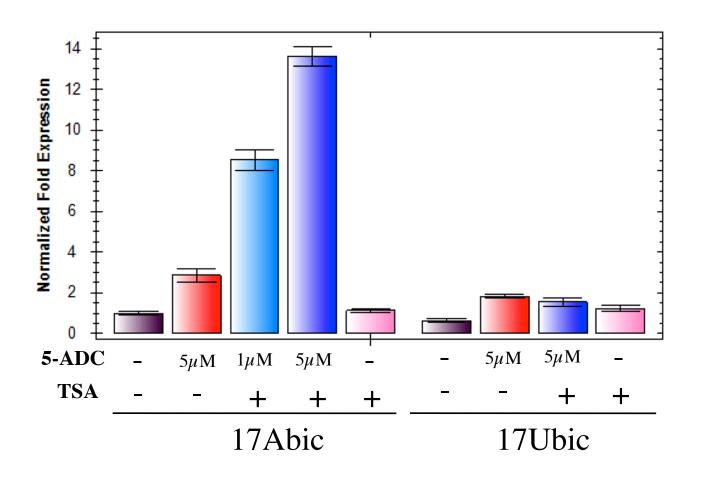
# Increased DUX4-fl expression appears necessary\* but alone is not sufficient for FSHD

## FSHD-derived myoblasts are epigenetically poised to express DUX4-fl



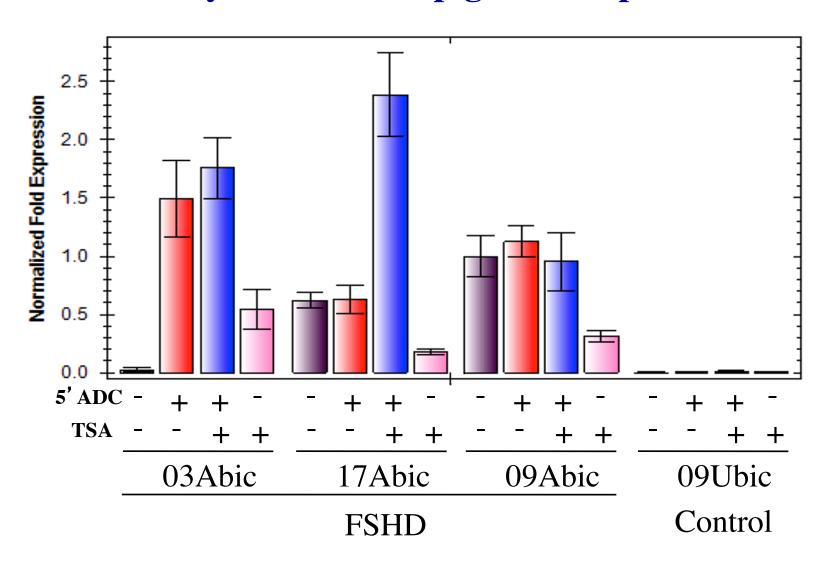
Decitabine treatment leads to DNA demethylation

## FSHD derived myogenic cells are epigenetically poised to express DUX4-fl mRNA



DUX4-fl qRT-PCR Analysis

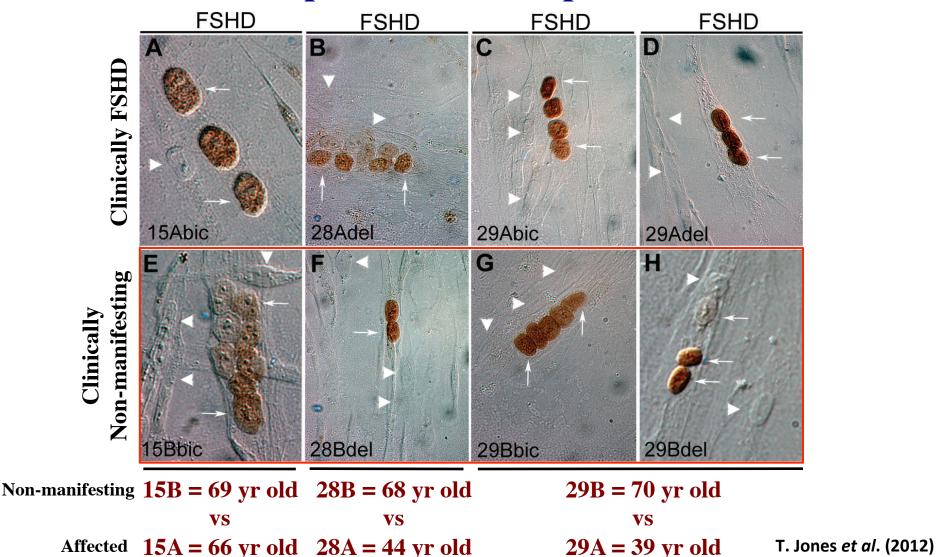
## FSHD subjects show individual variability in the stability of DUX4-fl epigenetic repression



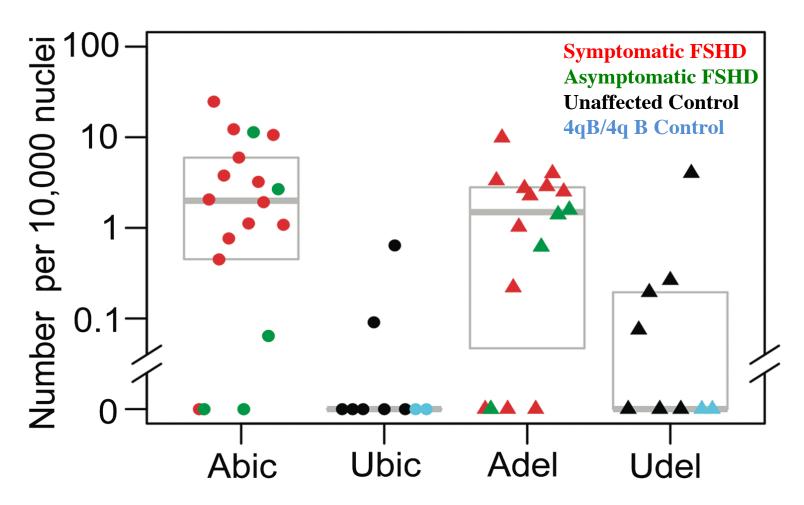
# The 4q35 D4Z4 in FSHD exists as differentially metastable epialleles among affected subjects → epigenetically poised for DUX4 expression

The 4q35 D4Z4 in normal subjects exhibits stable epigenetic repression

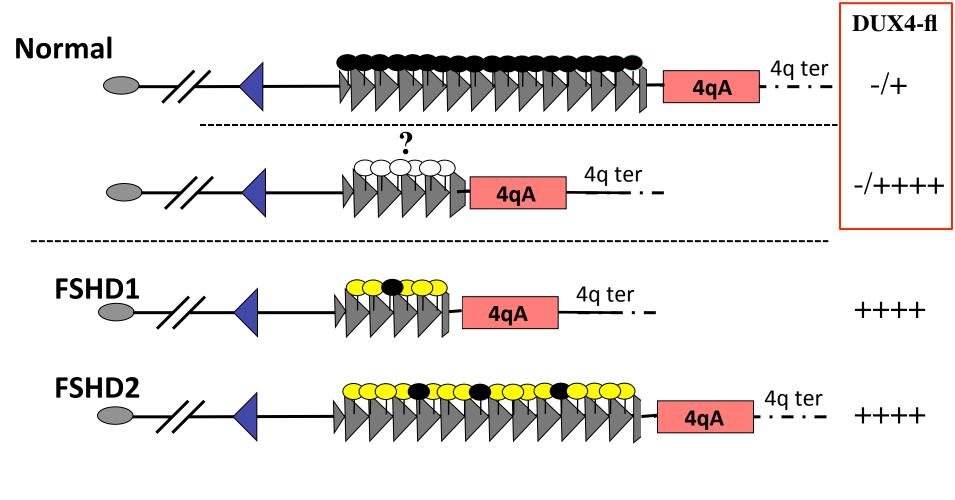
# Differentiated myogenic cells from genetically FSHD1 but clinically non-manifesting subjects express DUX4-FL protein



# DUX4-FL expression in myogenic cells from FSHD1 subjects that show no clinical manifestation of the disease suggests modifiers of disease

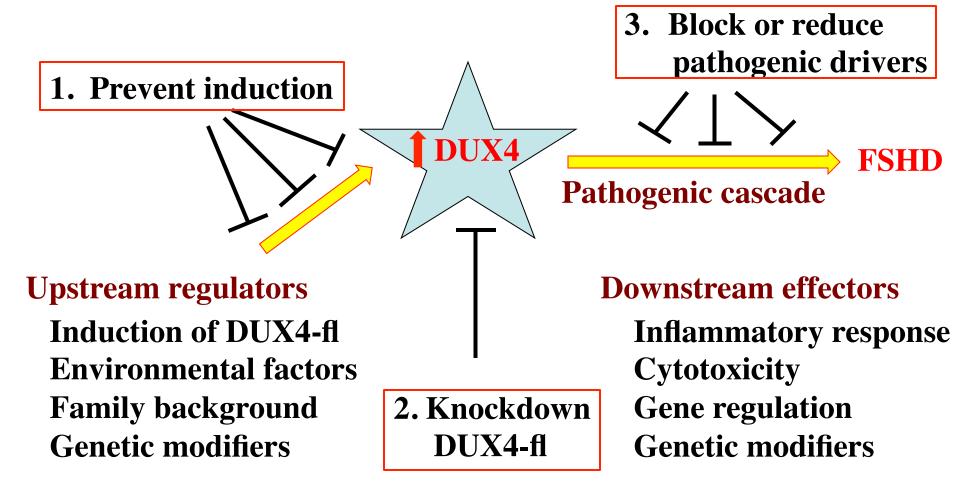


## DUX4 expression alone is not necessarily causal for FSHD

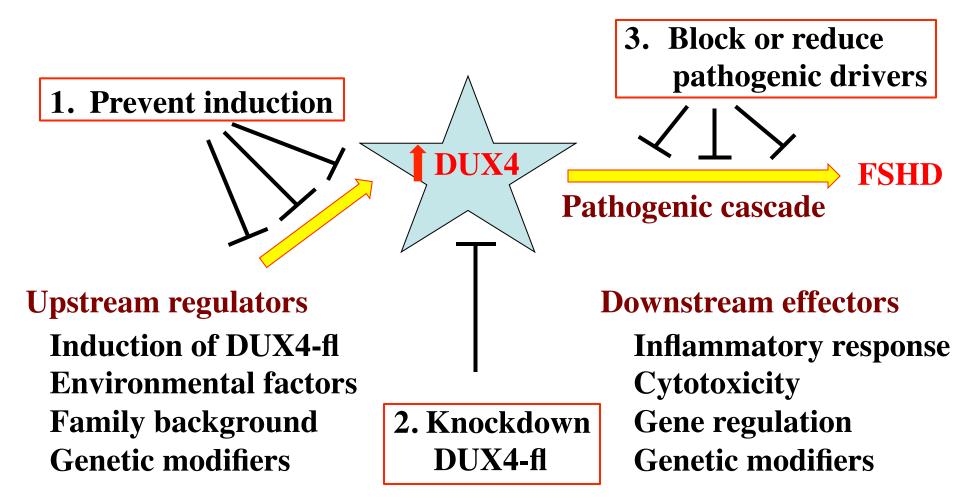


= Hypermethylated CpGs more heterochromatic = Hypomethylated CpGs more euchromatic

# Multiple therapeutic targets for FSHD



# Multiple therapeutic targets for FSHD



Key to therapy may lie with identifying the disease modifiers

# Analysis of DUX4 mRNA and protein expression in muscles and myogenic cells from FSHD subjects and unaffected relatives

At University of Massachusetts Medical School (and formerly BBRI)

Takako Jones\*, Jennifer Chen\*, Oliver King, Charles P. Emerson Jr.,
and Peter L. Jones

At Kennedy Krieger Institute and Johns Hopkins University Kathryn R. Wagner

At Children's Hospital – Boston and Harvard Medical School Fedik Rahimov and Louis M. Kunkel

At Boston University School of Medicine (and formerly BBRI)
Sachiko Homma, Mary Lou Beerman and Jeffrey Boone Miller











### Acknowledgements







Takako Jones









