U.S. DEPARTMENT OF HEALTH AND HUMAN SERVICES

National Institutes of Health

National Cancer Advisory Board

Understanding Telomere Biology Through Studies of Dyskeratosis Congenita

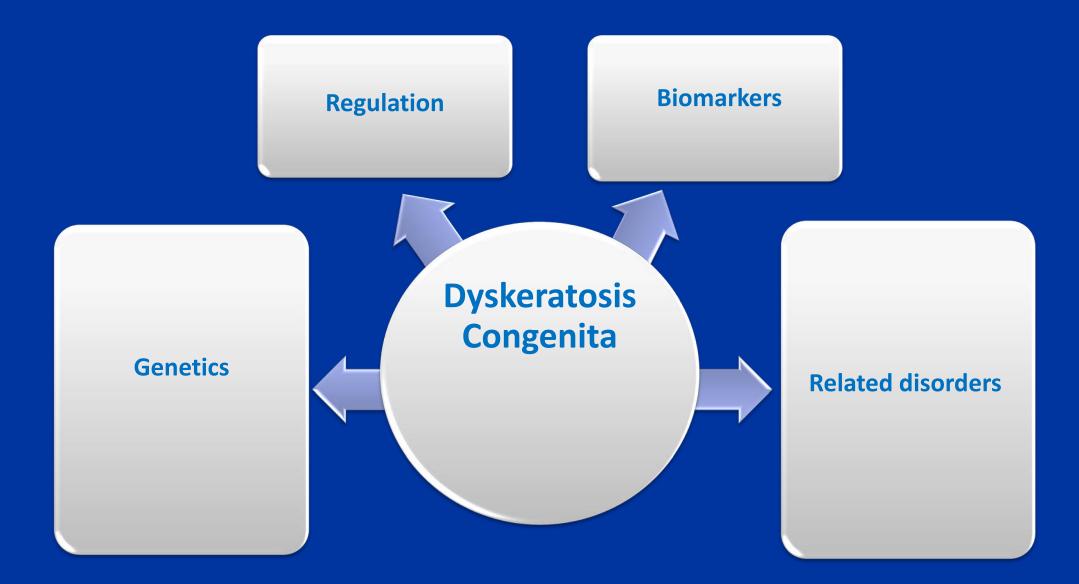
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Clinical Genetics Branch Division of Cancer Epidemiology and Genetics

December 7, 2010

Dyskeratosis congenita:

a highly informative model of telomere biology and cancer predisposition



Telomeres Preserve Chromosomal Integrity

- Long nucleotide (TTAGGG)_n repeats at chromosome ends
- Many proteins regulate telomere length and stabilize structure
- Shorten with each cell division
- Erosion causes genetic instability, cell crisis and cell death
- Cancer cells survive despite critically short telomeres and chromosomal instability

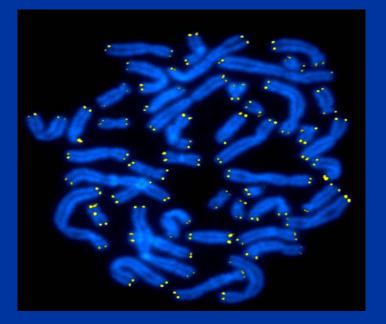


Photo: Dr. Peter Lansdorp

Evolutionary Conservation of Telomere Biology Genes

- *Hypothesis:* Genetic variation in telomere biology genes is limited because of their critical roles in genomic stability
- Population genetic study showed that these genes are highly conserved between ethnic groups and species

Savage SA et al, <u>Human Mutation</u> 2005;26(4):343-350

Dyskeratosis Congenita: a disorder of telomere biology

Nail Dystrophy

Oral Leukoplakia

Skin Pigmentation











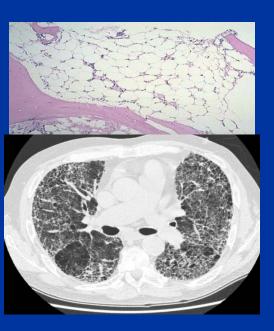
In addition to the triad...

Bone Marrow Failure

Pulmonary Fibrosis

Head and Neck Cancer Anogenital Cancer Leukemia

Liver Fibrosis Dental Disease Hyperhidrosis Esophageal Stenosis Urethral Stenosis Cerebellar Hypoplasia Tear duct stenosis



Microcephaly Osteoporosis Avascular Necrosis Developmental Delay Early Gray Hair Hair Loss Abnormally <u>short</u> telomeres are the unifying feature

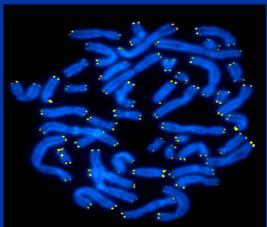
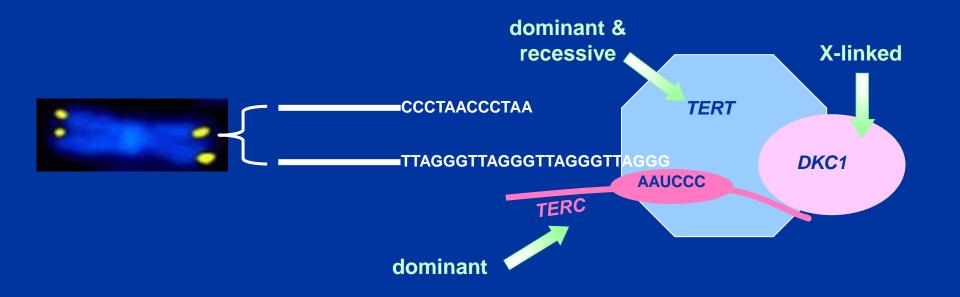


Photo: Dr. Peter Lansdorp

Traditional diagnosis: Diagnostic Triad <u>or</u> 1 of the triad, + BMF + 2 other findings, Vulliamy et al, *Blood*, 2006, 107(7):2680-5

Dyskeratosis Congenita Genetics: The Telomere Connection (c. 2006)



These mutations result in short telomeres, but only ~40% of patients have a mutation in one of these genes.

Inherited Bone Marrow Failure Syndromes



What is the NCI IBMFS Cohort and Who is Eligible Etiologic Investigation of Cancer Susceptibility in Inherited Bone Marrow Failure Syndromes (IBMFS)

- Systematic evaluation of affected individuals and their families in North America
 - Fanconi Anemia
 - Dyskeratosis Congenita (DC)
 - Diamond-Blackfan Anemia
 - Shwachman-Diamond Syndrome
 - And others
- Opened in January 2002
- http://marrowfailure.cancer.gov
- Principal Investigator: Blanche P. Alter, MD, MPH
- Staff Clinician: Neelam Giri, MD
- DC: Sharon Savage, MD

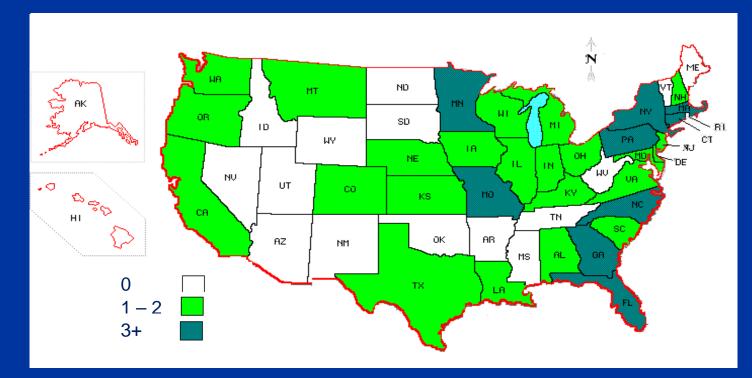
IBMFS Study: Levels of Participation

- Epidemiology Questionnaires
 - Family history
 - Individual history
- Medical Record Review
- Evaluation at the NIH Clinical Center
 - IBMFS Team
 - Genetic Counseling
 - Subspecialists
 - Biospecimens



Dyskeratosis Congenita Families

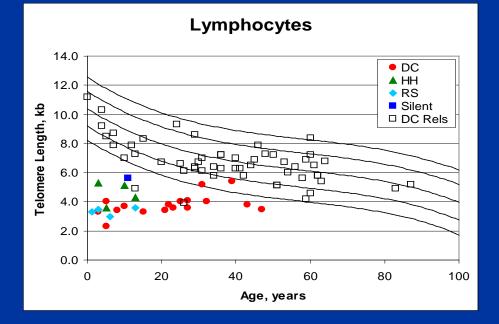
NCI Cohort	# Families	# Affected Individuals	# Healthy Relatives
Clinic	35	46	86
Field	11	24	101



Development of the First (and only) Diagnostic Test for DC

- Varied clinical presentations
- Signs and symptoms often progress with age
- <50% have an identifiable mutation
- Telomeres are very short for their age

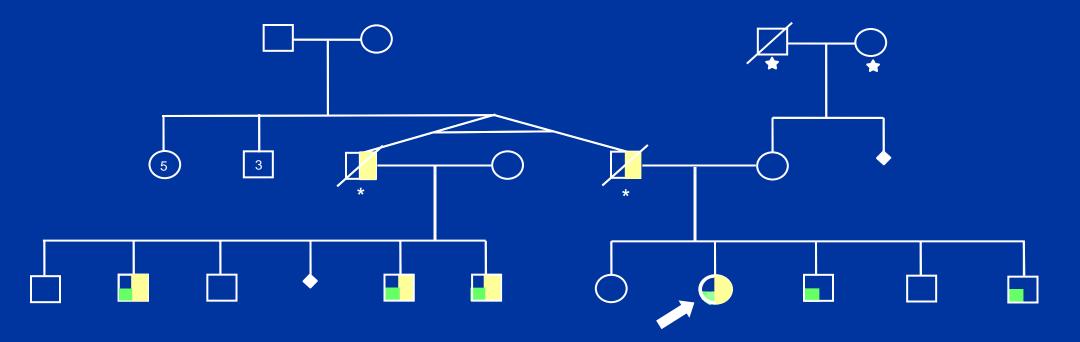
DC vs. healthy relatives



DC vs. other IBMFS	Sensitivity	Specificity
If 4/6 leukocyte subsets <1 st percentile	92%	98%

Genome-Wide Linkage Screen in DC

Heterogeneous, autosomal dominant, DKC1, TERC, and TERT normal



Clinical DC

Telomere length in WBC subsets <1st %ile

Spontaneous abortion

* Telomere length unknown

TINF2 mutations cause DC

- Evidence for linkage at 14q11.2 led to the identification of mutations in *TINF2*, a component of the shelterin telomere protection complex
 - Mutations present only in those with telomeres <1st percentile
 - No mutations in 298 healthy controls
 - Additional unrelated individuals with DC had mutations
- Mutations located in a small, highly conserved region of exon 6

Savage et al., Am J Hum Genet. 2008;82(2):501-9

Telomerase complex

TINF2

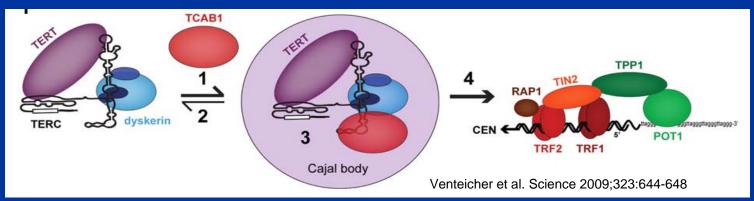
TINF2

Sequence Analysis of the Shelterin Telomere Protection Complex

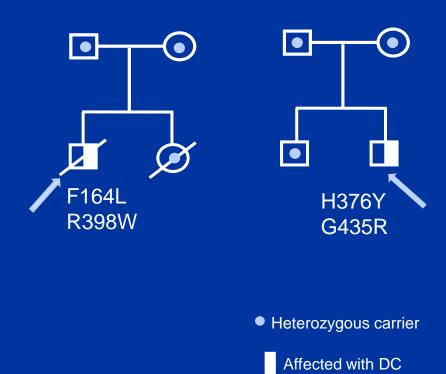
- TERF1, TERF2, POT1, TERF2IP, and ACD sequenced in 9 patients with DC, and 7 DC-like
- Variants in patients were compared to 380 control subjects
- Mutations in these genes do not appear to be a common cause of dyskeratosis congenita.

Savage et al, <u>J Med Genet</u> 2010: in press

TCAB1 in Dyskeratosis Congenita

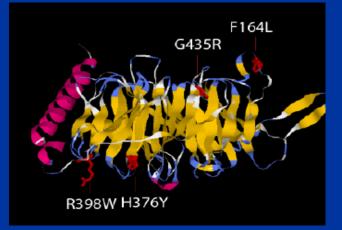


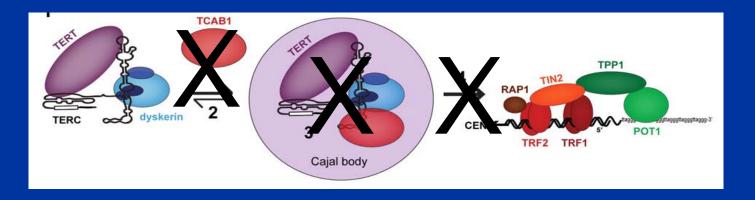
- Cajal body: sites of mRNA & rRNA processing in the nucleus
- *TCAB1* is located in Cajal bodies and is required for telomerase trafficking
- Mutations in TCAB1 cause DC
 - 2 of 9 families studied have autosomal recessive mutations



DC-associated TCAB1 Mutations Result in Defective Telomerase Trafficking

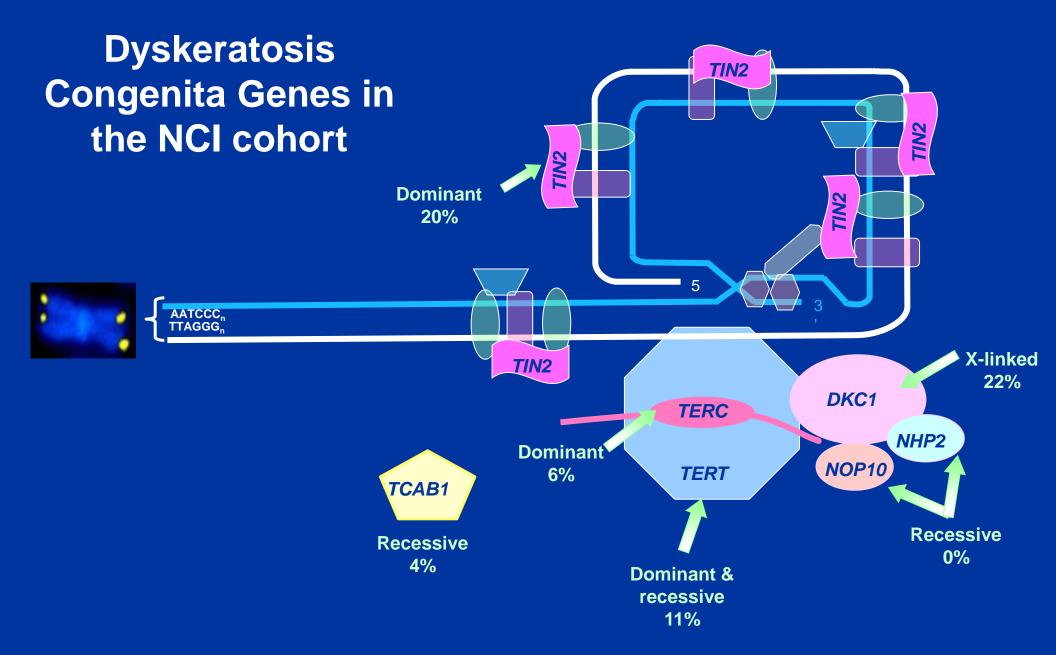
- Occur in conserved loop residues
- Reduce TCAB1 levels in Cajal bodies
- Prevents localization of the telomerase enzyme complex (*TERT*, *TERC*, *DKC1*) to Cajal bodies



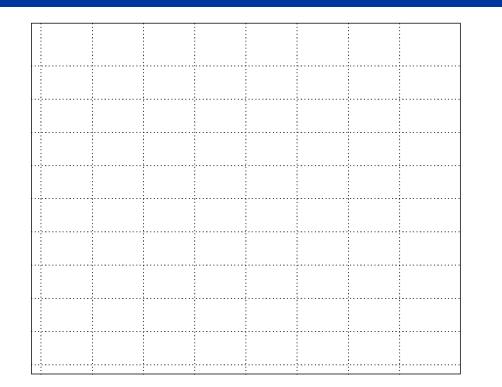


Zhong F, Savage SA, et al, Genes Devel 2010; in press

Study highlighted in the "Best of" session at the American Society of Hematology annual meeting



~ 1/3 of our families do not have a mutation in one of these genes



Refining the Phenotype of DC

- Detailed clinical characterization at the NIH Clinical Center
 - Dental
 - ENT
 - Ophthalmology
 - Radiology
 - Psychiatry
 - Pulmonology
 - And others



Atkinson et al, <u>Oral Dis</u> 2008;14(5):419-27.

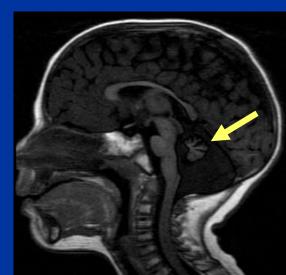


Tsilou et al, <u>Ophthalmology</u>. 2010;117(3):615-22

Tear duct present

Tear duct absent

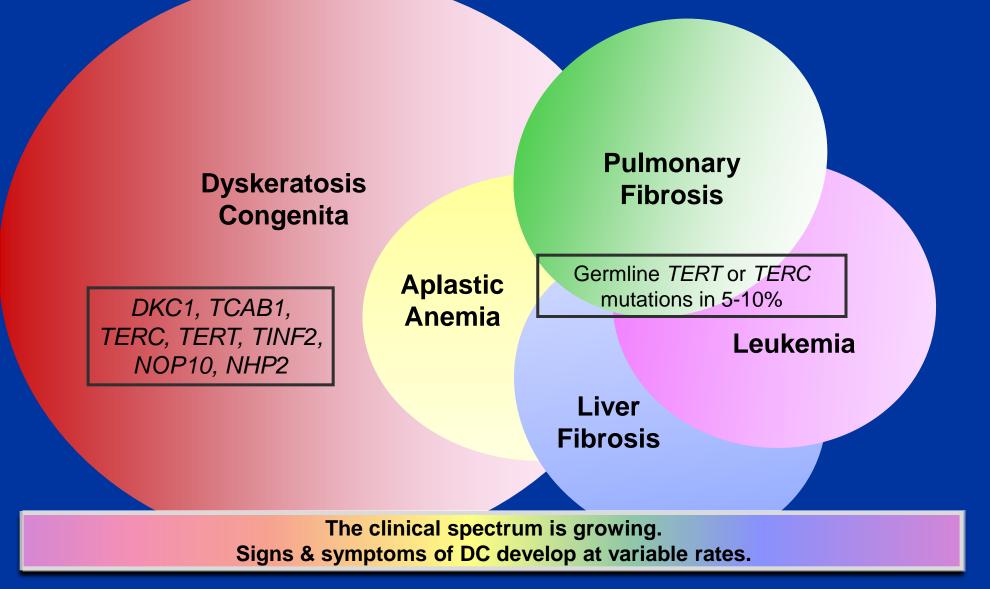
Unpublished data on MRI findings in the brain: John Butman, MD, Clinical Center, NIH, Neelam Giri, MD, Clinical Genetics Branch, DCEG, NCI



Major Contributions from NCl's Dyskeratosis Congenita Study

- Developed the diagnostic test
 - Telomere length
- Discovered 2 of the 7 causative genes
 - TINF2 : The first component of the shelterin telomere protection complex associated with any disease
 - TCAB1: The first study showing that telomerase mislocalization can cause disease
- Quantified cancer risk
- Refined the extent of medical complications
- Helped families create a support group: DC Outreach
- Created basis for population-based studies of the contribution of aberrations in telomere biology to cancer risk

Telomere biology disorders overlap clinically and genetically

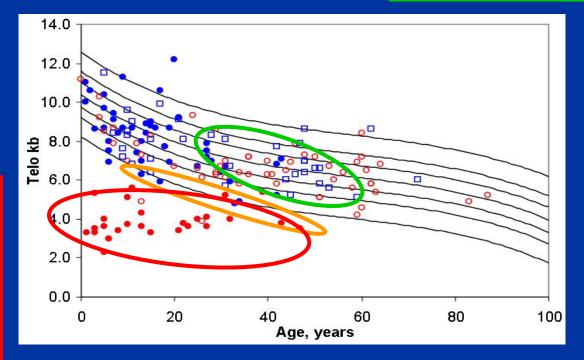


Savage and Bertuch, <u>Genetics in Medicine</u> 2010: in press

The Telomere Length Continuum

Normal range

Marker of common disease riskStatistically significant differences



Short, <10th%

•Aplastic anemia, leukemia, pulmonary fibrosis •Mutations in *TERT, TERC*

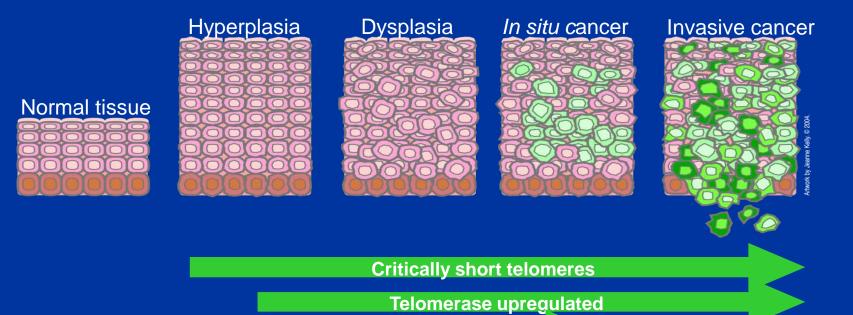
VERY short, <1st%

Dyskeratosis Congenita

•Rare, highly penetrant disease causing mutations

Telomere Length and Cancer Risk in the General Population

 Individuals in the general population with shorter germline telomeres may have increased cancer risk because of telomere dysregulation

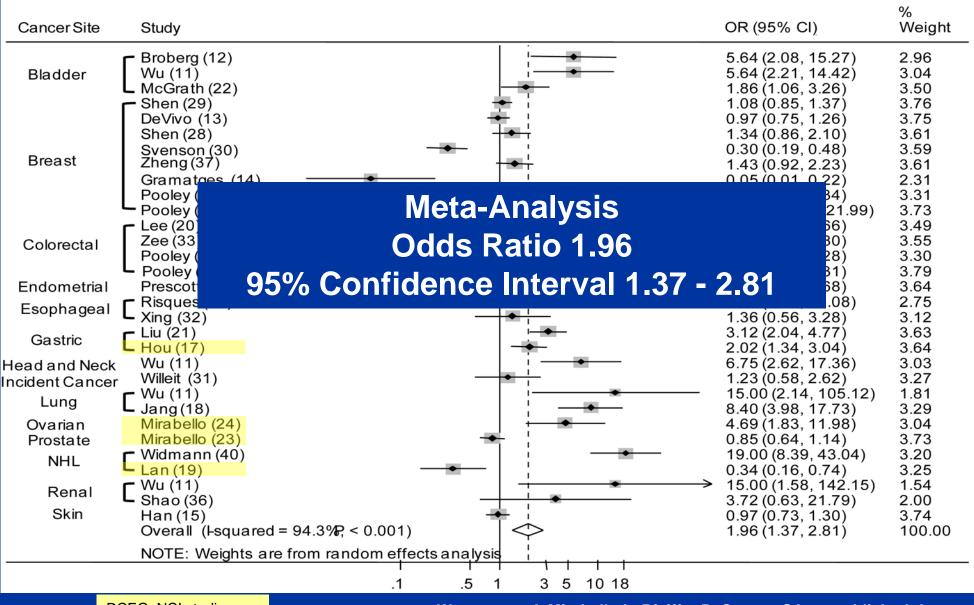


Cellular crisis bypassed

Chromosomal instability

Chromosomal abnormalities

Surrogate Tissue Telomere Length and Cancer Risk

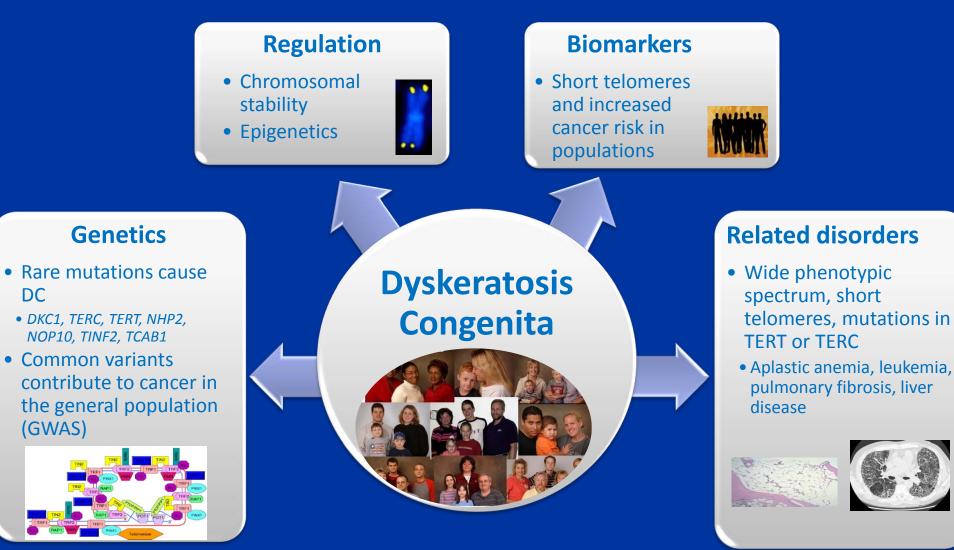


DCEG, NCI studies

Wentzensen I, Mirabello L, Pfeiffer R, Savage SA: unpublished data

Dyskeratosis congenita:

a highly informative model of telomere biology and cancer predisposition



Photos with permission

Acknowledgements & Collaborators

- Patients & Families
 - www.dcoutreach.com



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 - June Peters, CGC

Genomics

igodol

- Exome sequencing and gene discovery: CGF and LMT
- Copy number variants: P.
 NCI, CCR

Telomere Length Flow-FISH

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