



Glycosaminoglycans and diseases

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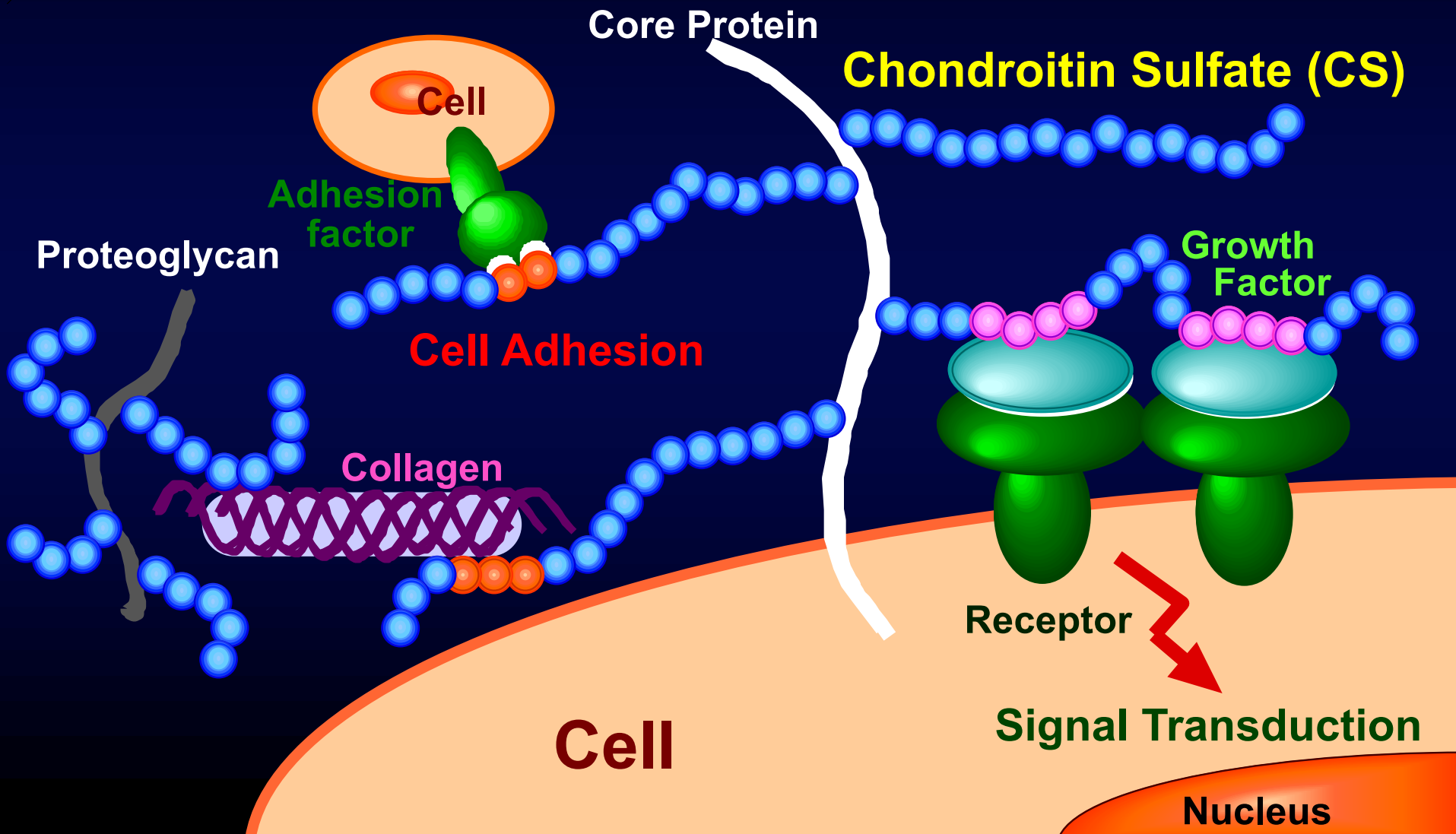
Faculty of Pharmacy,

Meijo University

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- Introduction
- Cancer and GAGs
- Spinal Cord Injury
- Infection of Pathogens via GAGs
- Mucopolysaccharidoses
- **Congenital Disorders Caused by Defects in Biosynthetic Enzymes of GAGs**

Functions of Cell Surface and Extracellular Matrix Proteoglycans



Glycosaminoglycans

- Chondroitin sulfate
- Dermatan sulfate
- Heparin
- Heparan sulfate
- Keratan sulfate
- Hyaluronan

Glycosaminoglycans

Polysaccharides composed of repeating disaccharide units containing an amino sugar and modified with sulfate groups.

Chondroitin Sulfate...Cartilage

Dermatan Sulfate...Skin, Aorta

Heparan Sulfate...Basement membrane

Heparin...Anticoagulant drug

Keratan Sulfate...Cornea

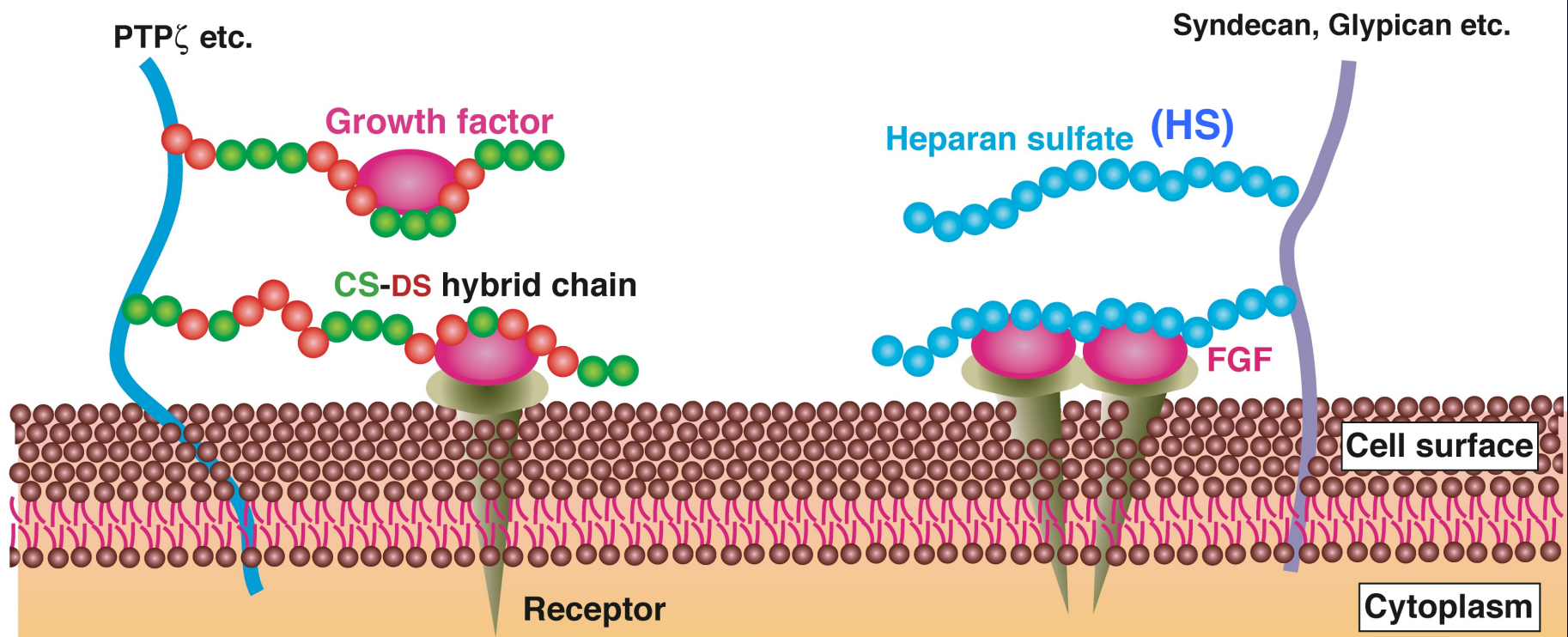
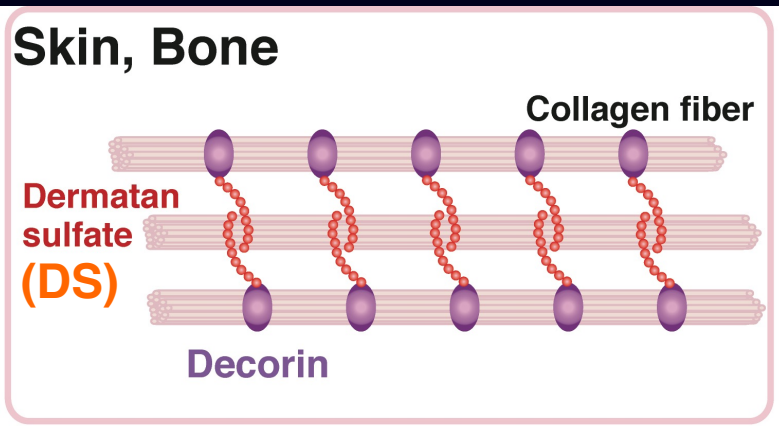
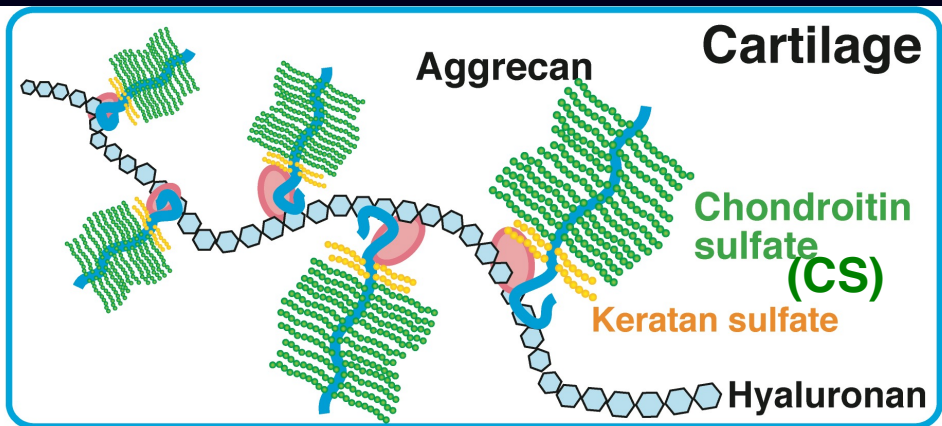
Functions of Proteoglycans

Interact with various bioactive proteins including cell growth factors, morphogens, cytokines, extracellular matrix components, nerve growth factors, coagulation factors, etc.



Exhibit various functions including cell growth, cell adhesion, cell migration, anticoagulation, morphogenesis, tissue regeneration etc.

Various functions of CS, DS, and HS-proteoglycans at cell surfaces and in extracellular matrix



Protein Factors interacting with GAGs

Cell Adhesion Molecules

CD44
L-Selectin
P-Selectin
RANTES
von Willebrand factor
MAC-1
N-CAM

Growth Factors/Morphogens

Fibroblast growth factors
Hepatocyte growth factor
Midkine/Pleiotrophin
Platelet derived growth factor
Vascular endothelial growth factor
Transforming growth factor- β
Glial-derived neurotrophic factor
Brain-derived neurotrophic factor
HB-EGF
Amphiregulin
Neuregulin
Insulin-like growth factor
Bone morphogenetic proteins
Sonic hedgehog
Wnts

Virus protein

Glycoprotein C

Coagulation

Heparin cofactor II
Antithrombin III
Factor Xa
Leuserpin
Thrombin

ECM Components

Tenascin-X
Opticin
Fibronectin
Collagens
Laminins
Tenascin
Thrombospondin
Vitronectin

Proteinases

Elastases
Cathepsin G

Chemokines

Interferon- γ
Interleukin-2, -3, -4, -5, -7, -8, -12
Macrophage inflammatory peptides 1
Monocyte chemoattractant protein-1
Secondary lymphoid tissue chemokine
Stromal cell-derived factor-1 β
Platelet factor 4
GM-CSF
TNF-alpha

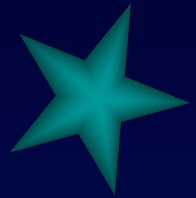
Others

β -Amyloid peptide
Cardiotoxins from spitting cobra venom
 α -Defensin
EGF-TM7 receptors CD97 and EMR2
Extracellular superoxide dismutase
Lipoprotein lipase
Thyroglobulin
Tissue plasminogen activator
Plasminogen activator inhibitor
Follistatin
Angiostatin
Endostatin
ApoB, ApoE

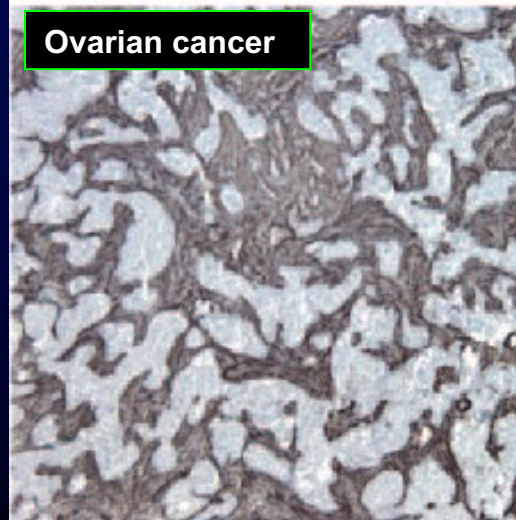
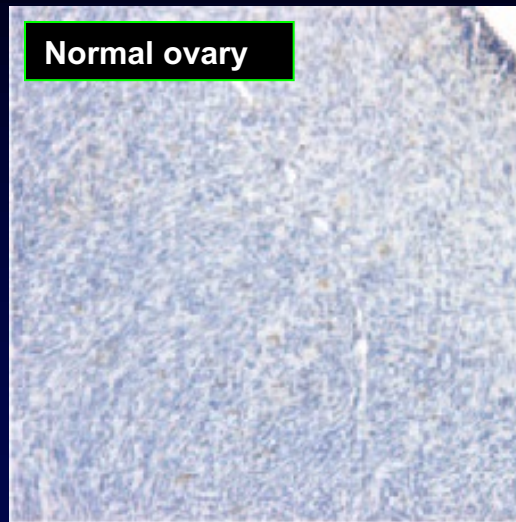
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Diseases Related to Proteoglycans and Glycosaminoglycans

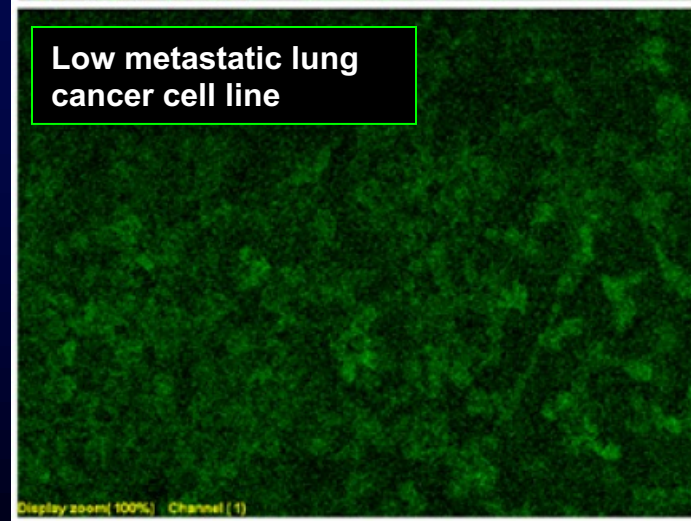
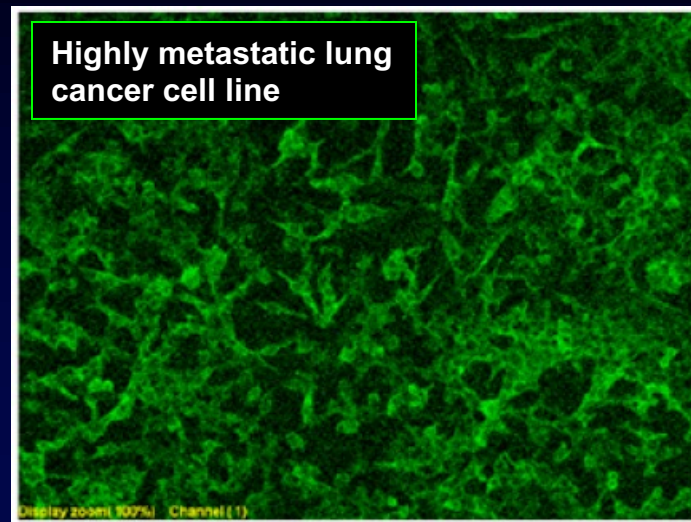
Cancer and Glycosaminoglycans



Immunostaining with an antibody against highly sulfated chondroitin sulfate

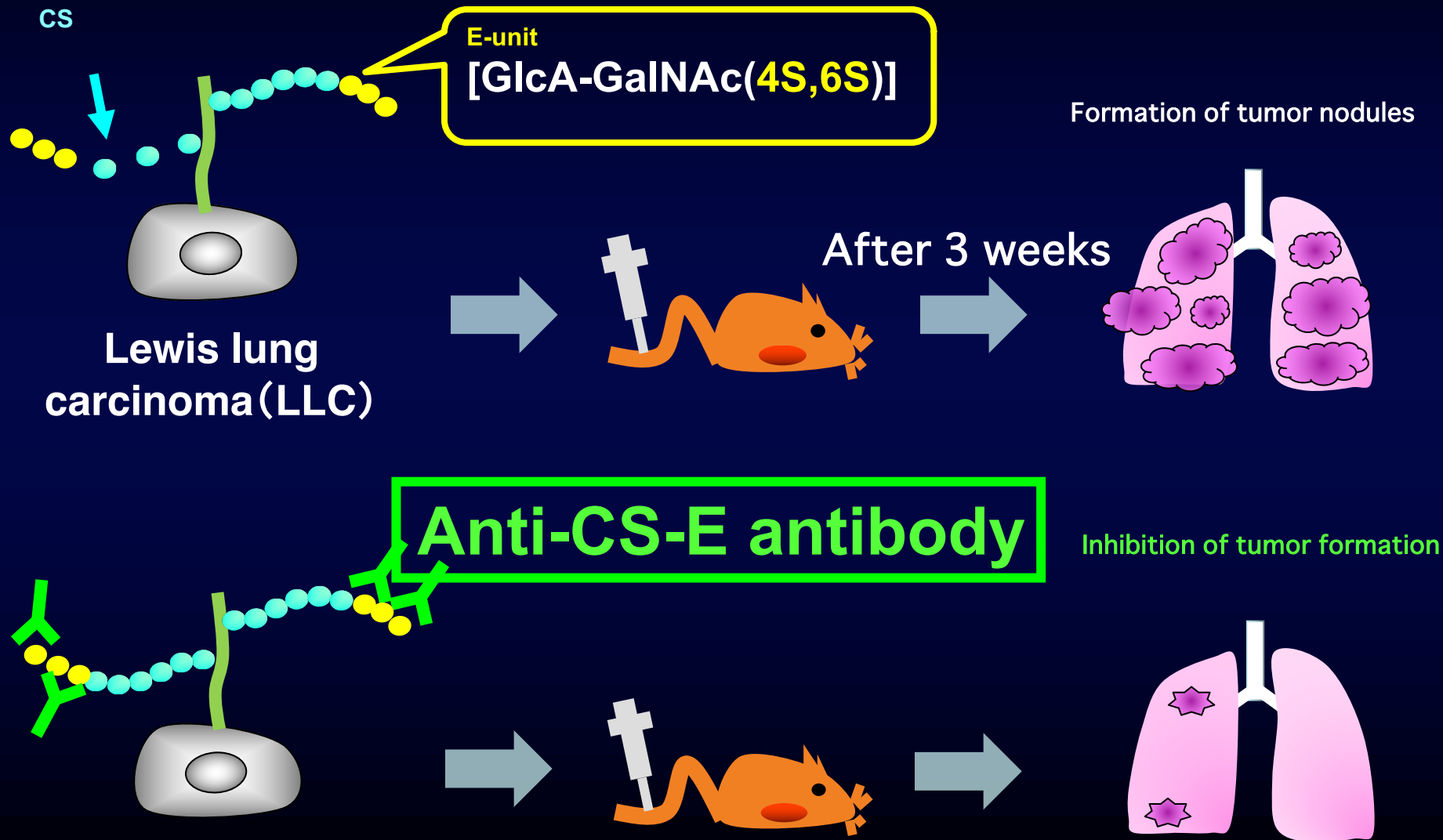


ten Dam *et al.*, *Am. J. Pathol.* (2007).



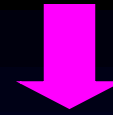
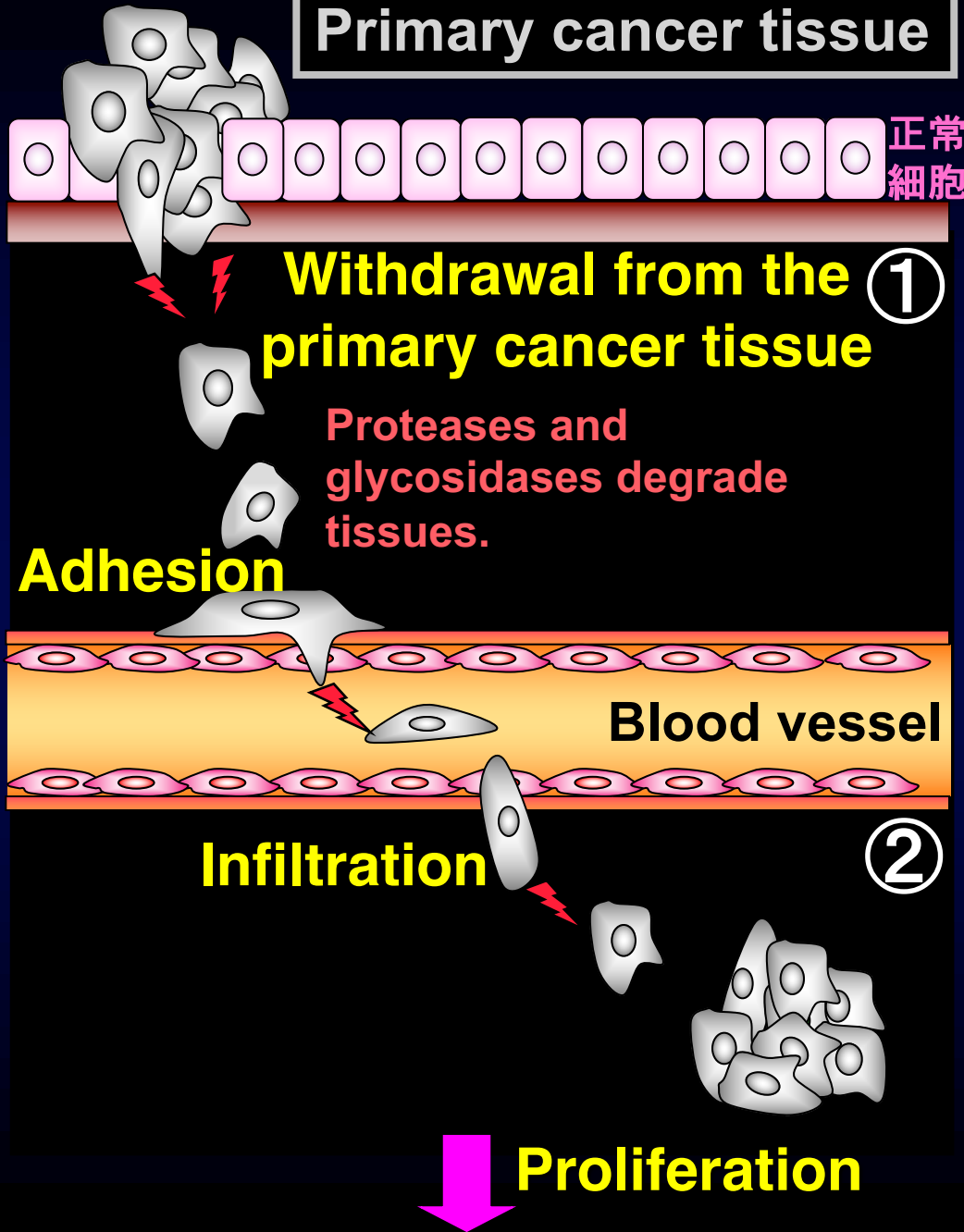
Strongly expressed in the highly metastatic cell line.
Li, *et al.*, *J. Biol. Chem.* (2008).

Inhibition of tumor formation by an antibody against highly sulfated chondroitin sulfate



Cancer cells

Primary cancer tissue

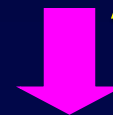


Blood vessel

Involvement of various growth factors

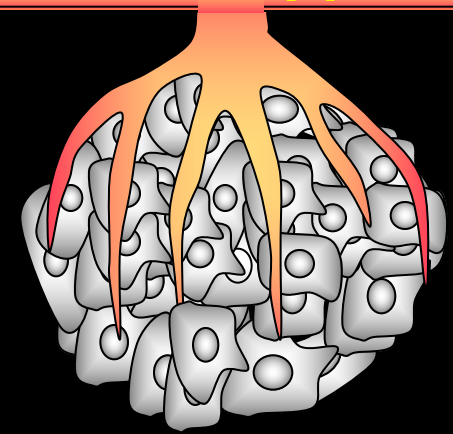
③

Angiogenesis (nutritional support)



Blood vessel

④



Cancer growth

MMPs: Matrix metalloproteinases

MMP1: Collagenase-1

MMP2: Gelatinase A

MMP3: Stromelysin-1

MMP7: Matrilysin-1

MMP8: Collagenase-2

MMP9: Gelatinase B

MMP10: Stromelysin-2

MMP11: Stromelysin-3

**MMP12: Macrophage
metalloelastase**

MMP13: Collagenase-3

MMP14: MT1-MMP

MMP15: MT2-MMP

MMP16: MT3-MMP

MMP17: MT4-MMP

MMP18: Collagenase-4

MMP19: Stromelysin-4

MMP20: Enamelysin

MMP21: X-MMP

MMP23: CA-MMP

MMP24: MT5-MMP

MMP25: MT6-MMP

MMP26: Matrilysin-2

MMP27: C-MMP

MMP28: Epilysin

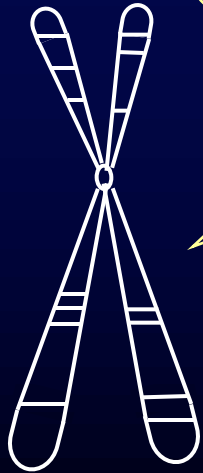
Human heparanases

4q22

HEPARANASE-1

10q23-24

HEPARANASE-2



chromosome

Spinal Cord Injury

- **Number of annual occurrences in Japan**
 - More than 5,000 (40/1,000,000)
- **Number of patients : More than 100,000**
- **Causes**
 - Traffic accidents
 - Sports injuries

In the adult centers the nerve paths are something fixed, ended and immutable. Everything must die, nothing may be regenerated. — **Cajal, 1928**

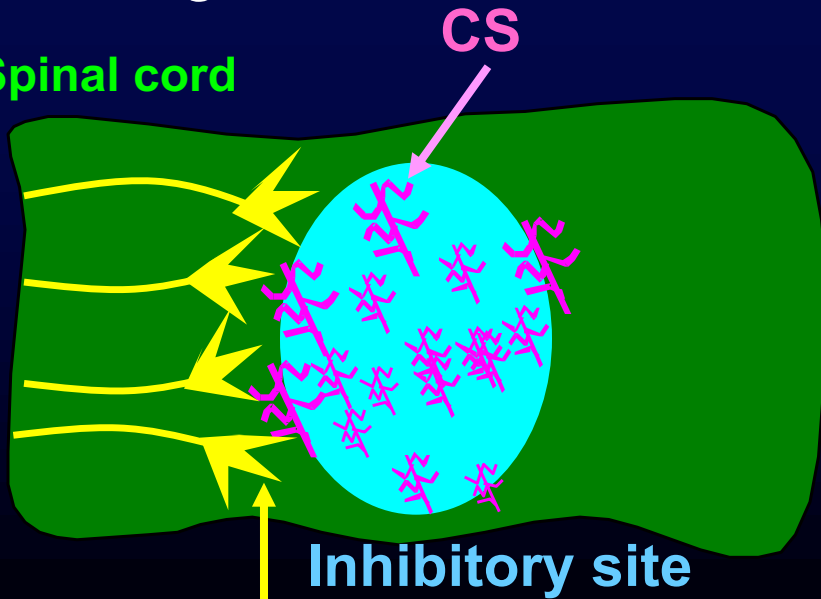
Removal of chondroitin sulfate (CS) promotes functional recovery after spinal cord injury

Bradbury et al., *Nature* 416, 636-640 (2002)

Alilain et al., *Nature* 475, 196-200 (2011)

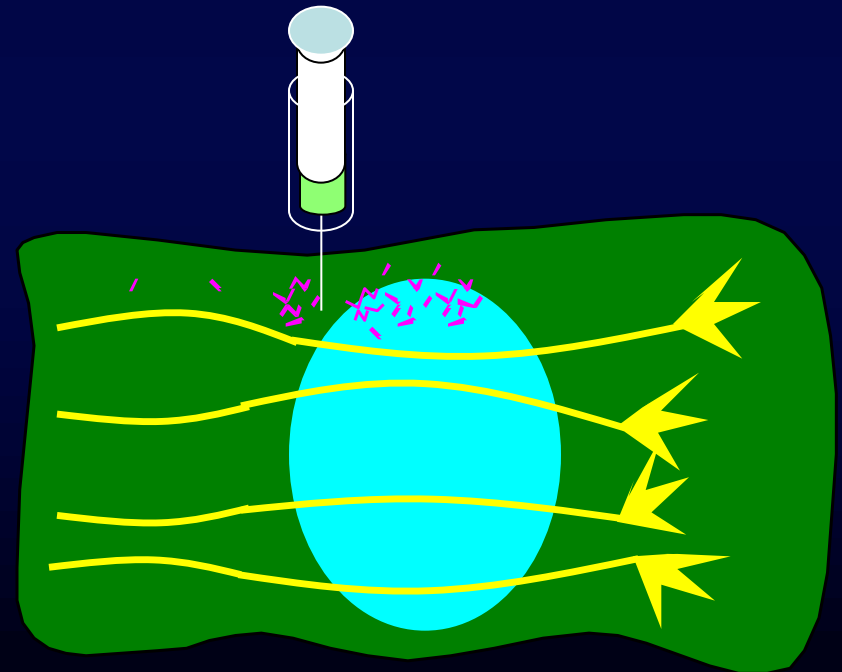
CS in a glial scar is inhibitory to axon growth.

Spinal cord



Regenerating Axons

Removal of CS by a bacterial degrading enzyme.



Infections of Pathogens via Glycosaminoglycans

Herpes Simplex Virus (HSV)

Human Immunodeficiency Virus (HIV)

Hepatitis C Virus

Dengue Virus

Japanese Encephalitis Virus

Severe Acute Respiratory Syndrome

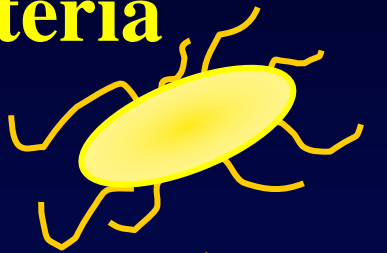
Coronavirus 2 (SARS-Cov-2)

Malaria Parasite

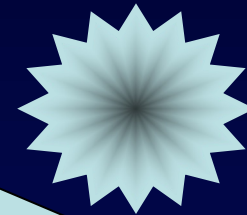
And more...

Glycoconjugates and infection by bacteria and viruses

Bacteria



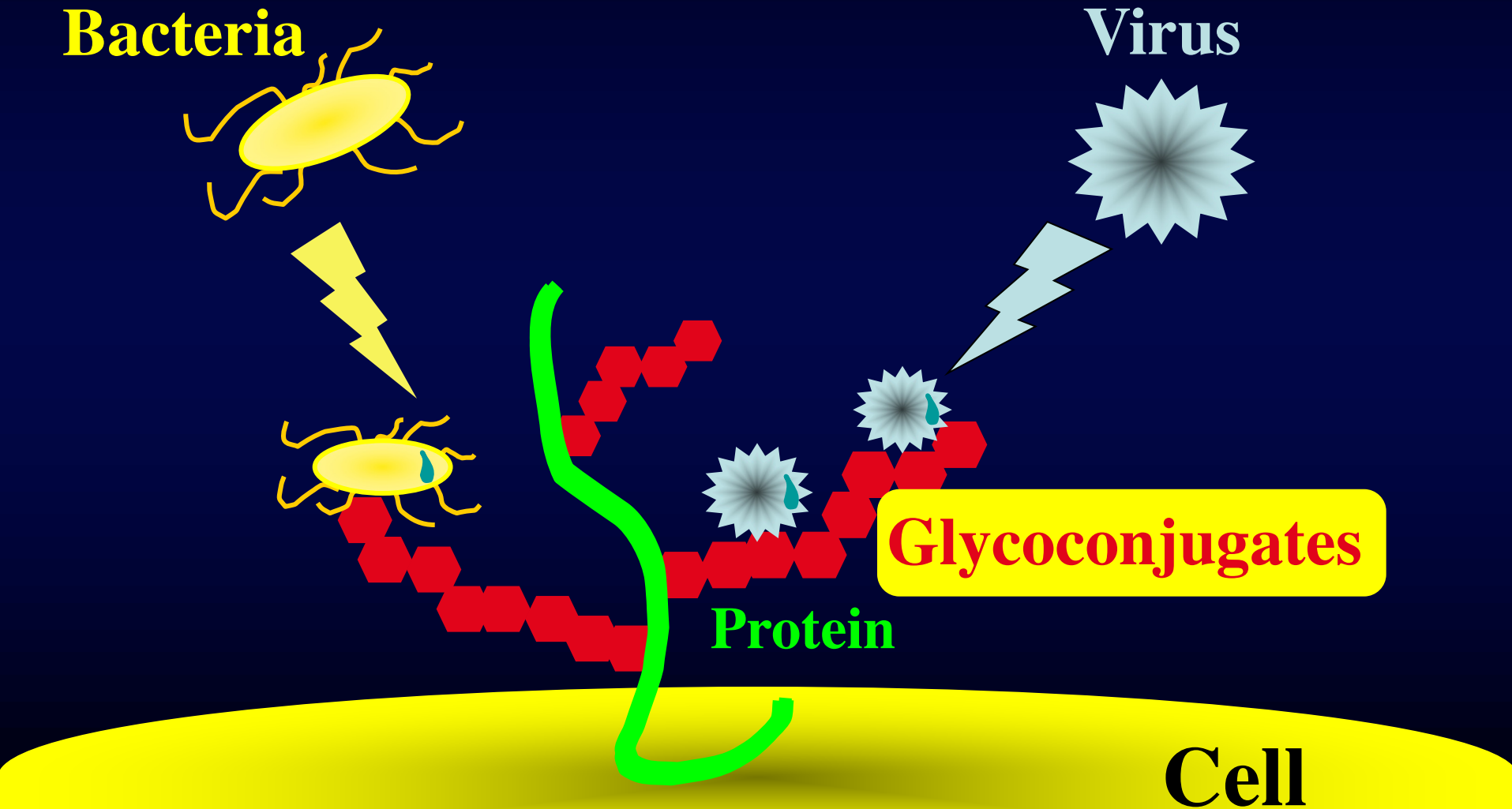
Virus



Glycoconjugates

Protein

Cell



Lysosome Diseases

Lysosome diseases, lysosomal storage diseases are a group of rare inherited metabolic disorders that result from defects in enzymes required for the metabolism in lysosome, leading to accumulation of the large molecules within the cell.

Various types of lysosome Diseases

Pompe disease

GM1 gangliosidosis

GM2 gangliosidosis

**(Tay–Sachs disease,
Sandhoff disease)**

Fabry disease

Farber disease

Gaucher disease

Niemann–Pick disease

Krabbe disease

Mucopolysaccharidoses

Multiple sulfatase deficiency

Sialidosis

Galactosialidosis

I-cell disease

Alpha-mannosidosis

Beta-mannosidosis

Fucosidosis

Aspartylglucosaminuria

Schindler disease/Kanzaki disease

Wolman disease

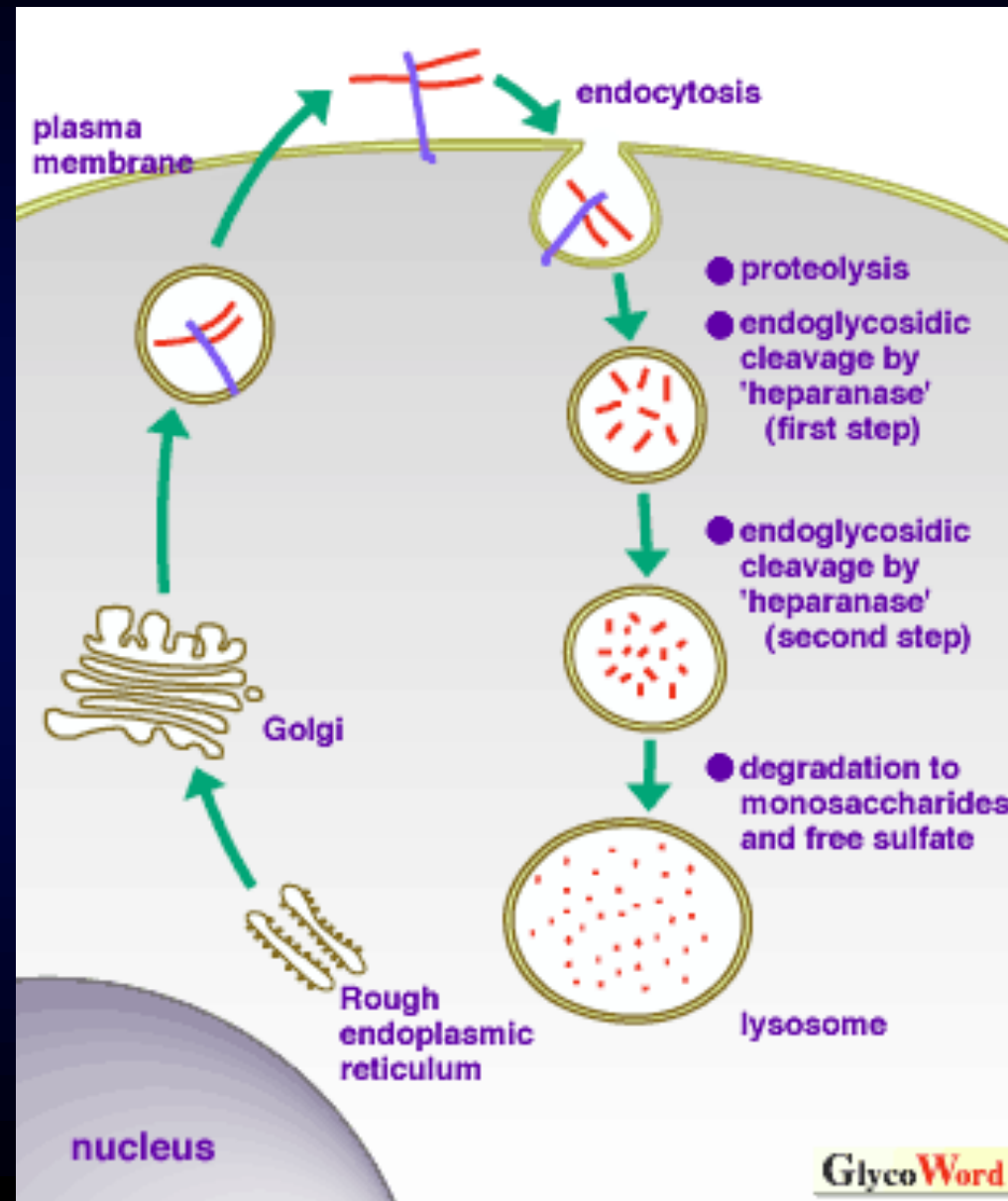
Danon disease

Sialic acid storage disease

Neuronal ceroid lipofuscinoses

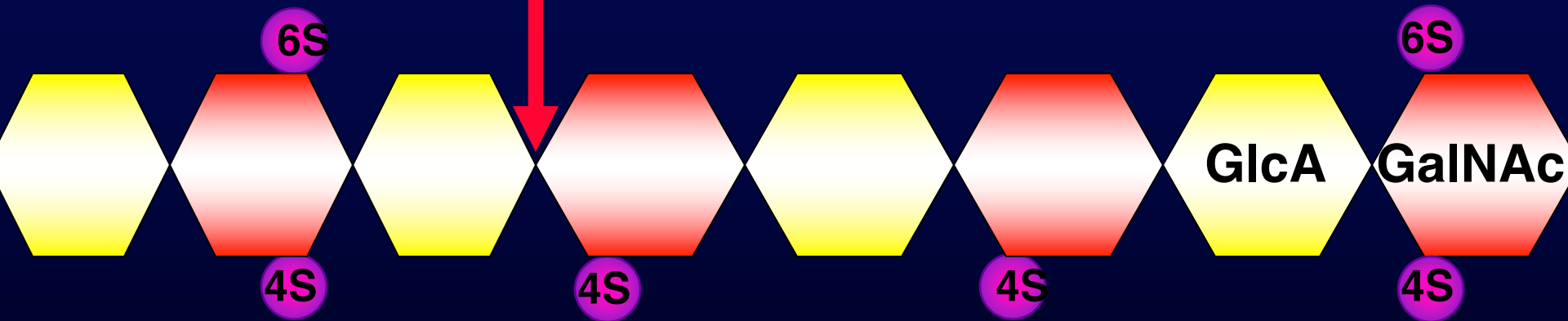
etc

Cellular catabolism of glycoproteins

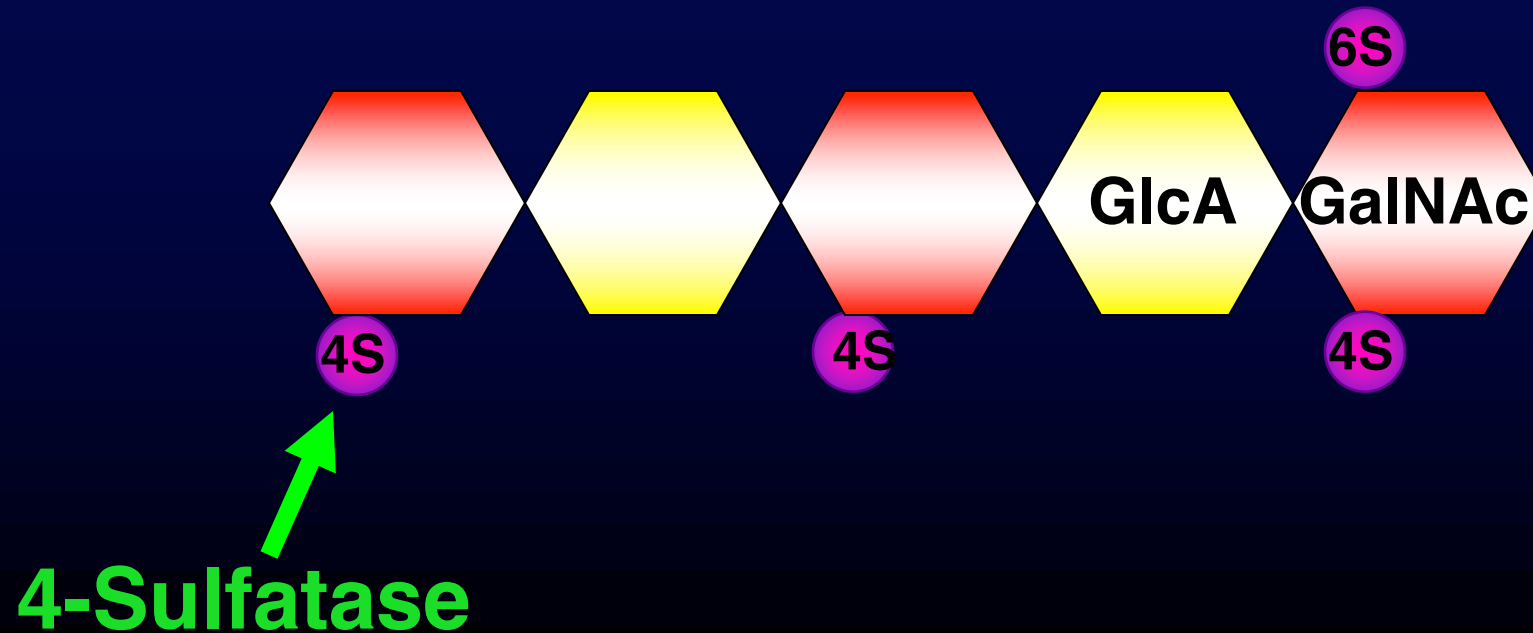


DEGRADATION OF CS/DS

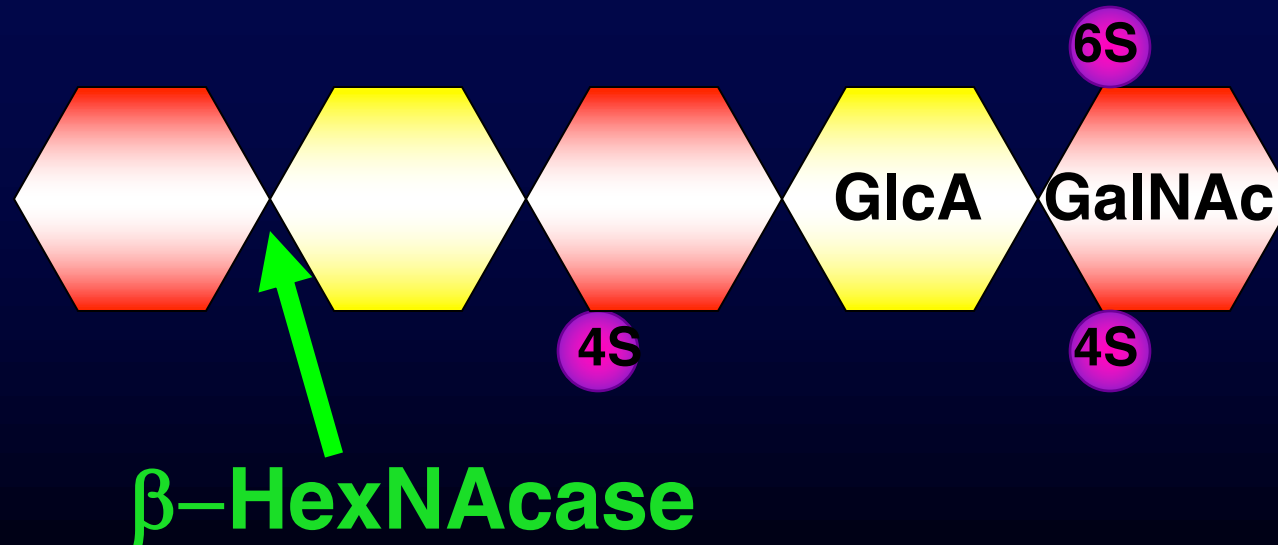
**ENDO-TYPE
ENZYME**



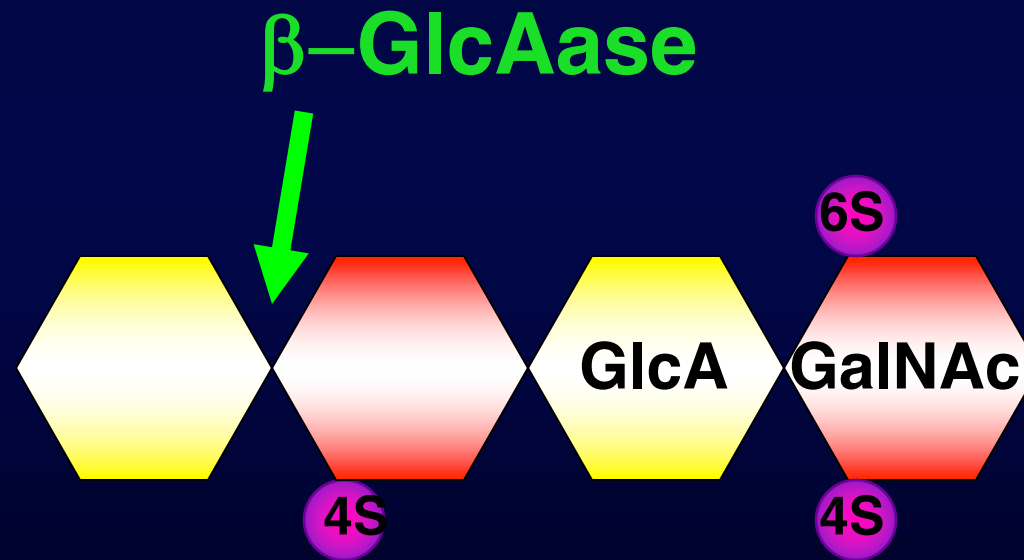
DEGRADATION OF CS/DS



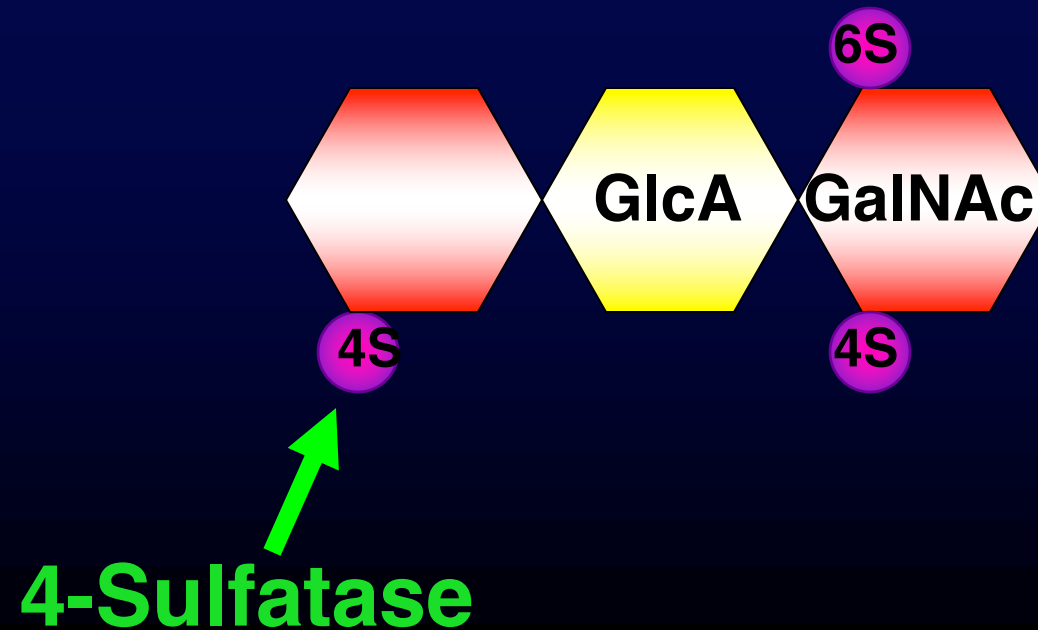
DEGRADATION OF CS/DS



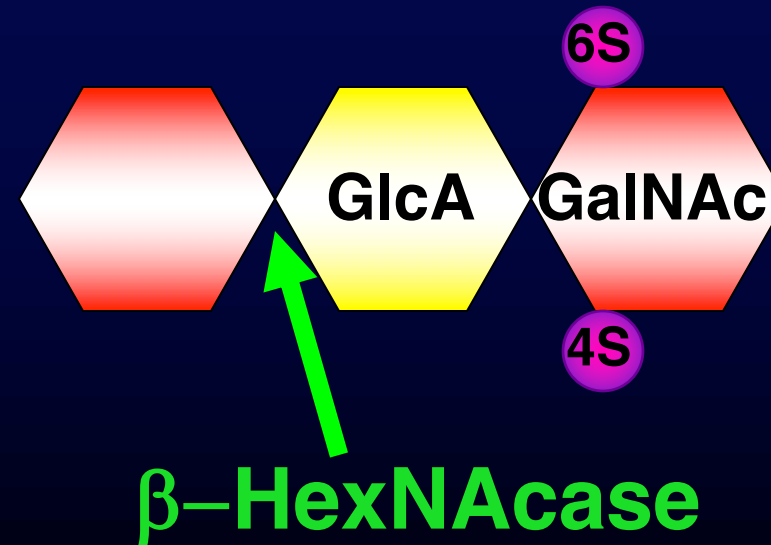
DEGRADATION OF CS/DS



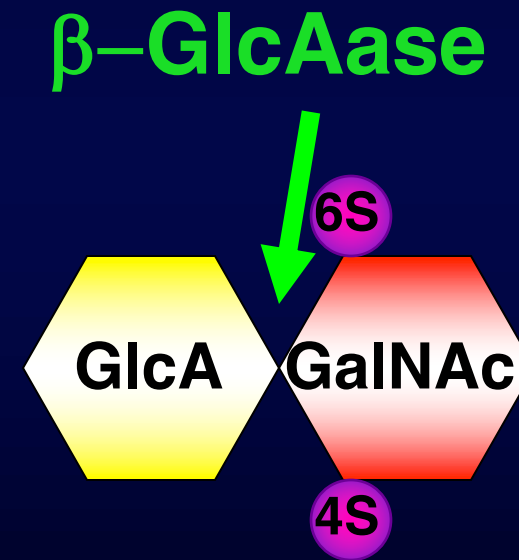
DEGRADATION OF CS/DS



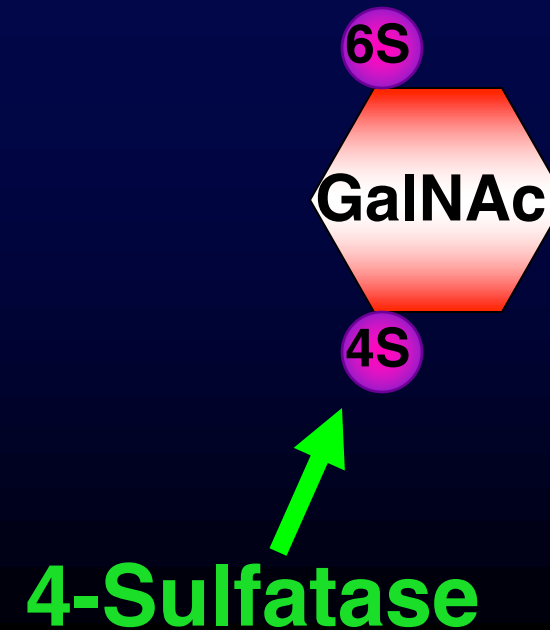
DEGRADATION OF CS/DS



DEGRADATION OF CS/DS



DEGRADATION OF CS/DS



DEGRADATION OF CS/DS

6-Sulfatase



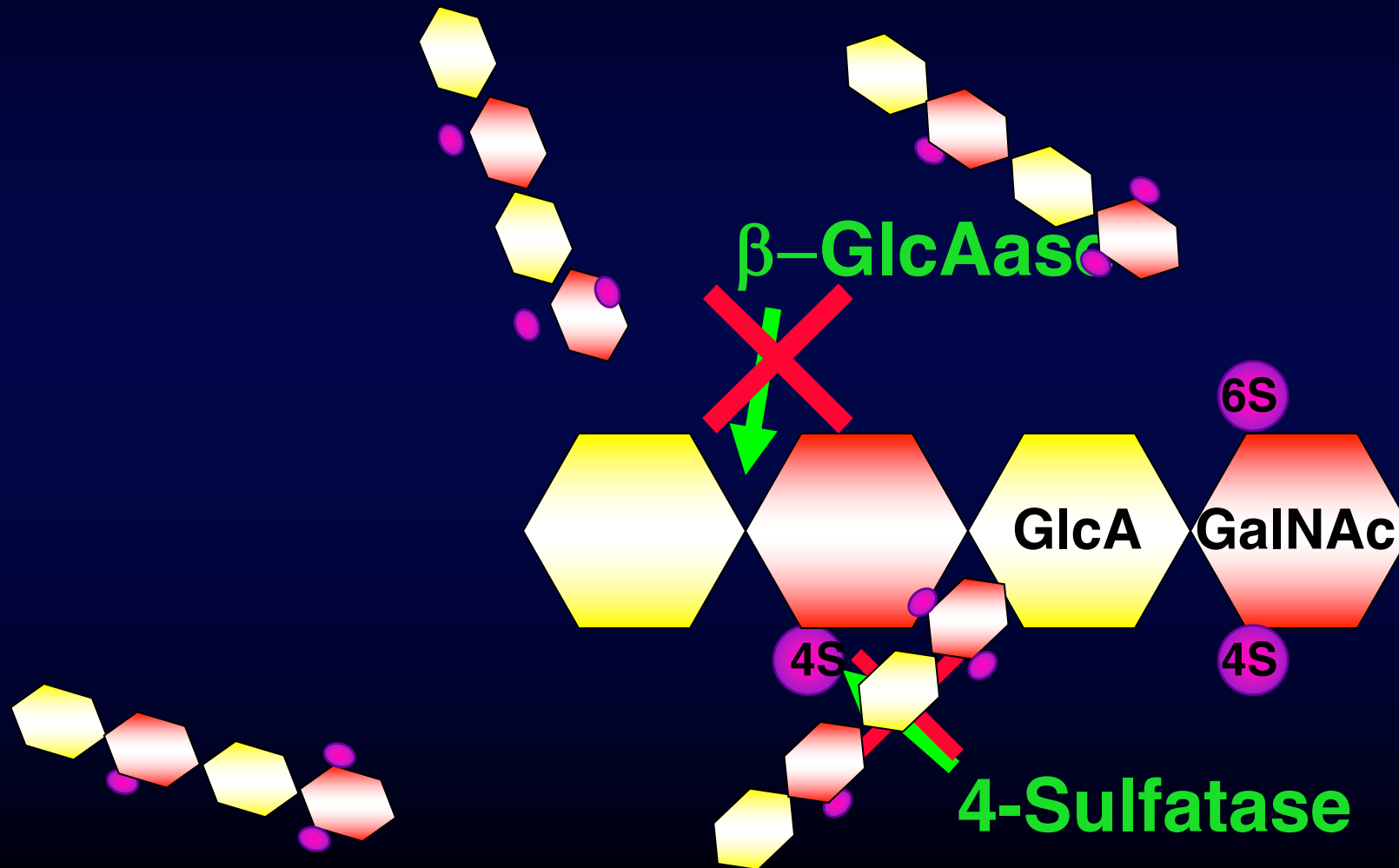
6S



DEGRADATION OF CS/DS



DEGRADATION OF CS/DS



ムコ多糖症(MPS)

Type	Eponym	Enzyme deficiency	Stored substrates
MPS I	Hurler	α -Iduronidase	DS, HS
	Scheie		
MPS II	Hunter	Iduronate 2-sulfatase	DS, HS
MPS III	Sanfilippo	Heparan sulfate degrading enzymes	HS
MPS IV	Morquio	Galactose-6-sulfatase	CS, DS, KS
MPS VI	Maroteaux-Lamy	<i>N</i> -Acetyl- <i>D</i> -galactosamine-4-sulphatase	DS
MPS VII	Sly	β - <i>D</i> -Glucuronidase	DS, HS

※ HS :Heparan sulfate

Type	Eponym	Enzyme Deficiency	Stored GAG	Pathological Conditions
MPS I	Hurler	α -L-iduronidase	HS, DS	Mental retardation, skeletal deformities, cloudy corneas, hepatosplenomegaly, joint contracture, hearing loss, short stature
	Scheie			Joint contracture, cloudy corneas, aortic closure
MPS II	Hunter (X-linked recessive inheritance)	iduronate-2-sulfatase	HS, DS	(Severe type) Mental retardation, hepatosplenomegaly (Mild type) Joint contracture
MPS III	Sanfilippo (type A) (type B) (type C) (type D)	HS- <i>N</i> -sulfatase α - <i>N</i> -acetylglucosaminidase acetyl-CoA: α -glucosaminide <i>N</i> -acetyltransferase α - <i>N</i> -acetylglucosamine-6-sulfatase	HS	Mild skeletal changes, mental retardation
MPS IV	Morquio (type A) (type B)	galactose-6-sulfatase (<i>N</i> -acetylgalactosamine-6-sulfatase) β -galactosidase	KS, C6S KS	Skeletal deformities (anterior thoracic protrusion, vertebral posterolateral), cloudy corneas, aortic regurgitation
MPS VI	Maroteaux - Lamy	<i>N</i> -acetylgalactosamine 4-sulfatase	DS	Skeletal deformities, cloudy corneas, hepatosplenomegaly, heart disorder
MPS VII	Sly	β -D-glucuronidase	HS, DS	Skeletal deformities, cloudy corneas, mental retardation

Type	Eponym	Enzyme Deficiency	Stored GAG	Pathological Conditions
MPS I	Hurler	α -L-iduronidase	HS, DS	Mental retardation, skeletal deformities, cloudy corneas, hepatosplenomegaly, joint contracture, hearing loss, short stature
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MPS III	Sanfilippo (type A) (type B) (type C) (type D)	HS- <i>N</i> -sulfatase α - <i>N</i> -acetylglucosaminidase acetyl-CoA: α -glucosaminide <i>N</i> -acetyltransferase α - <i>N</i> -acetylglucosamine-6-sulfatase	HS	Mild skeletal changes, mental retardation
MPS IV	Morquio (type A) (type B)	galactose-6-sulfatase (<i>N</i> -acetylgalactosamine-6-sulfatase) β -galactosidase	KS, C6S KS	Skeletal deformities (anterior thoracic protrusion, vertebral posterolateral), cloudy corneas, aortic regurgitation
MPS VI	Maroteaux - Lamy	<i>N</i> -acetylgalactosamine 4-sulfatase	DS	Skeletal deformities, cloudy corneas, hepatosplenomegaly, heart disorder
MPS VII	Sly	β -D-glucuronidase	HS, DS	Skeletal deformities, cloudy corneas, mental retardation

MPS IX

Deficiency: Hyaluronidase 1

Storage: HA in plasma

Symptoms: Mild phenotype
(Periarticular masses and
mild short stature)

Treatment of lysosome disease

1. Symptomatic treatment
2. Enzyme replacement therapy (ERT)
3. Bone marrow transplantation
4. Gene therapy

Enzyme Replacement Therapy, ERT

ERT is a medical treatment which replaces an enzyme that is deficient or absent in the body. Increasing the concentration of the missing enzyme within the body improves the body's normal cellular metabolic processes and reduces substrate concentration in the body.

Problems on ERT

1. Less available to certain areas in the body (brain, bone, and heart)
2. Less efficient to be incorporated into cells
3. Less half life in the body
4. Need to be treated earlier in life time
5. Unwanted immune response against the enzyme
6. Expensive costs

Congenital disorder of glycosylation (CDG)

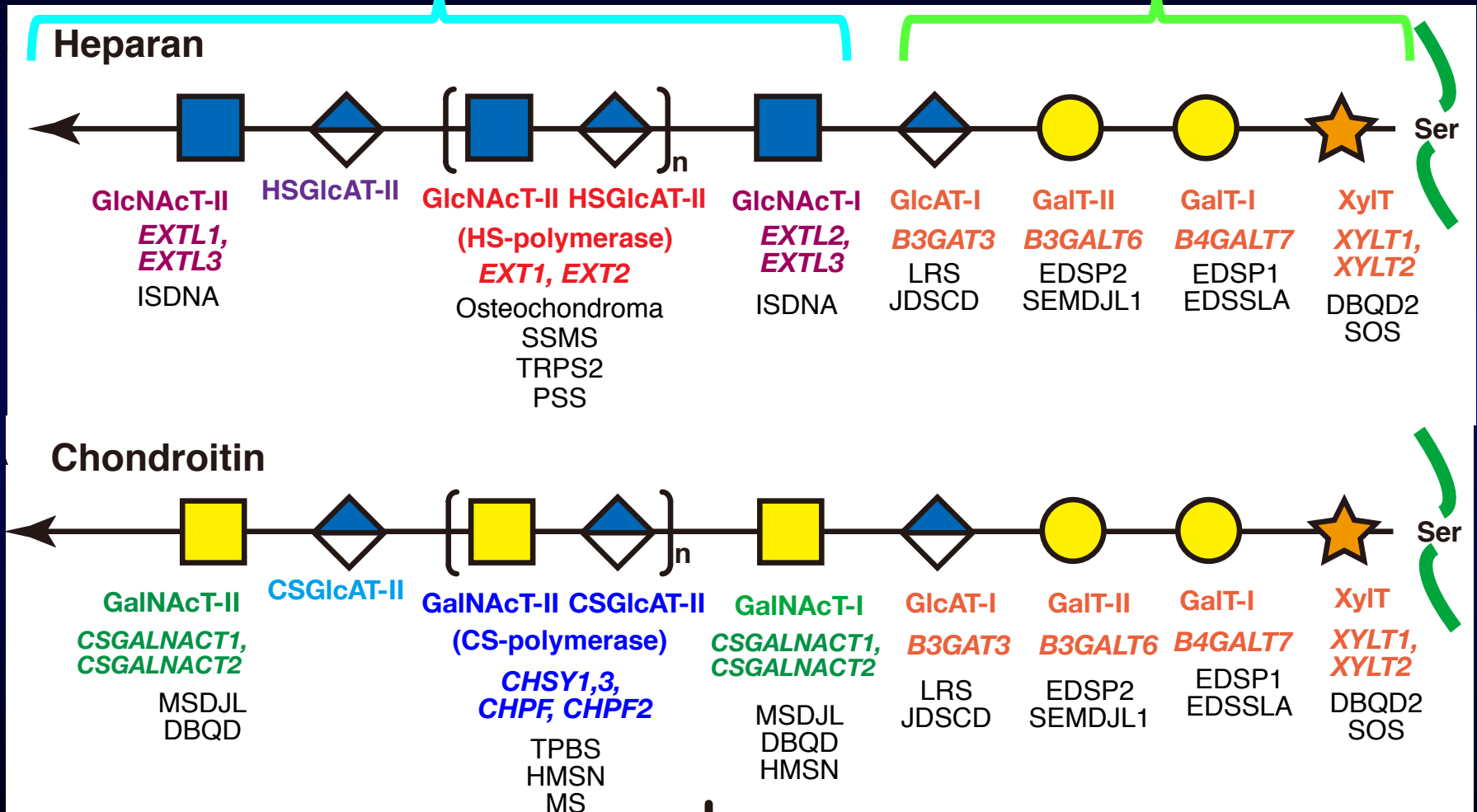
It is caused by a deficiency of glycosyltransferase, which is involved in the biosynthesis of sugar chains on glycoproteins.

Common symptoms of this disease include hypotonia in infancy, poor weight gain, severe psychomotor developmental delay, characteristic facial features, epilepsy, ophthalmologic abnormalities such as esotropia, skin symptoms such as fatty deposits in the buttocks and nipple retraction, pericardial effusion, cardiomyopathy, liver dysfunction, and abnormalities in blood clotting factors.

Biosynthesis of HS and CS backbones

Repeating disaccharide region

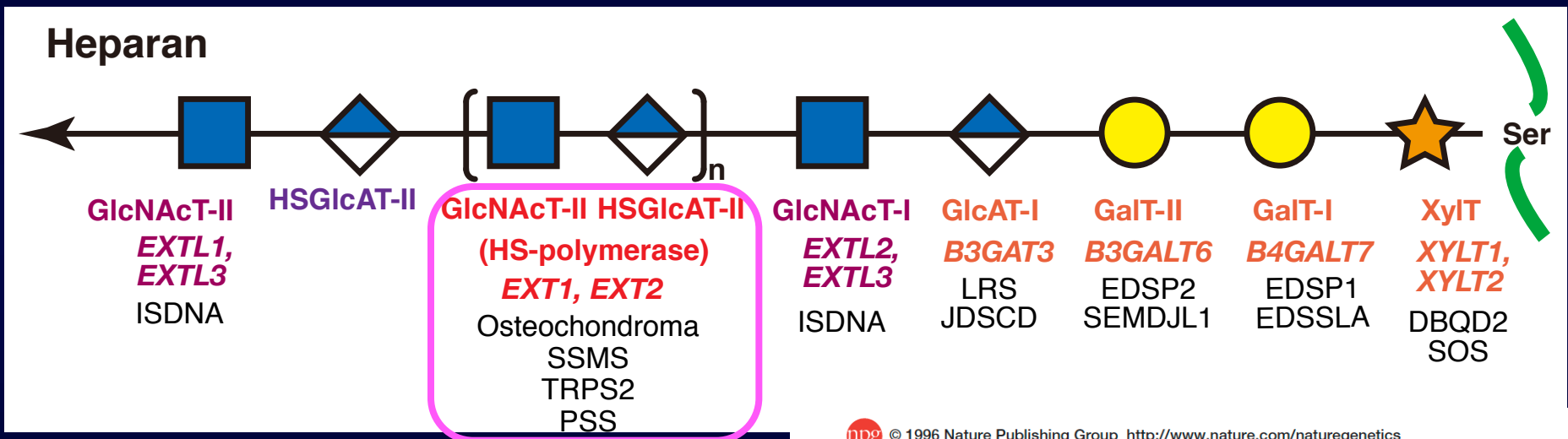
Linker tetra-saccharide region



Human genetic disorders caused by mutations in *EXT1* or *EXT2*

Repeating disaccharide region of HS

Linker tetra-saccharide region



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article

Cloning of the putative tumour suppressor gene for hereditary multiple exostoses (*EXT1*)

Jung Ahn^{*1}, Hermann-Josef Lüdecke^{*2}, Steffi Lindow², William A. Horton³, Brendan Lee³, Michael J. Wagner¹, Bernhard Horsthemke² & Dan E. Wells¹

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article

The *EXT2* multiple exostoses gene defines a family of putative tumour suppressor genes

Dominique Stickens^{1*}, Gregory Clines^{1*}, David Burbee^{1,3}, Purita Ramos¹, Sylvia Thomas¹, Deborah Hogue⁴, Jacqueline T. Hecht⁴, Michael Lovett^{1,3} & Glen A. Evans^{1,2,3}

Hereditary multiple exostosis is caused by mutations in HS biosynthetic enzymes EXT1/EXT2

Hereditary multiple exostosis is an autosomal dominant skeletal disorder characterized by the presence of cartilage capped bony outgrowths mainly located at the juxtaepiphyseal region of the long bones.

It is caused by mutations in EXT1 and EXT2.

The heterooligomeric complex of EXT1 and EXT2 represents the biologically relevant form of the HS polymer modification

Proteoglycan Linkeropathy

•Proteoglycan Linkeropathy is a collective term for diverse connective tissue disorders caused by mutations in the glycosyltransferases responsible for the biosynthesis of the linker region tetrasaccharide, which is characterized by bone, skin, connective tissue, and heart defects.

Genetic disorders

Affected genes

Desbuquois dysplasia type 2

XYLT1

Spondyloocular syndrome

XYLT2

Spondylodysplastic

B4GALT7,

Ehlers-Danlos syndrome

B3GALT6

Larsen-like syndrome

GlcAT-I

Human genetic disorders caused by mutations in enzymes responsible for biosynthesis of CS, DS, and HS

Genetic disorders	Affected genes	Refs.
Spondyloepiphyseal dysplasia, Omani type	C6ST-1	Am. J. Med. Genet. 2008
Severe osteopenia and fractures	GlcAT-I	BMC Med. Genet. 2016; J. Med. Genet. 2020
A mild skeletal dysplasia and joint laxity	GalNAcT1	Hum. Mutat. 2017; 2020
Musculocontractural Ehlers-Danlos syndrome	D4ST-1 DSE	Hum. Mutat. 2010; Clin. Biochem. 2017; BBA 2019 Mol. Genet. Genomic Med. 2020; Hum. Mutat. 2022
A novel type of spondylo-epi-metaphyseal dysplasia	EXTL3	J. Hum. Genet. 2017
Pseudodiastrophic dysplasia	CANT1	J. Med. Genet. 2020
Desbuquois dysplasia type 2 Pseudotorsion dysplasia	XYLT1 XYLT2	In preparation

① Ehlers-Danlos syndrome caused by mutations in D4ST-1

② A novel type of spondylo-epi-metaphyseal dysplasia caused by mutations in EXTL3

① Ehlers-Danlos syndrome caused by mutations in D4ST-1

② A novel type of spondylo-epi-metaphyseal dysplasia caused by mutations in EXTL3

Ehlers-Danlos syndrome

- **A heritable connective tissue disorder characterized by joint and skin laxity as well as tissue fragility.**
(aged appearance, hypermobile joints, short stature, craniofacial dysmorphism etc.)
- **Previously, the Ehlers-Danlos syndromes were classified in a system of six subtypes as follows.**

Types	Affected genes
Classic type	<i>COL5A1, COL5A2</i>
Vascular type	<i>COL3A1</i>
Arthrochalasia type	<i>COL1A1, COL1A2</i>
Dermatosparaxis type	<i>ADAMTS2</i>
Hypermobility type	<i>TenascinXB</i>
Kyphoscoliosis type	<i>Lysyl hydroxylase</i>

Clinical symptoms of Ehlers-Danlos syndrome (Kosho type)



Others :

Skin hyperextension

Wrinkled skin

Defective heart valve

Subcutaneous hematoma

Identification of the mutations in Ehlers-Danlos syndrome

P281L
Y293C

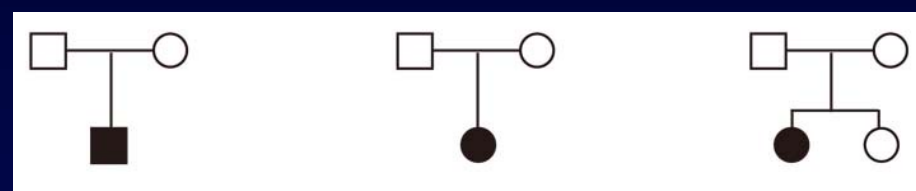
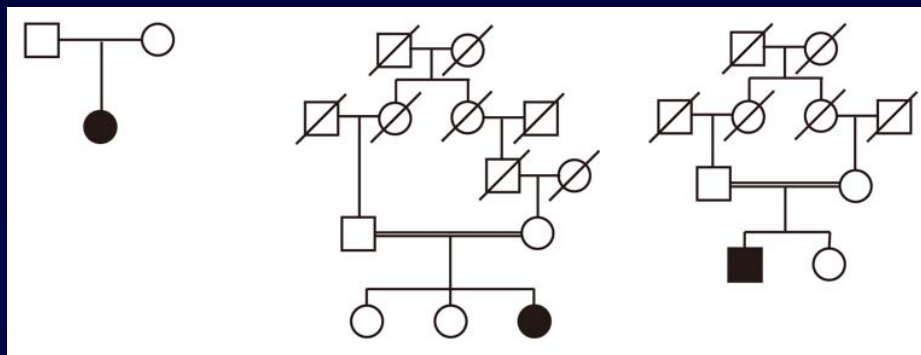
P281L
(homo)

P281L
(homo)

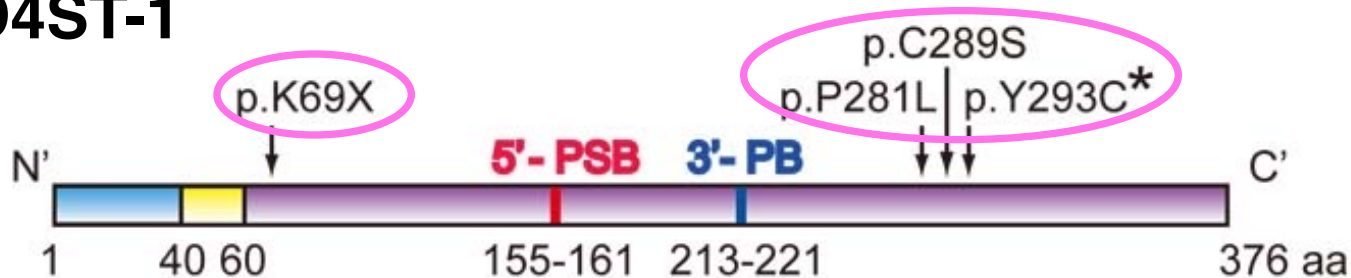
K69X
P281L

P281L
C289S

P281L
Y293C



D4ST-1



5'-PSB, 5'-phosphosulfate binding motif
3'-PB, 3'-phospho binding motif

Ehlers-Danlos syndrome

- **A heritable connective tissue disorder characterized by joint and skin laxity as well as tissue fragility.**
(aged appearance, hypermobile joints, short stature, craniofacial dysmorphism etc.)

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Kyphoscoliosis type	<i>Lysyl hydroxylase</i>

Ehlers-Danlos syndrome

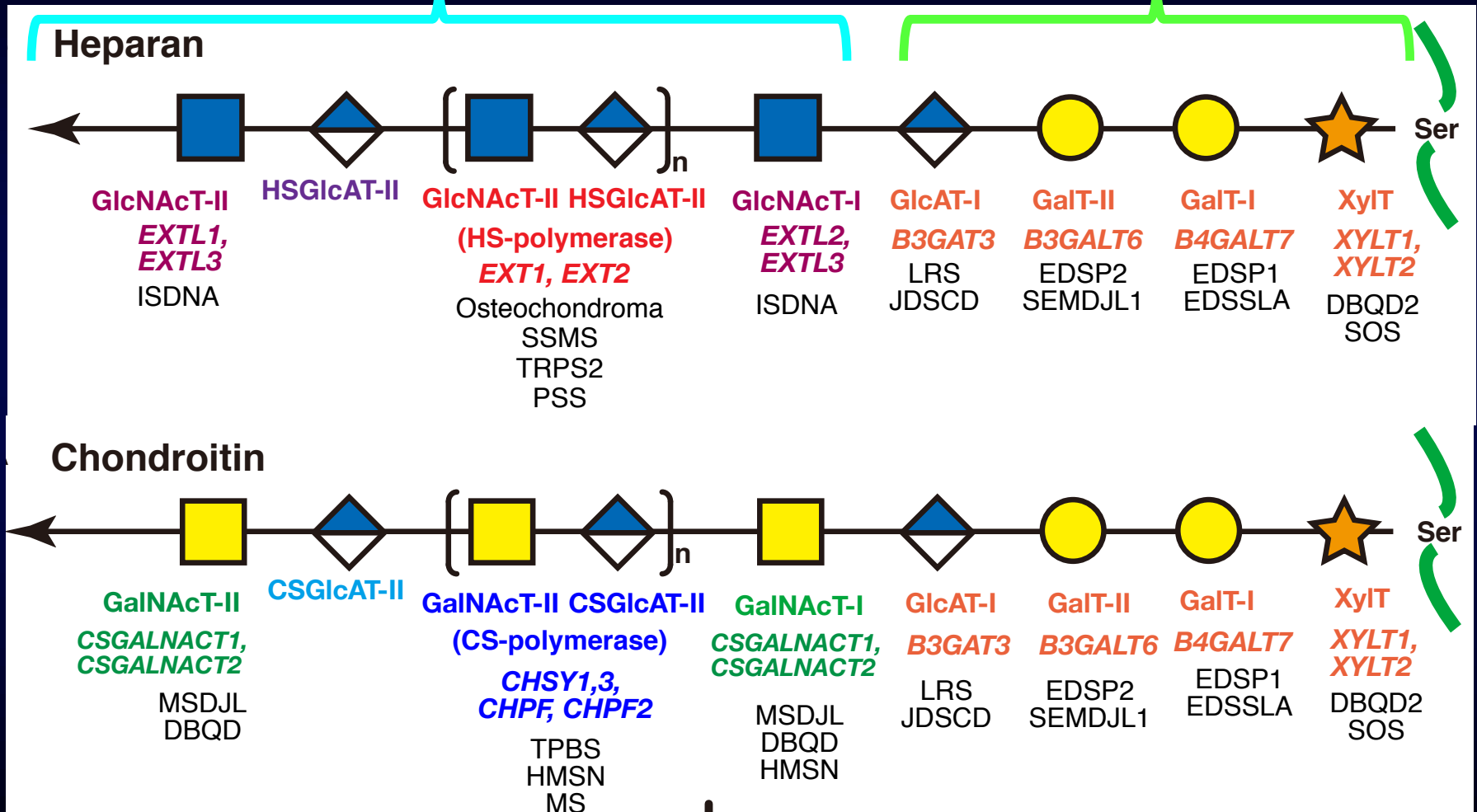
- The International EDS Consortium proposes a revised EDS classification, which recognizes 13 subtypes. (Malfait *et al.* American Journal of Medical Genetics, 2017)

New Types	Affected genes
Classical-like type	<i>TenascinXB</i>
Cardiac-valvular type	<i>COL1A2</i>
Brittle Corbea Syndrome	<i>ZNF469, PRDM5</i>
Spondylodysplastic type	<i>B4GALT7, B3GALT6, SLC39A13</i>
Musculocontractural type	<i>DSE, D4ST-1</i>
Myopathic type	<i>COL12A1</i>
Periodontal type	<i>C1R</i>

Biosynthesis of HS and CS backbones

Repeating disaccharide region

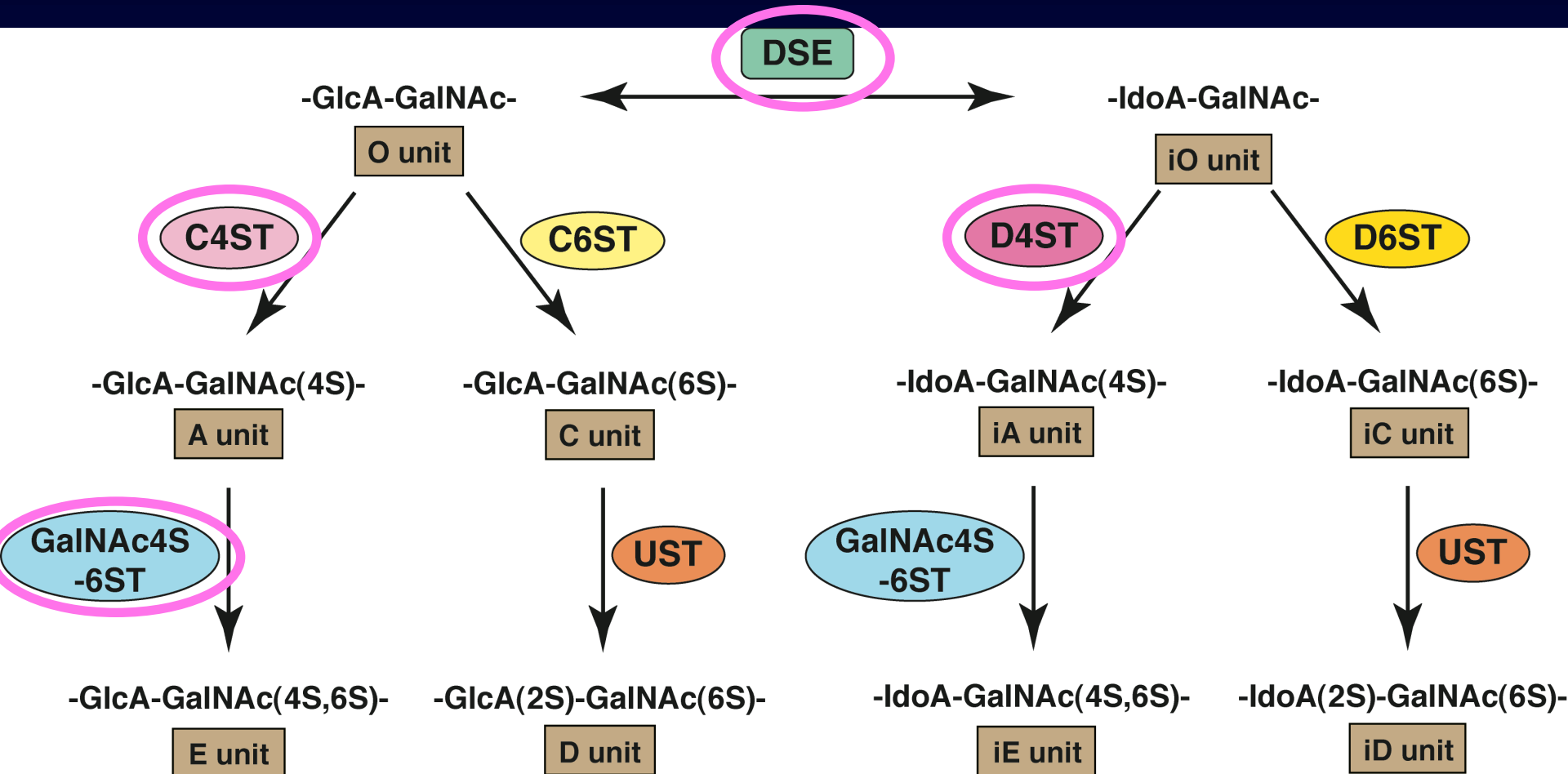
Linker tetra-saccharide region



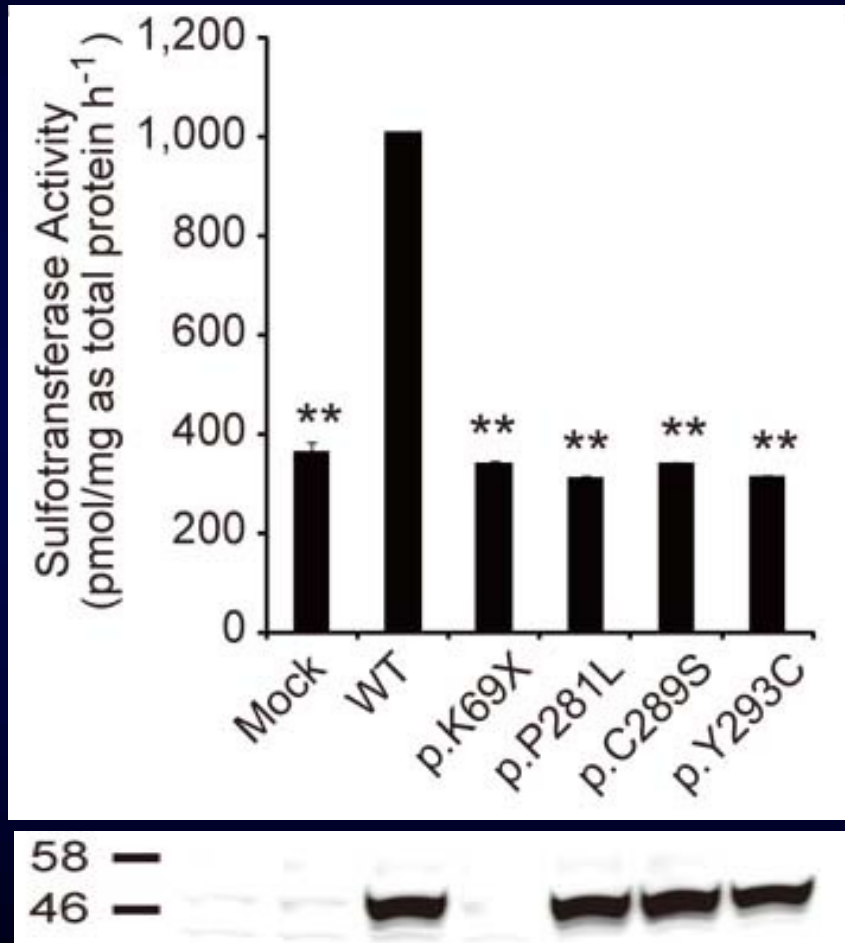
Schematic pathways of the modifications of CS and DS

CS

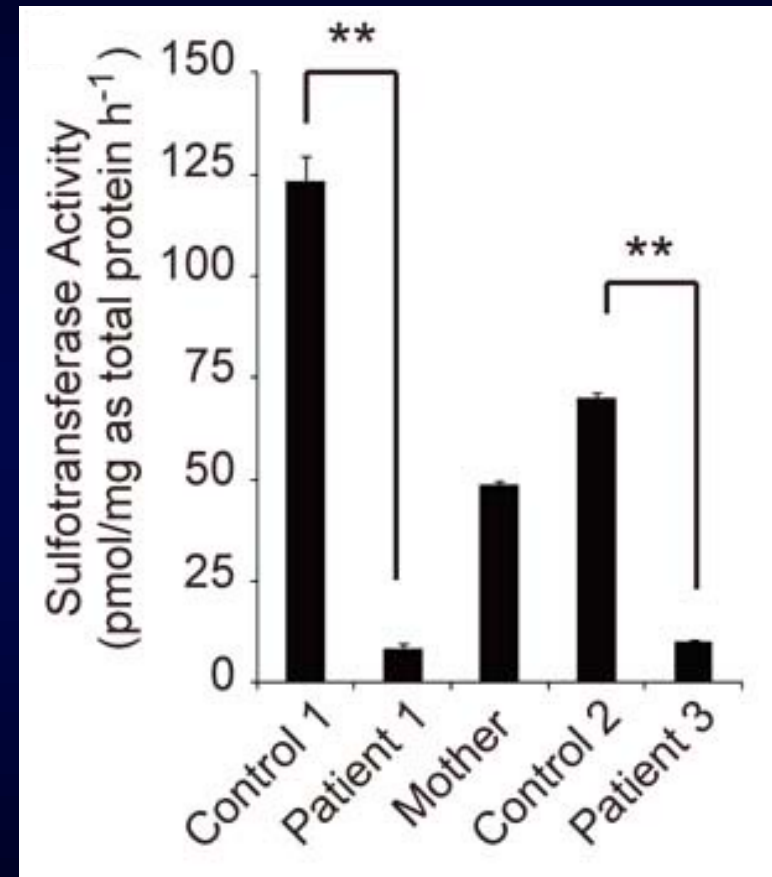
DS



Sulfotransferase activities of the recombinant D4ST-1 and the fibroblasts from patients



** $P < 0.001$



Patient 1, P281L/Y293C

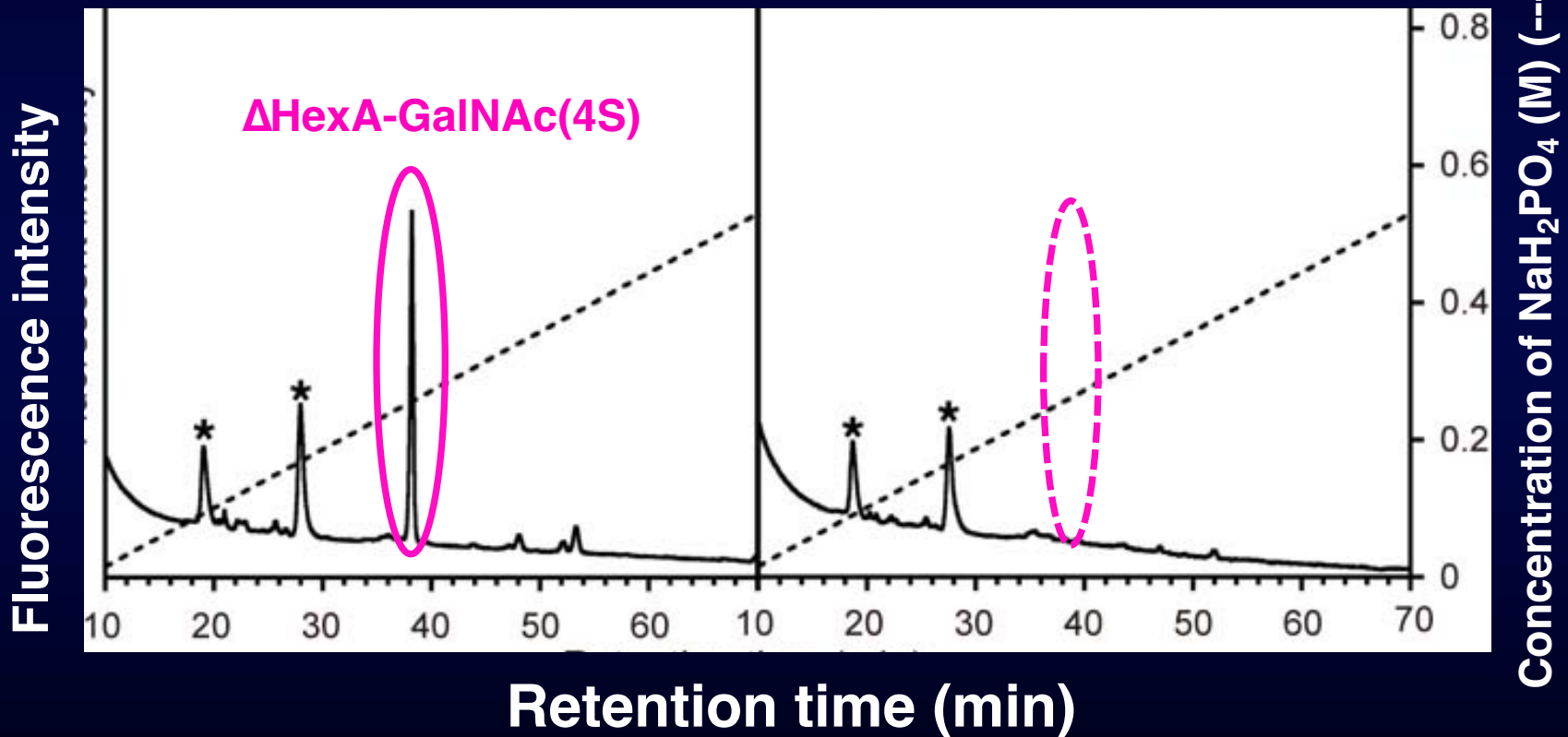
Patient 3, P281L (homo)

** $P < 0.001$

Chromatograms of the DS disaccharide from the skin fibroblasts

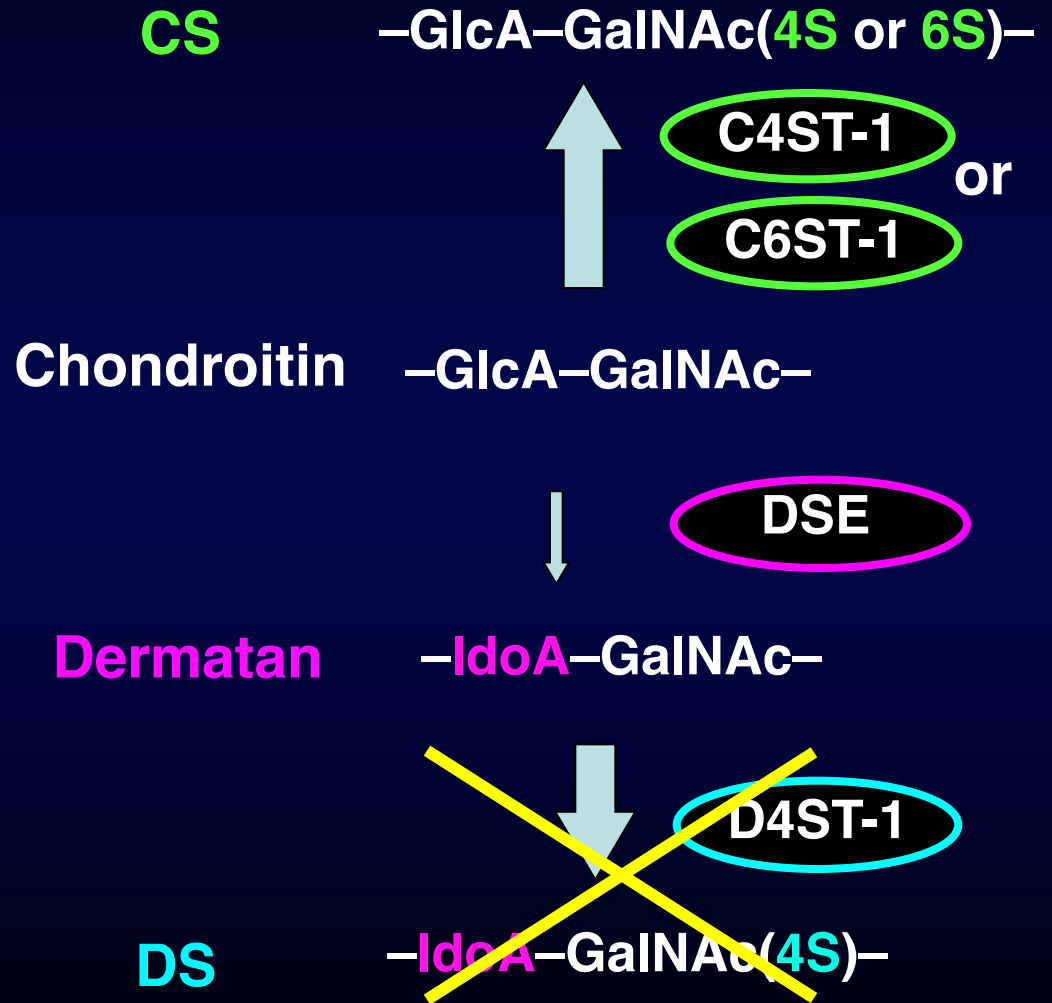
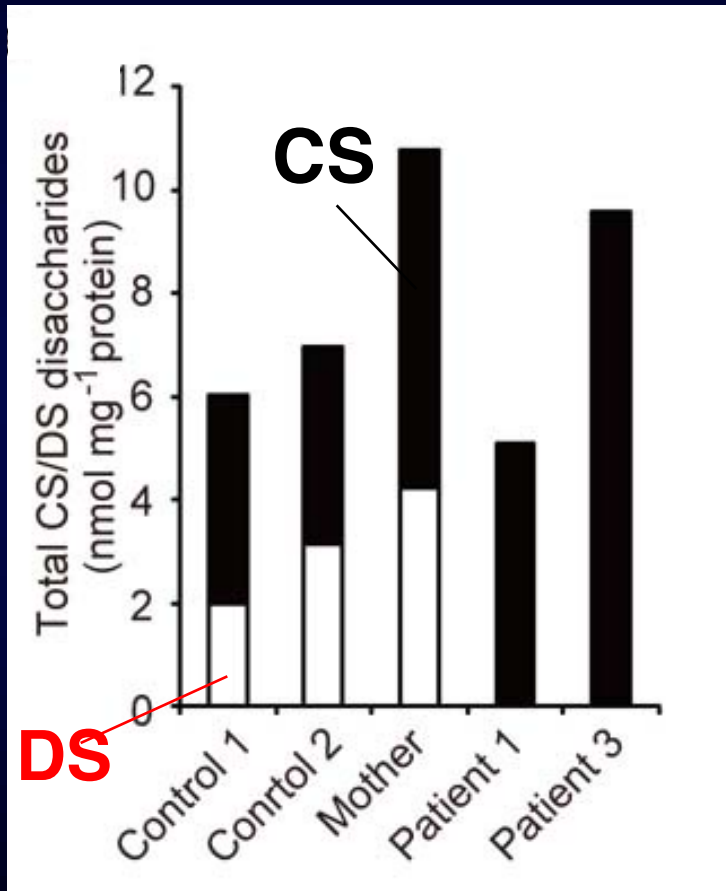
Control 2

Patient 3 (P281L)

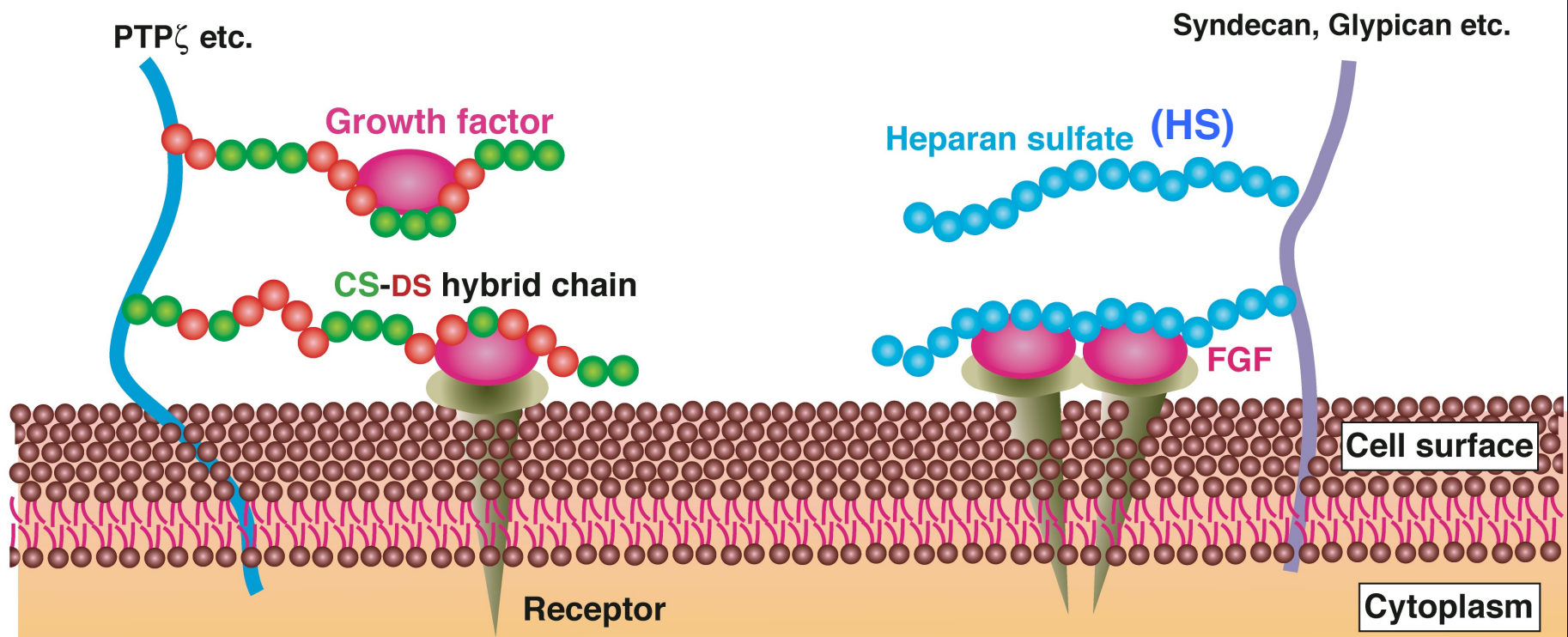
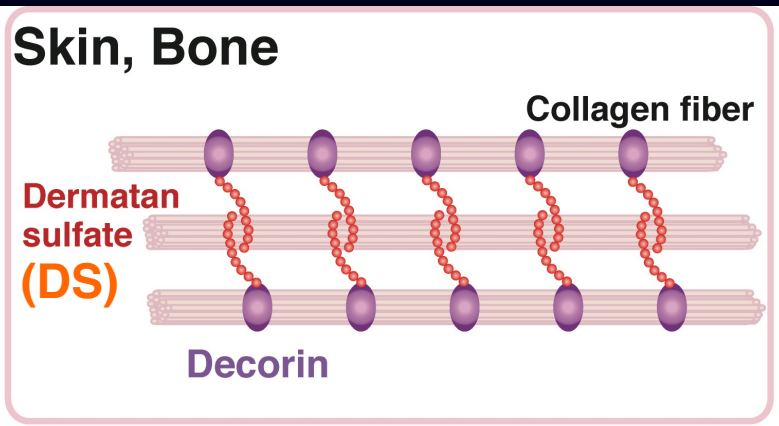
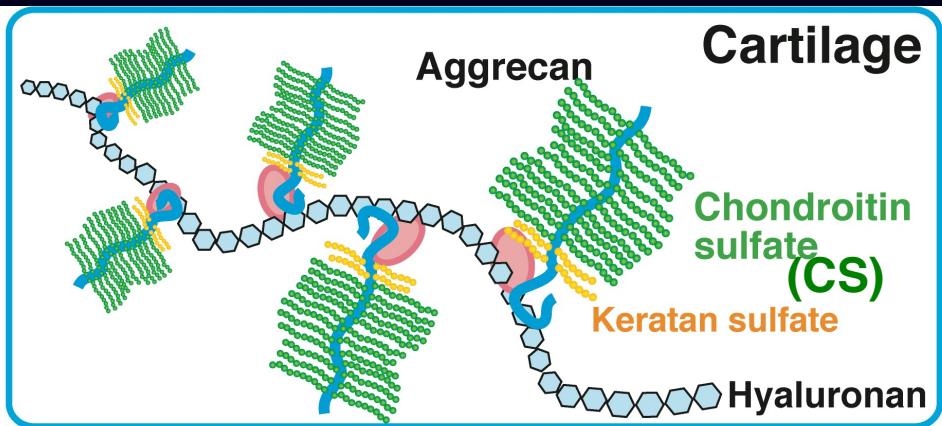


*, impurity

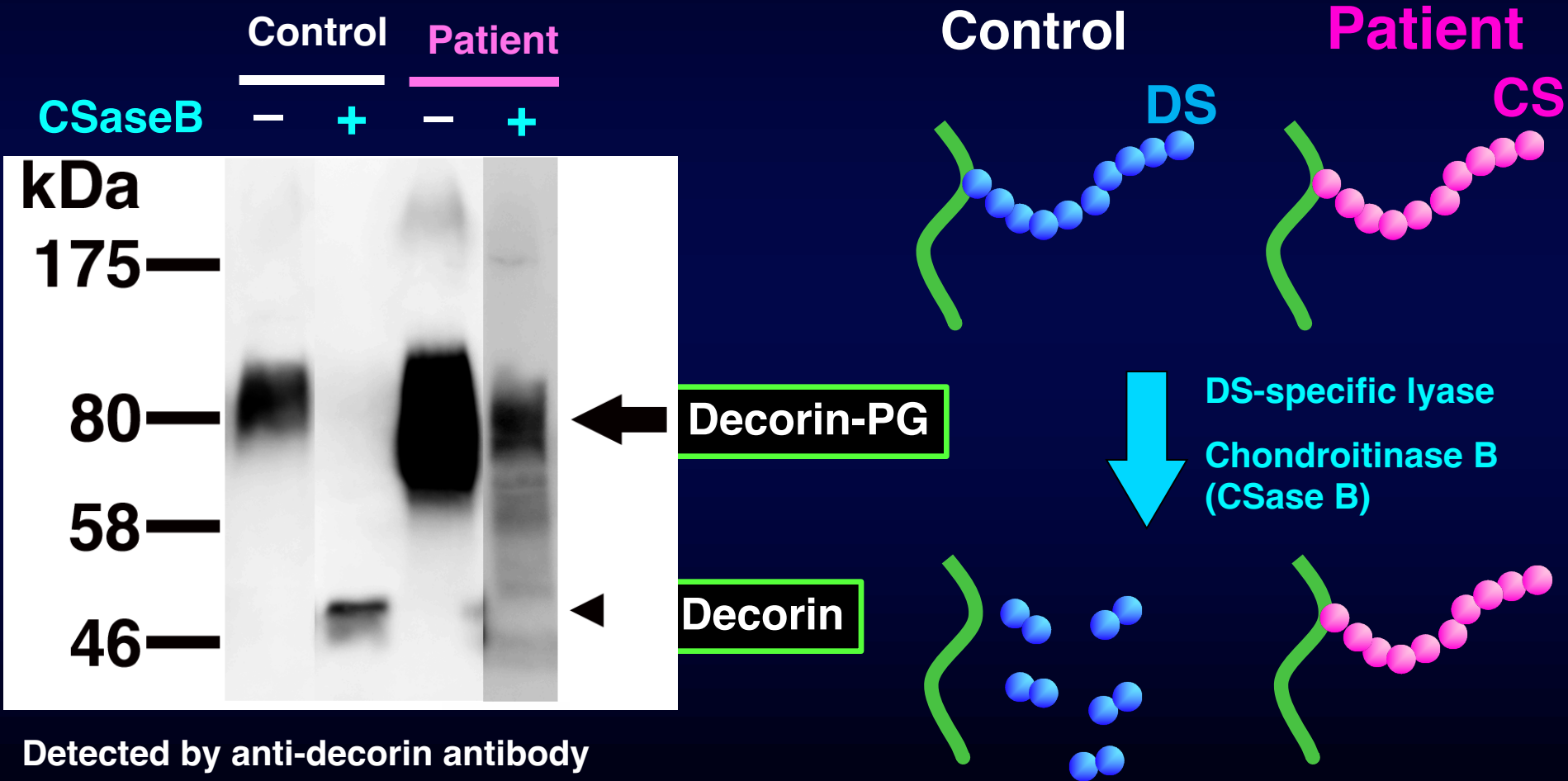
Total amount of CS and DS in fibroblasts from patients and healthy subjects



Various functions of CS, DS, and HS-proteoglycans at cell surfaces and in extracellular matrix



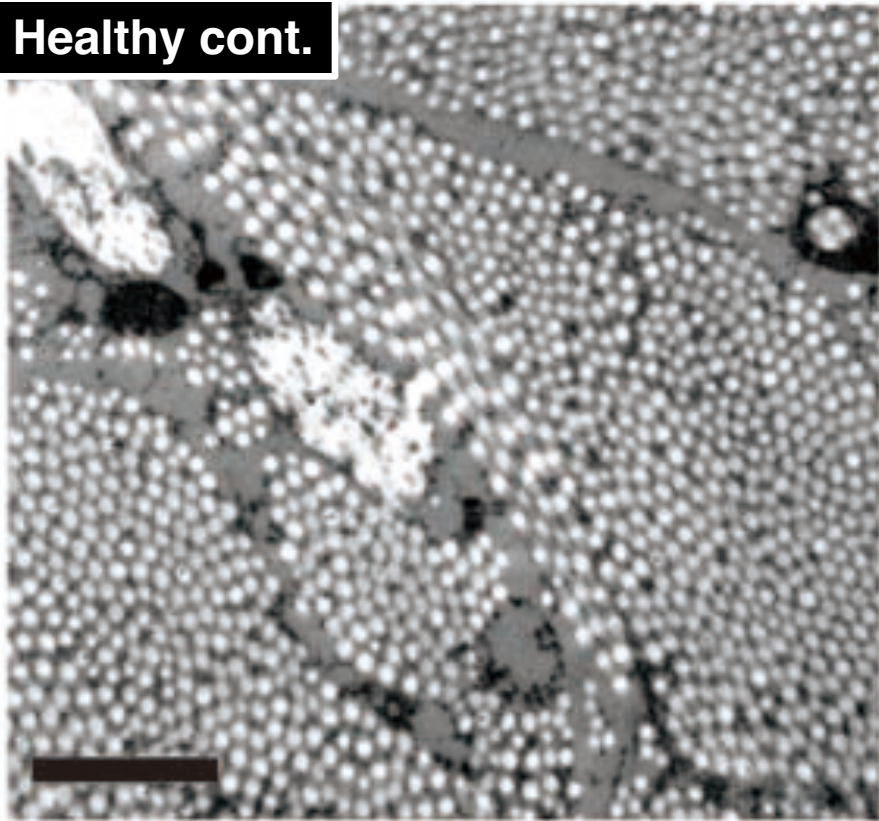
Western blotting of decorin proteoglycan (PG) secreted by fibroblasts from a patient



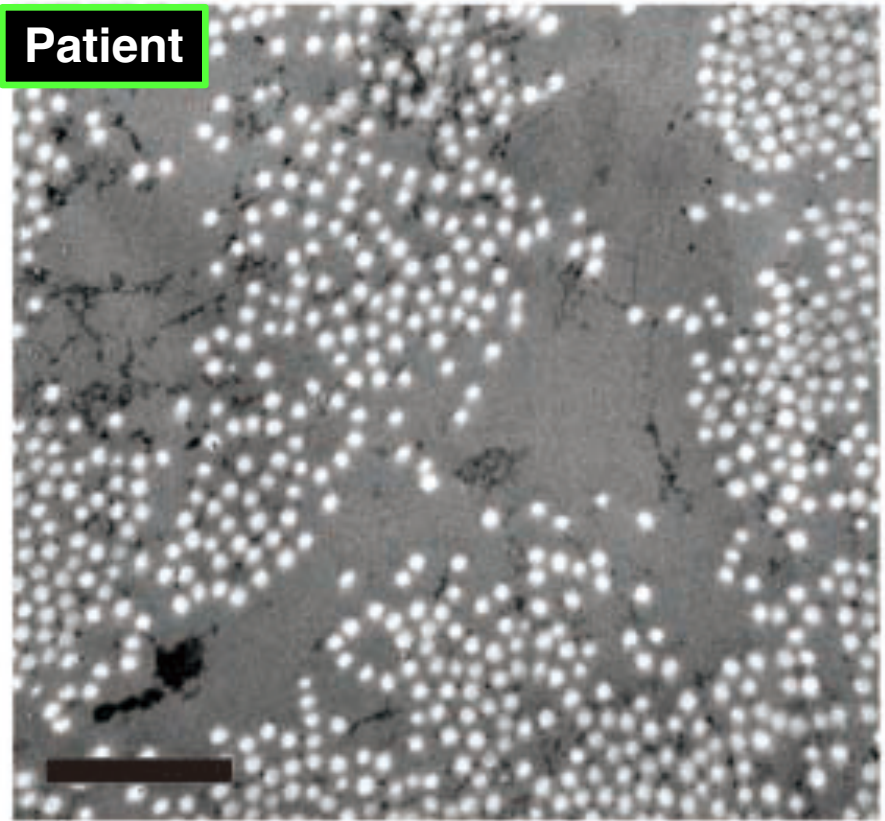
The collagen fibers in the patient

Electron microscopy

Healthy cont.



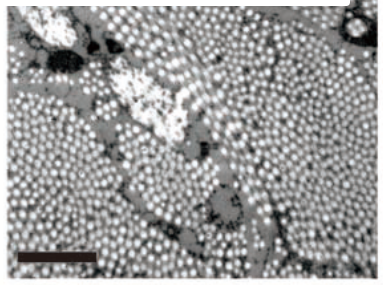
Patient



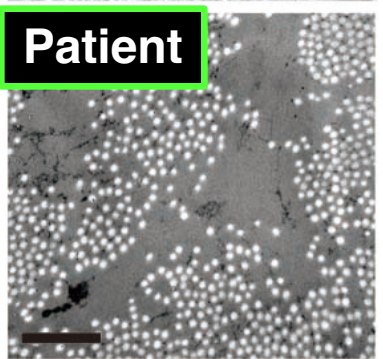
The collagen fibers in the patient

Electron
microscopy

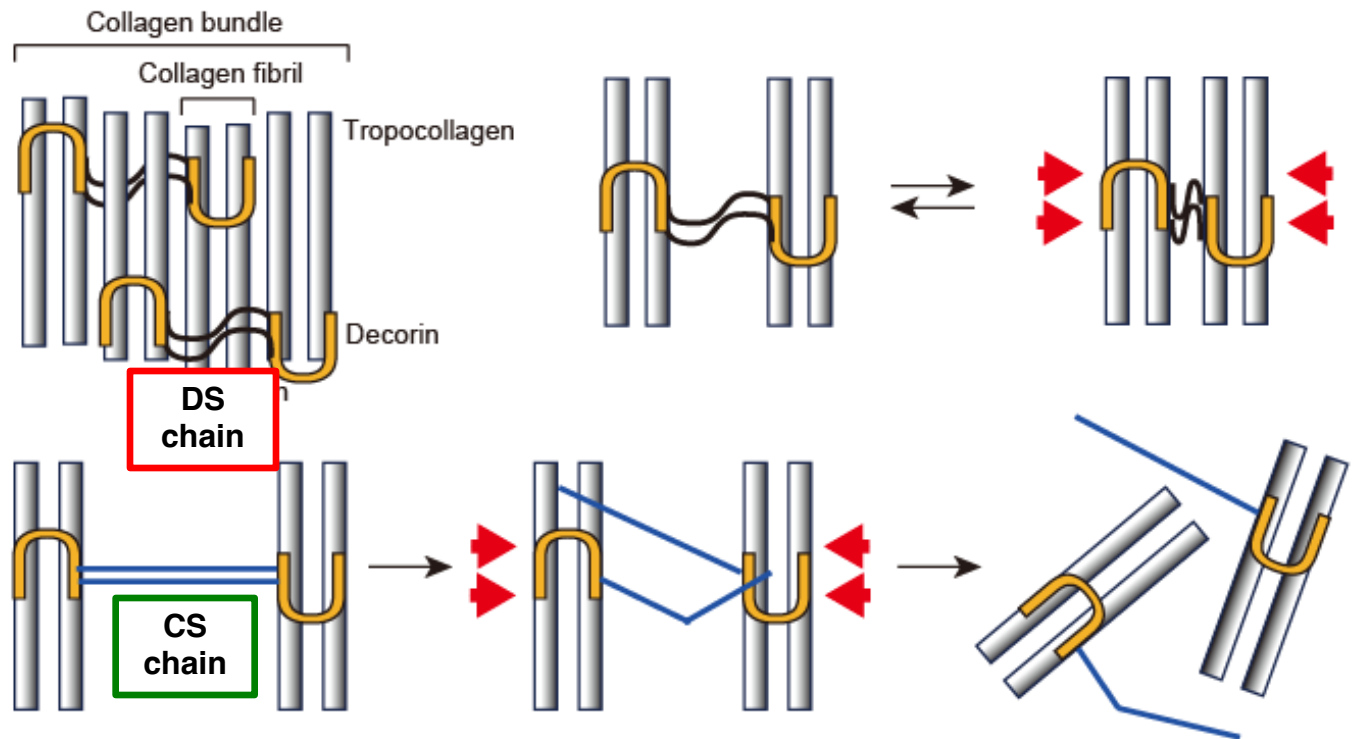
Healthy cont.



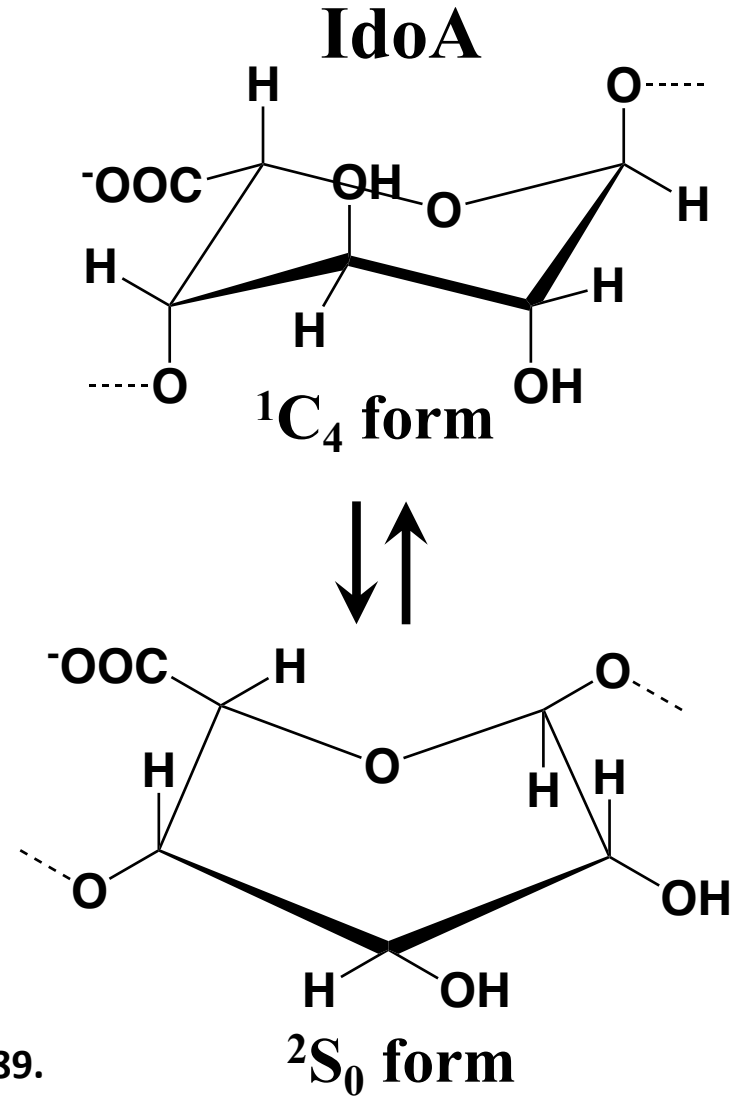
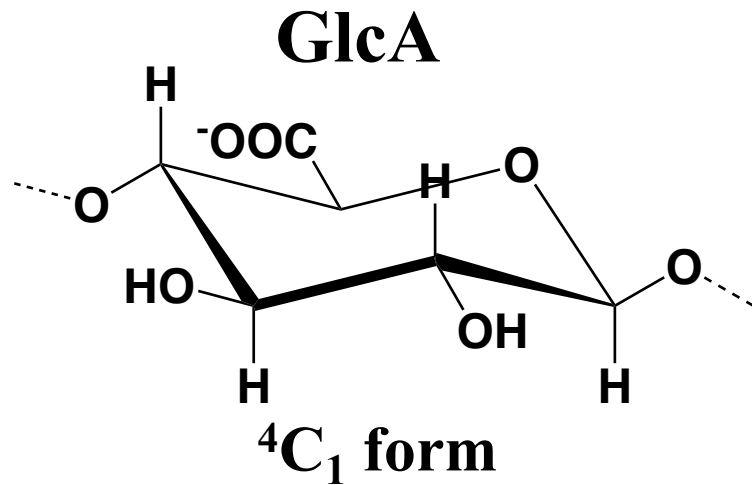
Patient



The structural alterations of collagen fibers
by mechanical compression in normal and affected states



Conformation of GlcA and IdoA

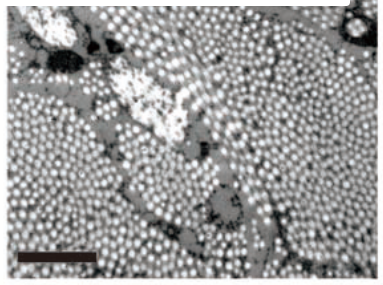


"Heparin" (Lane and Lindahl eds.) , 1989.

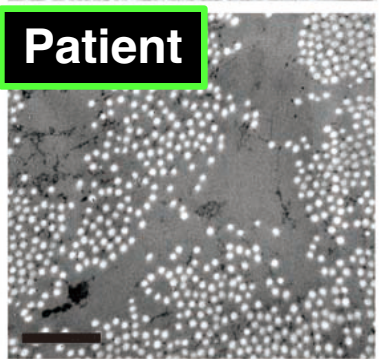
The collagen fibers in the patient

Electron
microscopy

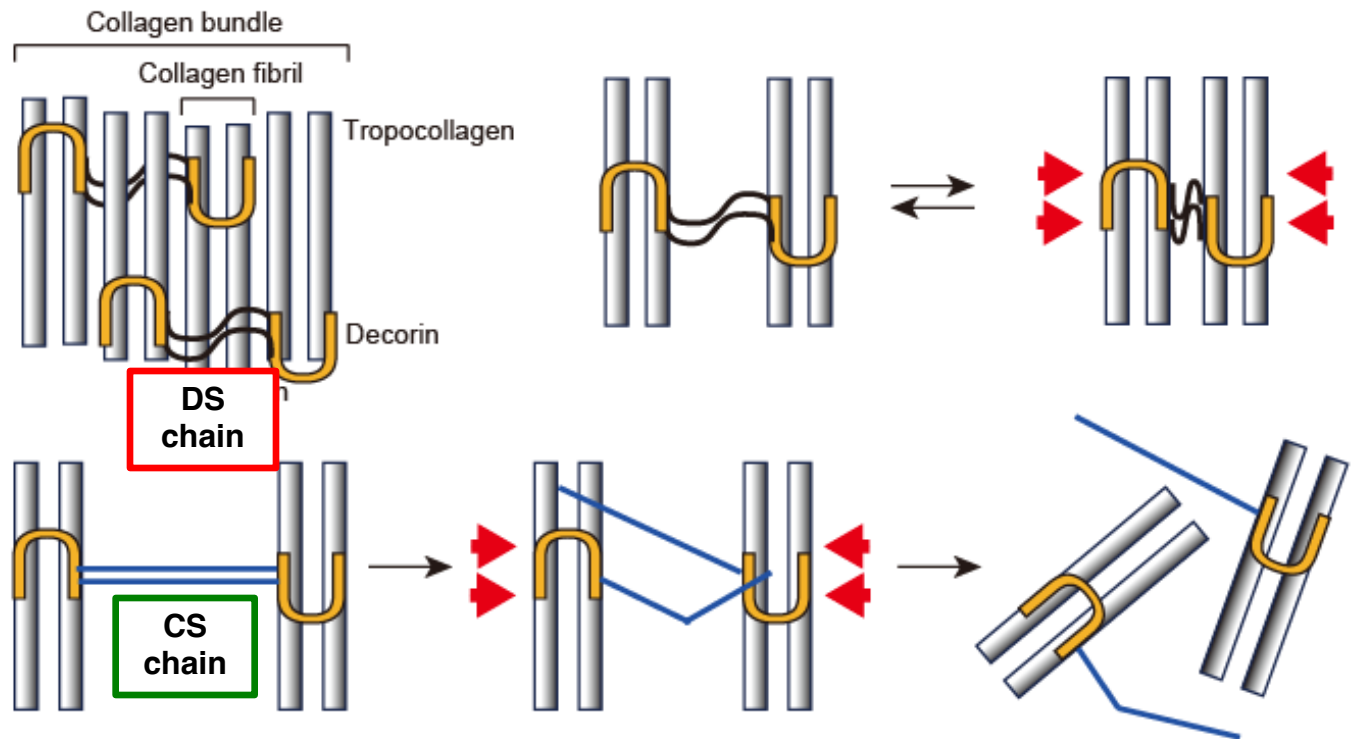
Healthy cont.



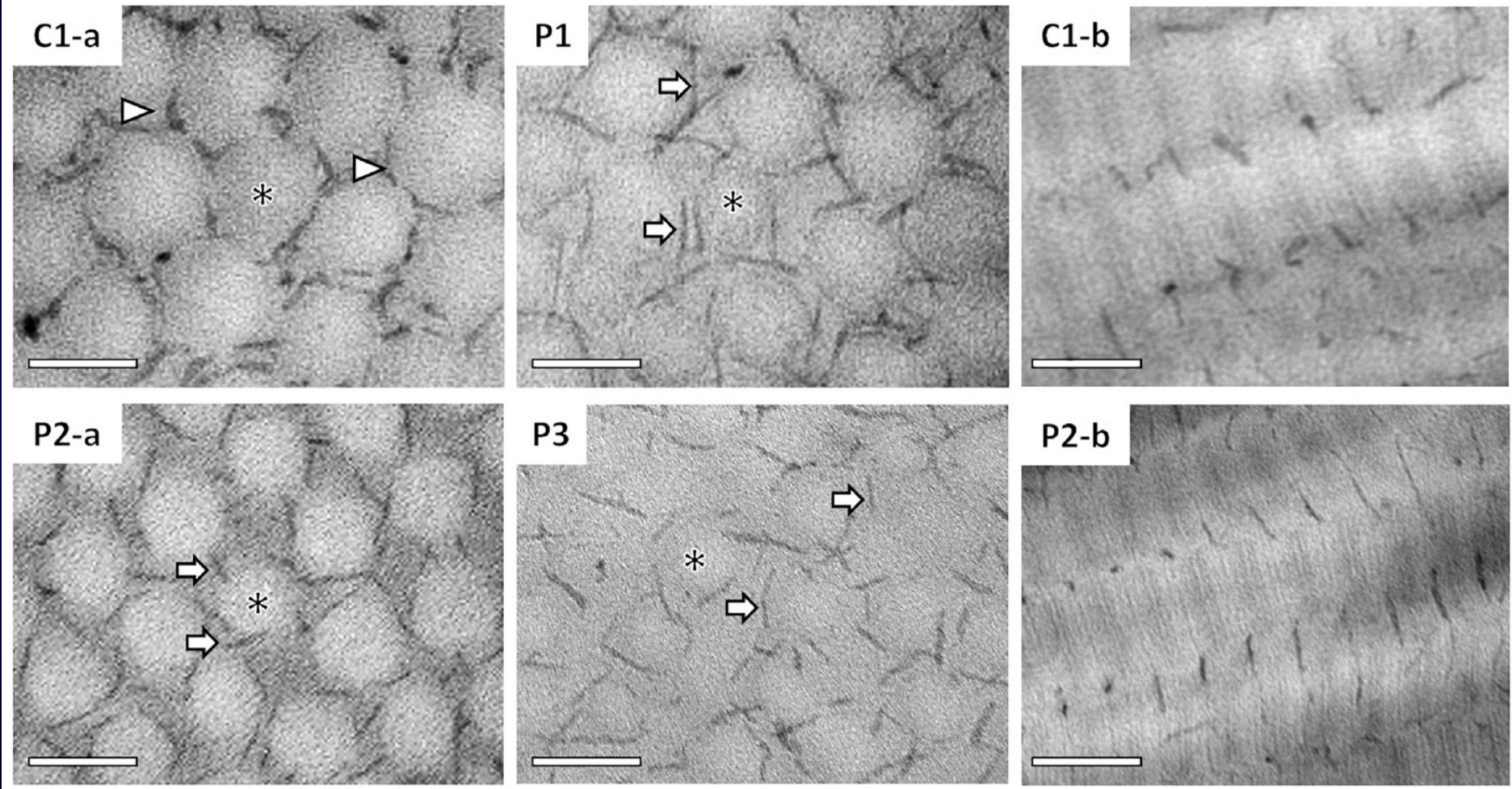
Patient



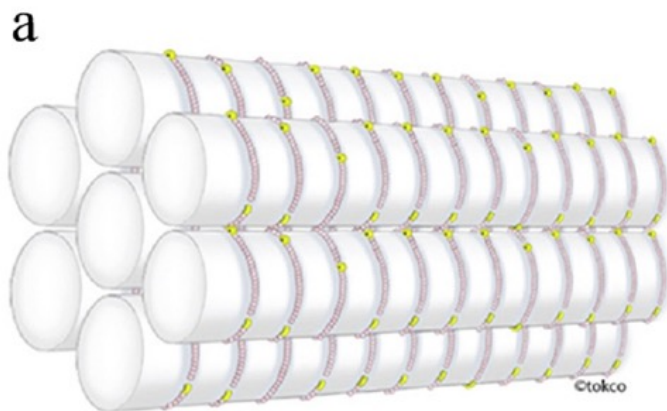
The structural alterations of collagen fibers
by mechanical compression in normal and affected states



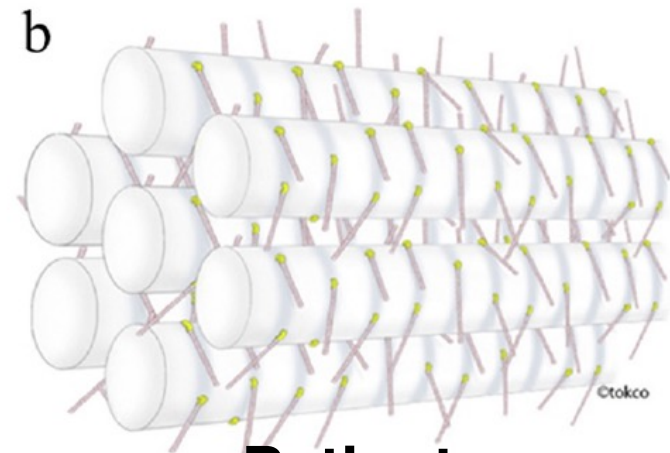
Structural alteration of glycosaminoglycan side chains in the skins of patients



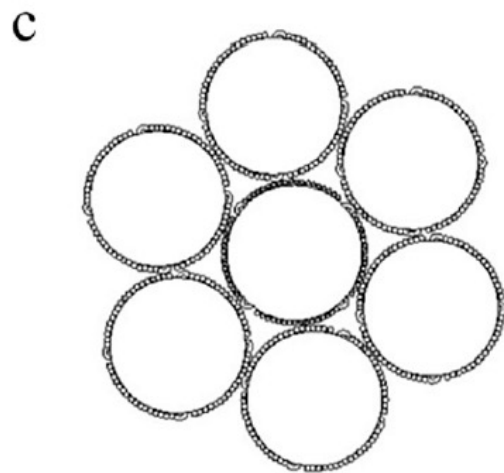
Structure of collagen fibrils and GAG chains in the skin of healthy individuals and patients



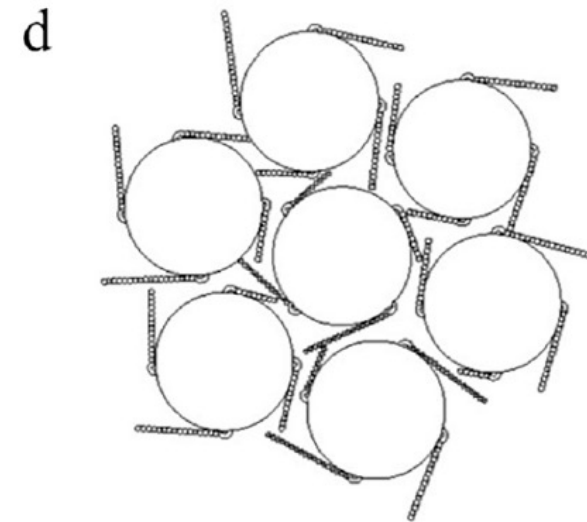
Healthy subjects



Patients



DS chains adhere to collagen fibrils.

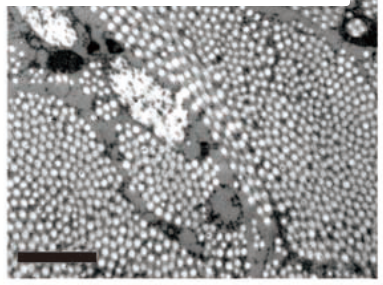


CS chains spread across interfibrillar spaces.

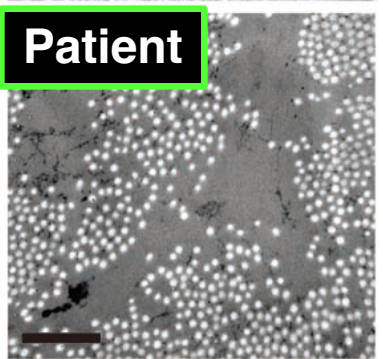
The collagen fibers in the patient

Electron
microscopy

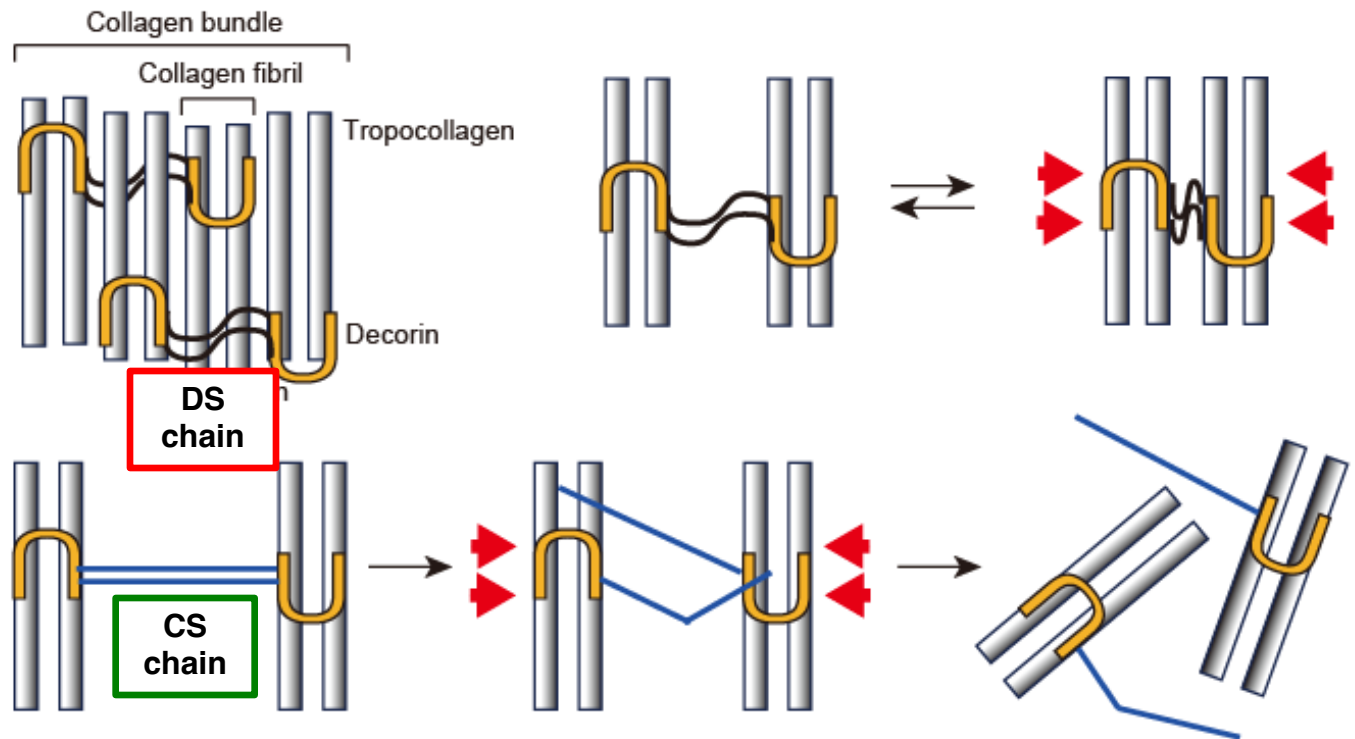
Healthy cont.



Patient



The structural alterations of collagen fibers
by mechanical compression in normal and affected states



Urine samples from EDS patients with mutation in *D4ST1*

Patient	Age (y)	Sex	D4ST1	References
1	11	F	P281L/Y293C	P1 in Kosho <i>et al.</i> AJMG 2010
2	29	F	P281L/P281L	P2 in "
3	32	M	P281L/P281L	P3 in "
4	20	F	P281L/C289S	P5 in "
5	4	F	P281L/Y293C	P6 in "
6	41	F	F209S/P281L	Kono <i>et al.</i> Acta Derm Venereol 2016
7	10	M	F209S/P281L	P2 in Shumizu <i>et al.</i> AJMG 2011
8	3m	M	F209S/P281L	Brother of patient 7

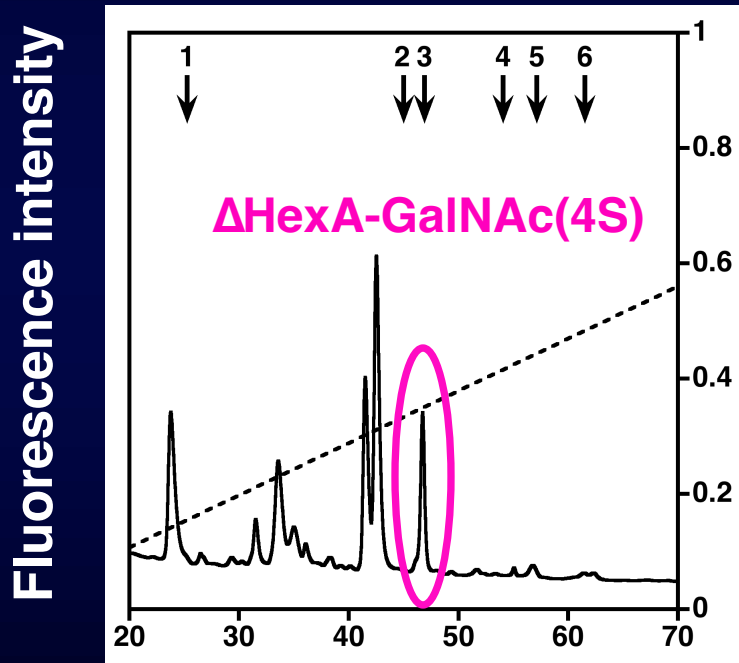
Urines from healthy subjects (15 samples)

M: 6m, 10, 11, 30, 31y

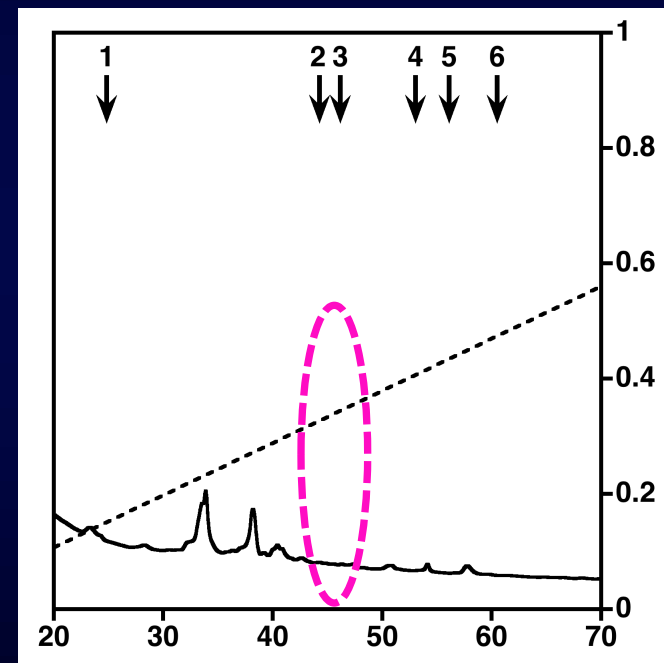
F: 3, 10, 12, 18, 21, 29, 39, 43y

Analysis of DS in urine from a patient and a healthy subject

Control 3



Patient 2 (P281L/P281L)

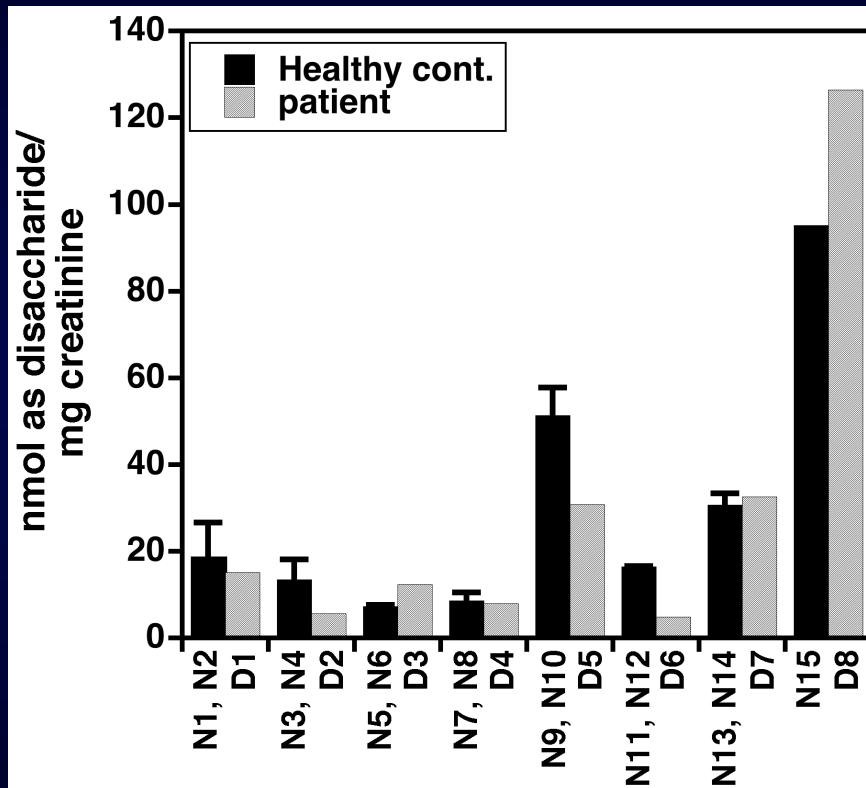


Retention time (min)

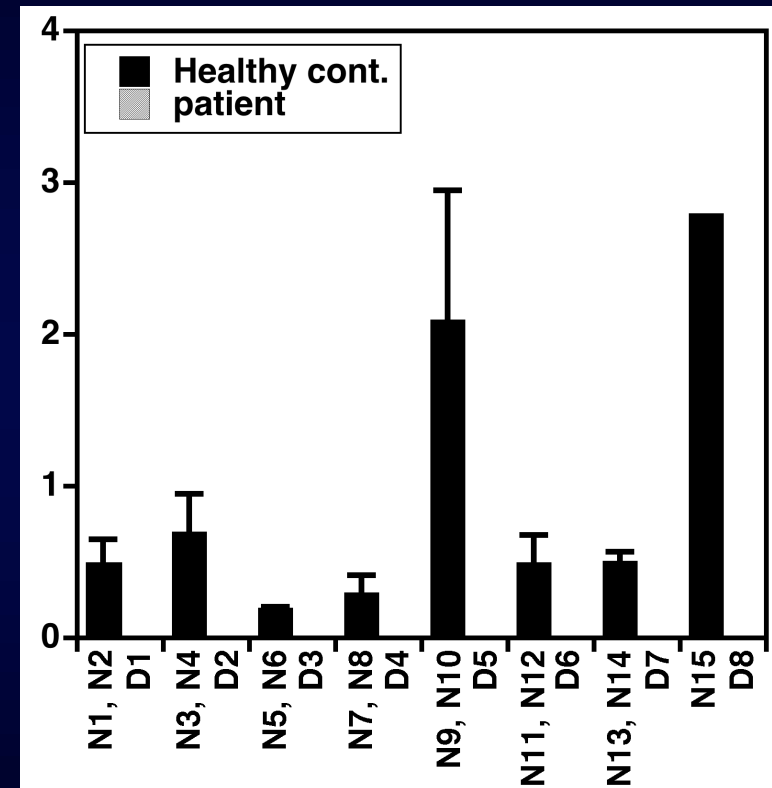
Concentration of NaH₂PO₄ (M) (----)

Total disaccharides of urinary CS and DS from the EDS patients with mutation in *D4ST1*

Chondroitin sulfate



Dermatan sulfate



This result proposes the usefulness of a urinary disaccharide compositional analysis of CS/DS chains as a non-invasive screening method for this disorder.

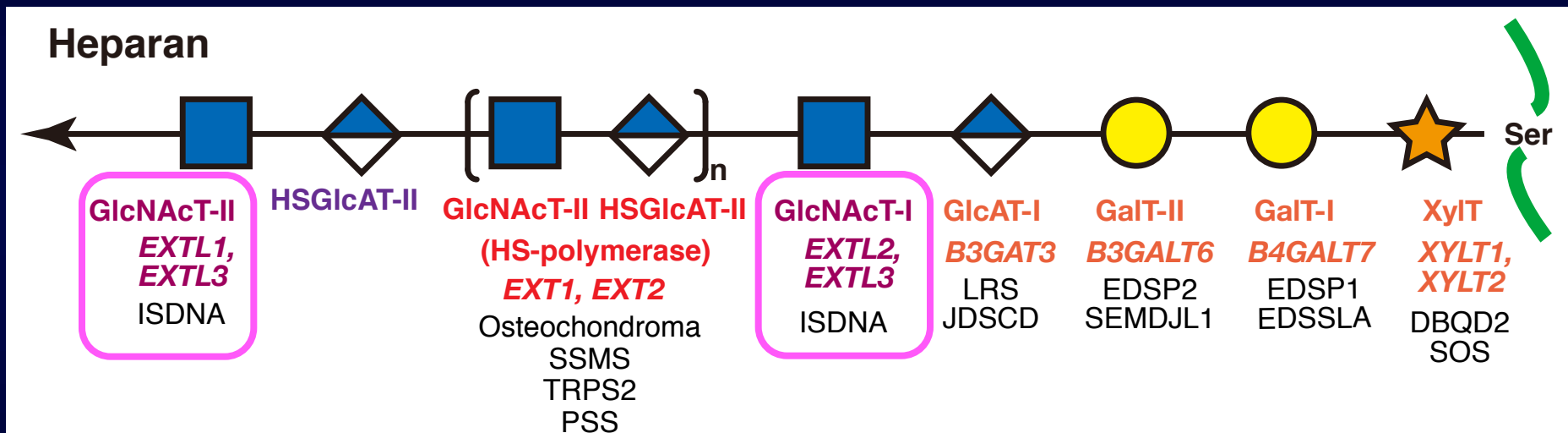
① Ehlers-Danlos syndrome caused by mutations in D4ST-1

② A novel type of spondylo-epi-metaphyseal dysplasia caused by mutations in EXTL3

Biosynthesis of HS backbone

Repeating disaccharide region of HS

Linker tetra-saccharide region



Human tumor suppressor *EXT* gene family members *EXTL1* and *EXTL3* encode α 1,4-*N*-acetylglucosaminyltransferases that likely are involved in heparan sulfate/heparin biosynthesis

Byung-Taek Kim, Hiroshi Kitagawa, Jun-ichi Tamura, Toshiyuki Saito, Marion Kusche-Gullberg, Ulf Lindahl, and Kazuyuki Sugahara

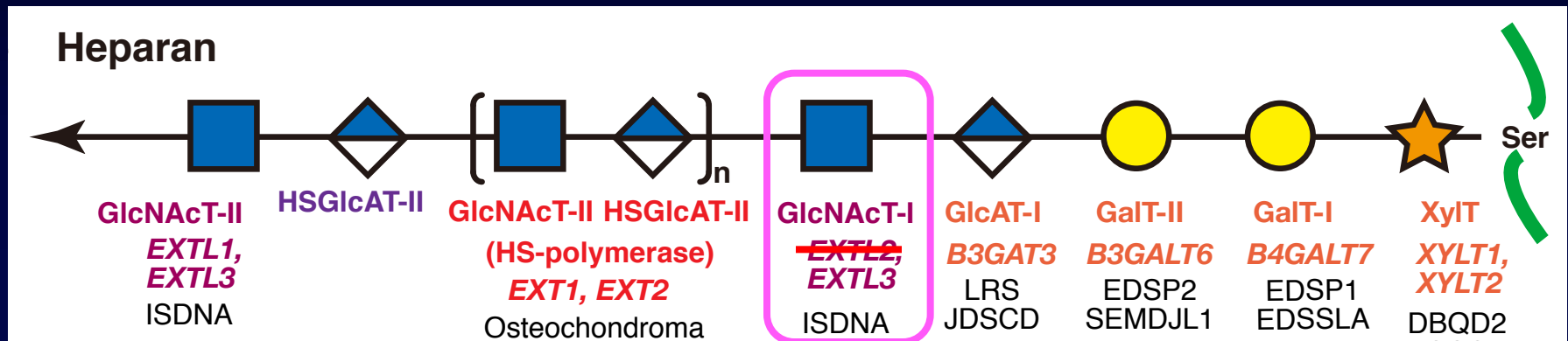
Systemic *Extl3*-knockout mice are embryonic lethal

	<i>Extl3</i> ^{+/+}	<i>Extl3</i> ^{+/-}	<i>Extl3</i> ^{-/-}	Total
E6.5	3 (19 %)	8 (50 %)	5 (31 %)	16
E7.5	4 (33 %)	4 (33 %)	4 (33 %)	12
E8.5	6 (22 %)	16 (59 %)	5 (19 %)	27
E9.5	12 (26 %)	33 (69 %)	1 (2 %)	46
E10.5	10 (31 %)	22 (69 %)	0	32
5-9 weeks (male)	21 (31 %)	47 (69 %)	0	68
5-9 weeks (female)	14 (31 %)	31 (69 %)	0	45

Biosynthesis of HS backbone

Repeating disaccharide region of HS

Linker tetrasaccharide region



EXTL3 seems to be responsible for initiating the chain elongation of HS polysaccharides.

Congenital diseases caused by mutations in the human EXTL3 gene

Guo *et al.*, J Hum Genet, 2017

Oud *et al.*, Am J Hum Genet, 2017

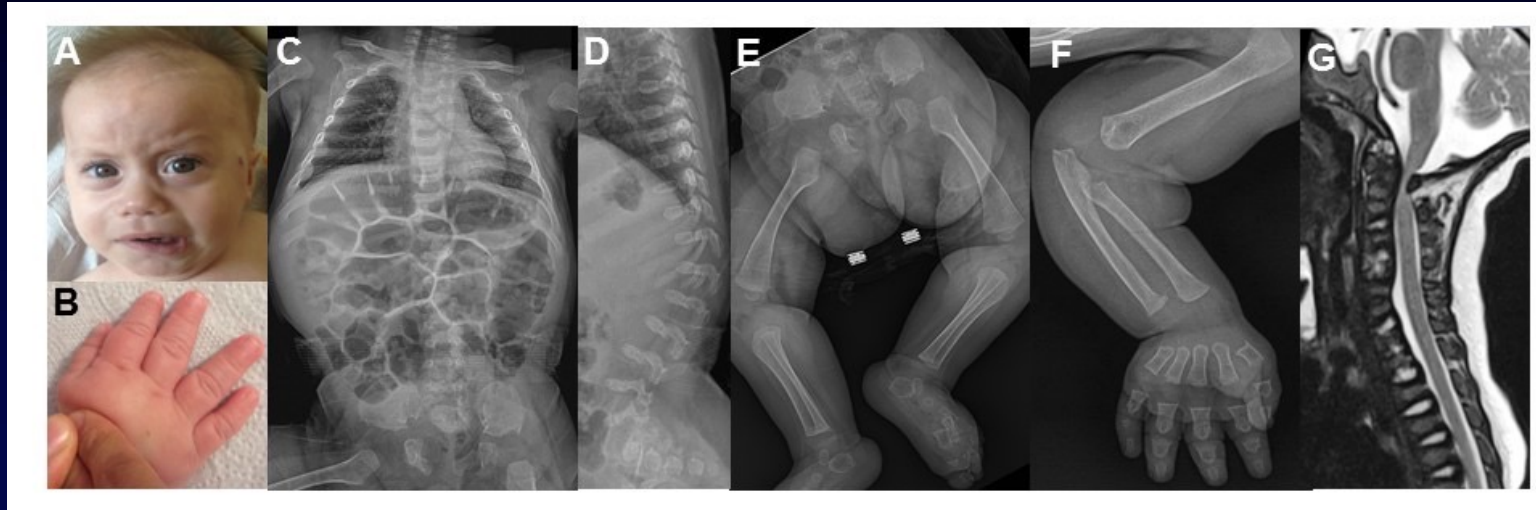
Volpi *et al.*, J Exp Med, 2017

Affected individuals presented with

- 1) various skeletal abnormalities,**
- 2) neurodevelopmental defects, and**
- 3) T-cell immunodeficiencies.**

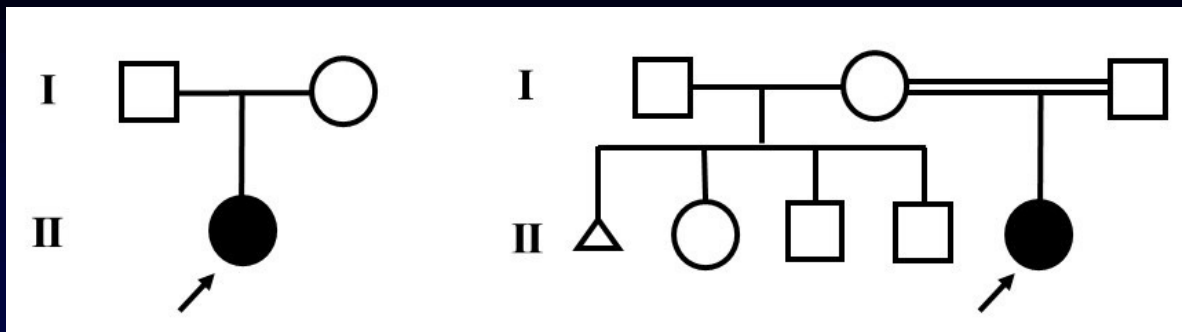
Spondyloepimetaphyseal dysplasia (SEMD) with immunodeficiency/ Neuro-immuno-skeletal dysplasia syndrome caused by mutation in *EXTL3*

Patient 1 (5 months)

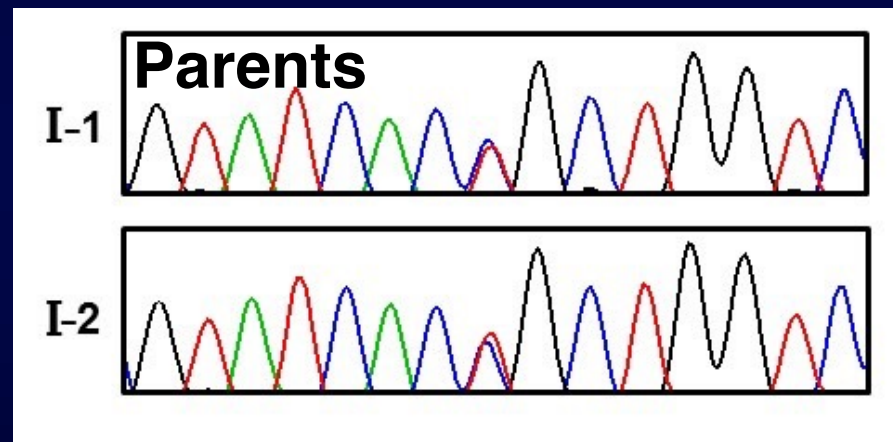
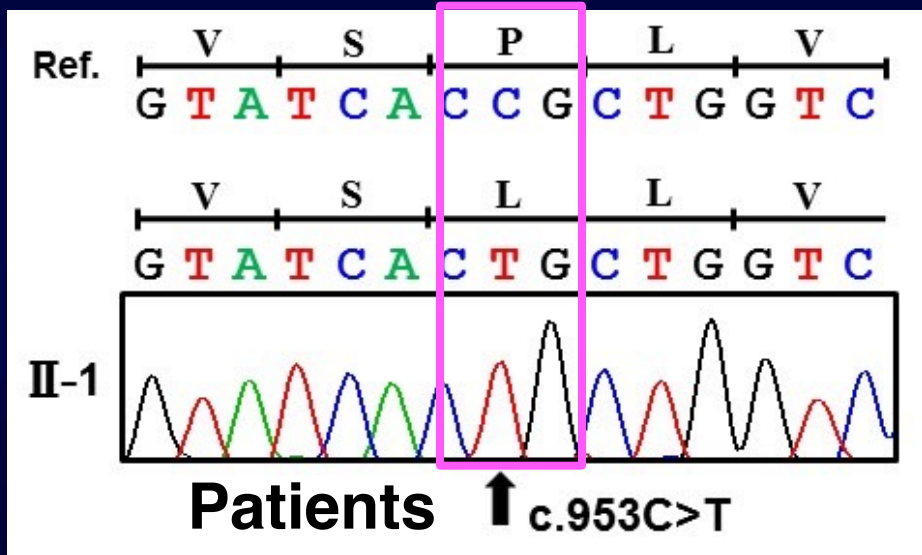


- Clinical pictures (A, B): frontal bossing, prominent eye, depressed nasal bridge, micrognathia and severe brachydactyly
- Radiographs (C-G): broad thorax, severe platyspondyly with kyphosis, broad ilia, broad ischia and pubes, broad metaphyses of the long tubular bones

EXTL3 mutation



EXTL3

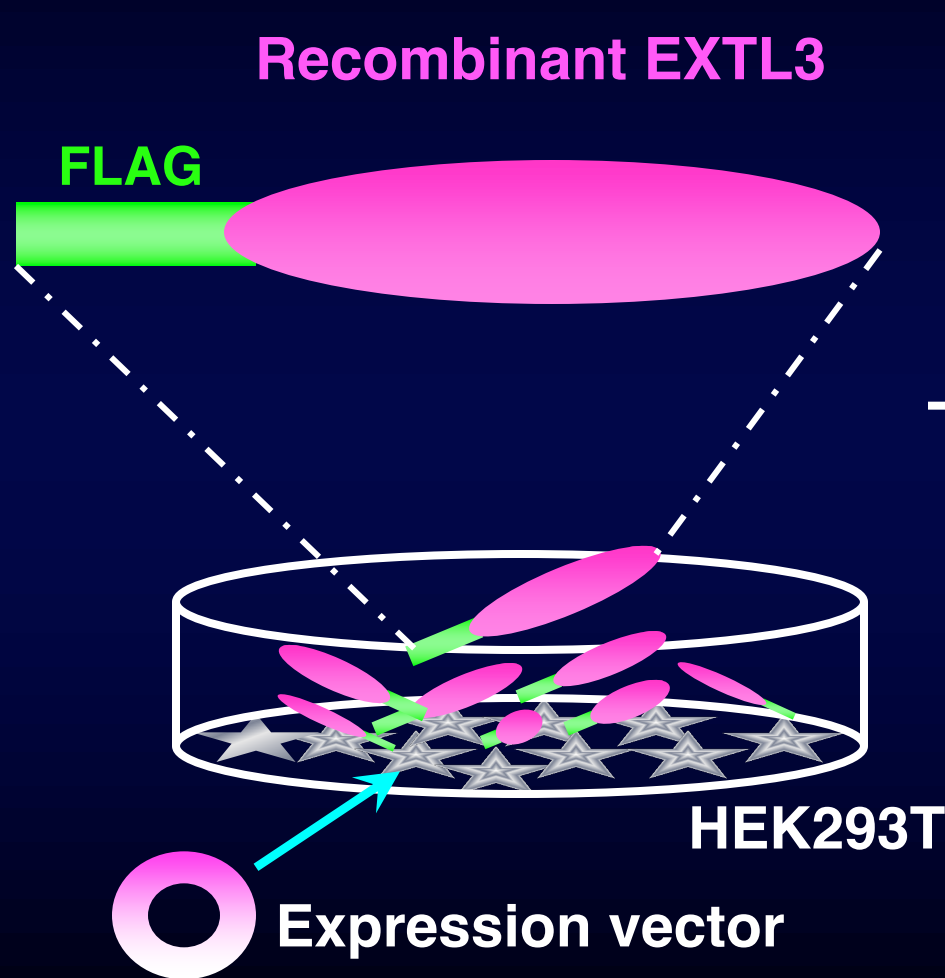


P318L

human
mouse
rat
bovine
Xenopus
zebrafish

267	RTDGHNEVI	INLSRKS	SDTONLL	YNVST	GRAMVA	QSTFY	TVQYR	PGFDL	VVSP	IVHAM	SEPNF	MEIPP	QVPV	KRKYLF	TFQ	GEKIE	SLRSS
267	RTDGHNEVI	INLSRKS	SDTONLL	YNVST	GRAMVA	QSTFY	YAAQY	RAGFDL	VVSP	IVHAM	SEPNF	MEIPP	QVPV	KRKYLF	TFQ	GEKIE	SLRSS
267	RTDGHNEVI	INLSRKS	SDTONLL	YNVST	GRAMVA	QSTFY	YAAQY	RAGFDL	VVSP	IVHAM	SEPNF	MEIPP	QVPV	KRKYLF	TFQ	GEKIE	SLRSS
267	RTDGHNEVI	IISLRKS	SDTONLL	YNVST	GRAMVA	QSTFY	YAAQY	RPGFDL	VVSP	IVHAM	SEPNF	MEIPP	QVPV	KRKYLF	TFQ	GEKIE	SLRSS
267	RTDGHNELI	INLSRK	SETQNY	LYNVST	GRAMIA	QSTFY	YDSQY	RPGFDI	VVSP	IVHAM	SEPNF	LEIPP	QVPV	KRKYLF	SFQ	GEKIE	SLHSS
263	RSDGHNELL	VHLSIN	SLTONF	LYNVST	GRAAVA	QSTFF	ERYQY	REGFDL	VVSP	IVHAL	SEPNF	LEVPP	QVPV	KRKYLF	TFQ	GEKV	ESLRSS

Expression of recombinant EXTL3 (Wild-type, P318L) and measurement of the enzyme activity



Purified EXTL3	10 μ L
UDP-[3 H]GlcNAc	8.3 μ M
Heparan sulfate	10 μ g
MES-NaOH (pH 6.5)	50 mM
ATP	0.17 mM
MnCl ₂	10 mM
<hr/>	
Total	30 μ L

37 °C for 2h

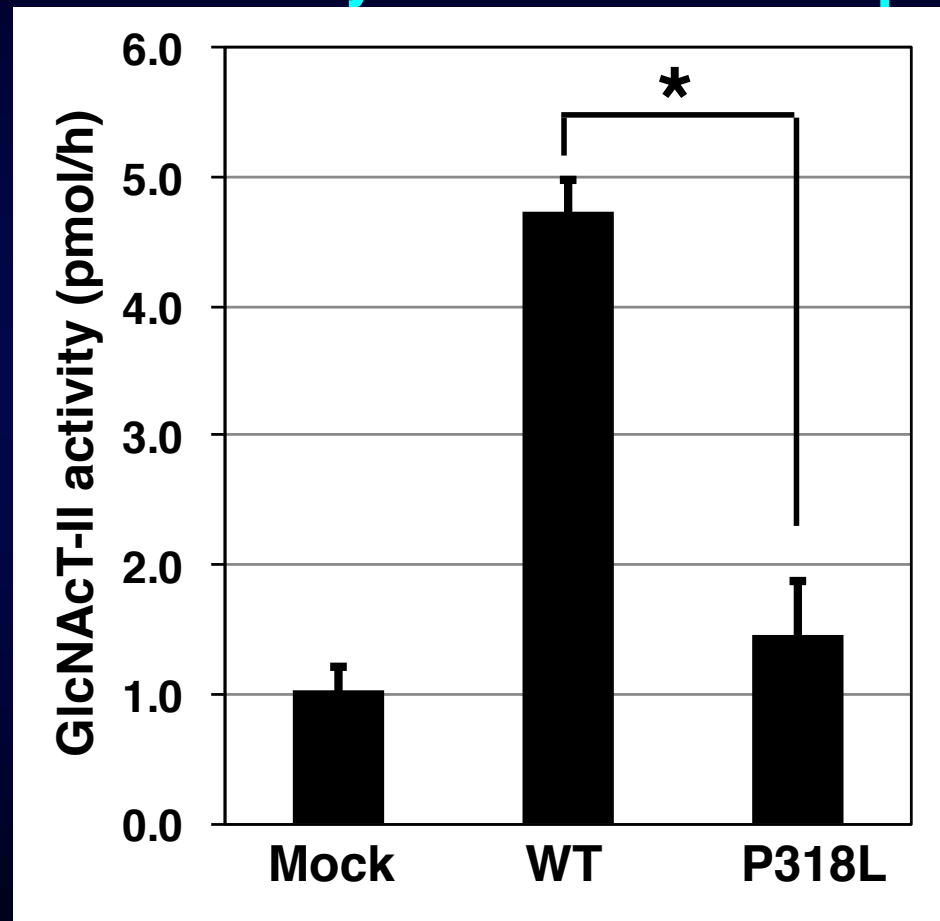
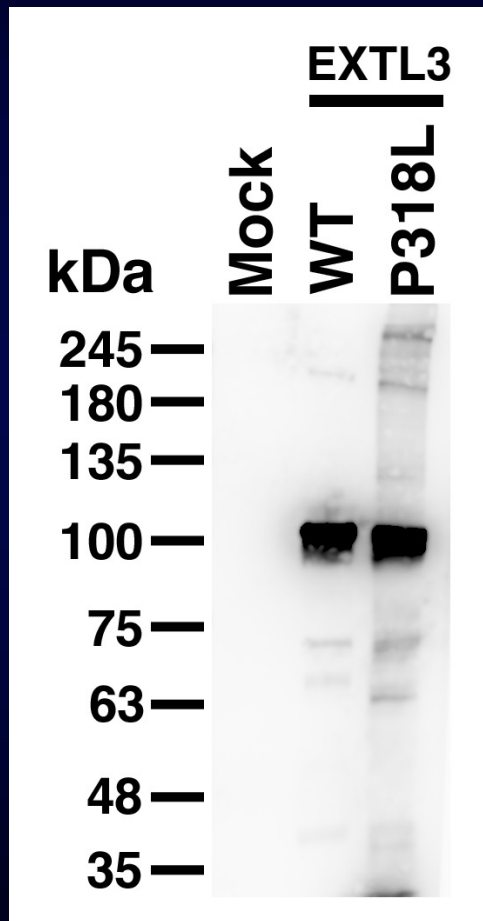
Gel filtration

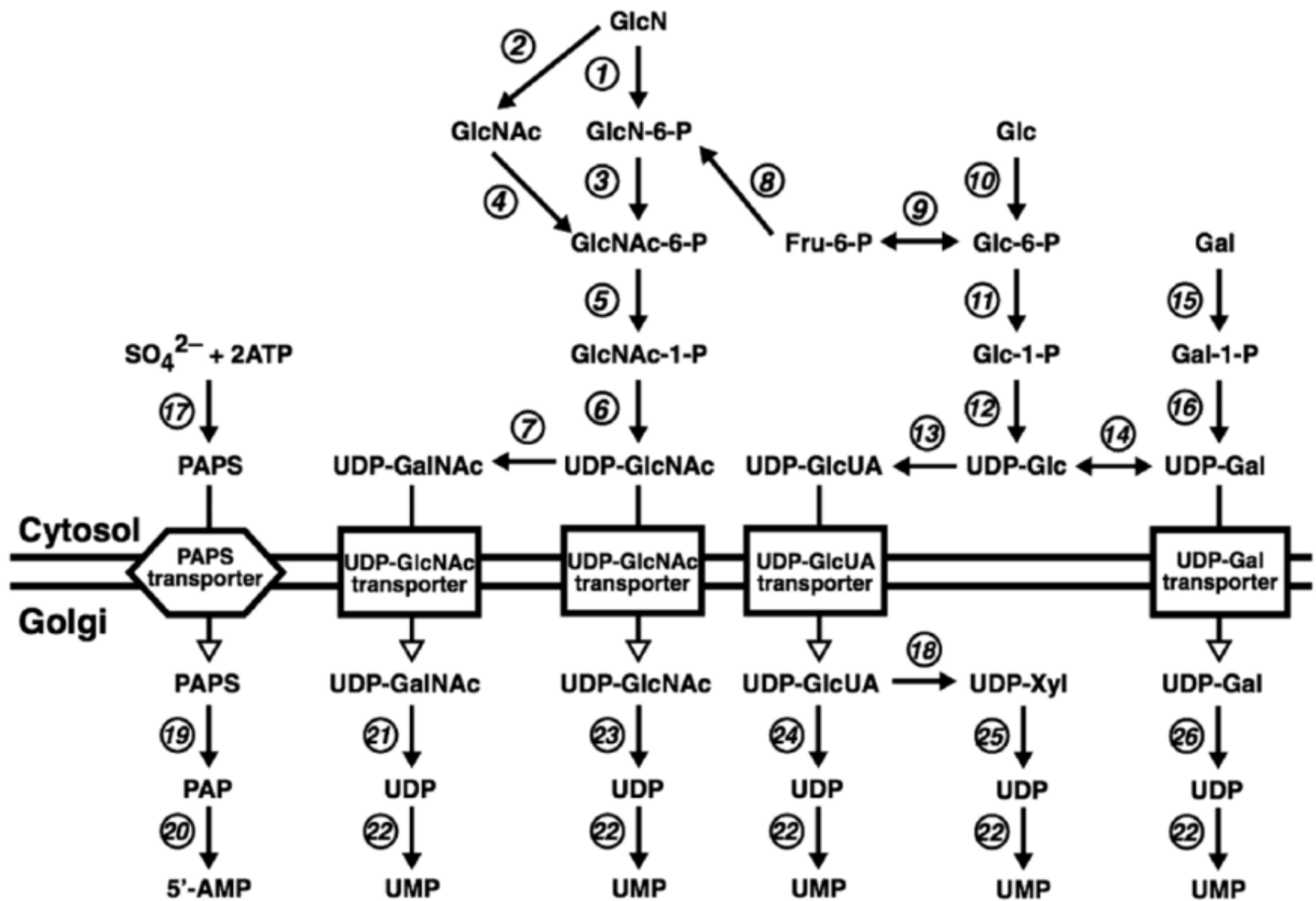
[3 H]GlcNAc-HS polymer

Liquid Scintillation Counting

GlcNAc-transferase activity of recombinant EXTL3 (Wild-type, P318L)

Western blotting GlcNAc-T activity of recombinant proteins





CONCLUSIONS

- **Glycosaminoglycans are involved in various diseases including cancer, spinal cord injury, and infectious diseases.**
- **Genetic disorders caused by deficiency of catabolism of glycosaminoglycans (mucopolysaccharidoses) have been well investigated.**
- **Genetic disorders caused by deficiency of anabolism of glycosaminoglycans have recently been found one by one.**
- **There is an urgent need to develop treatments for such genetic diseases.**

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Istanbul, Turkey)

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