

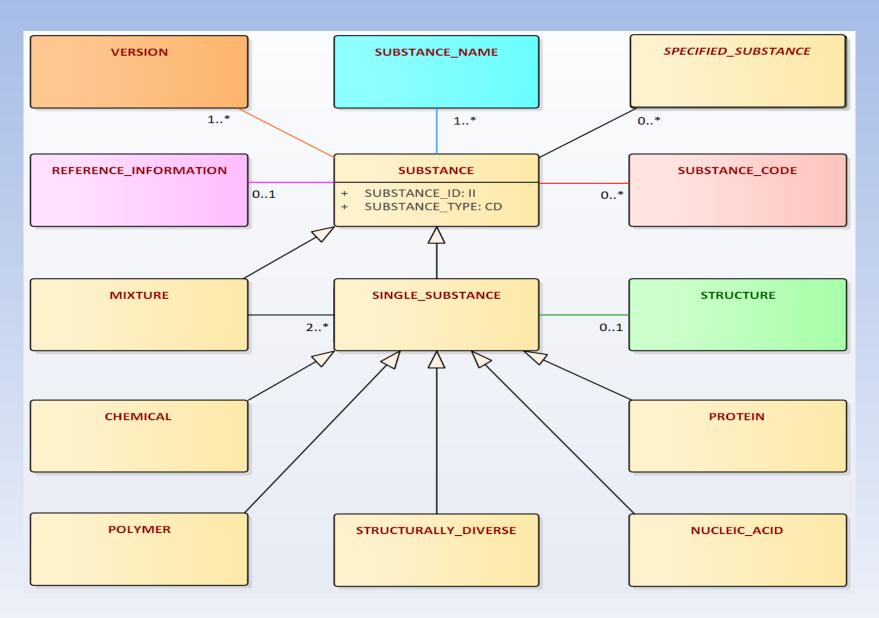
Signature fields (Minimum requirements) for a Protein Substance Registration

GInAS Meeting, 30 October 2019 Herman Diederik

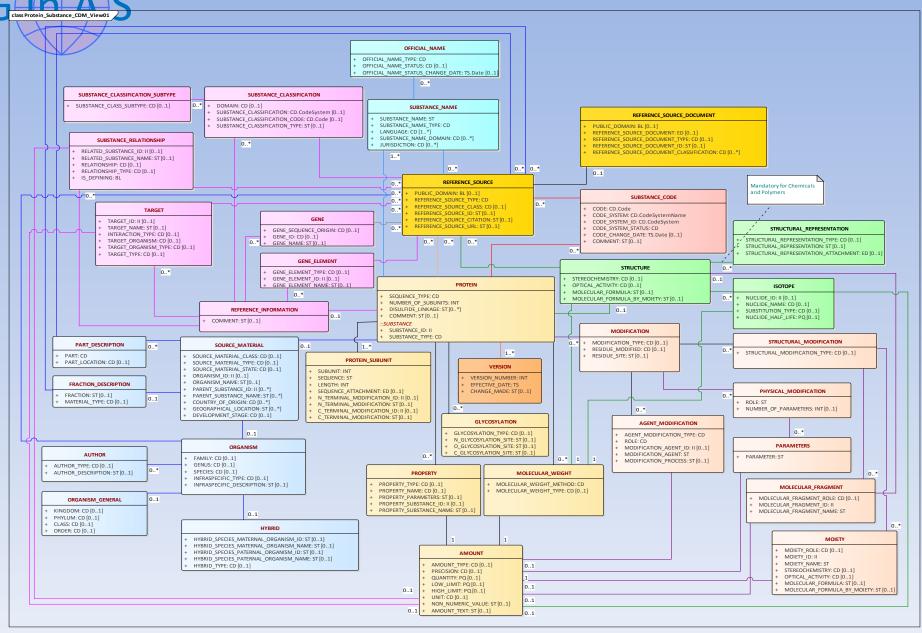
(h.diederik@cbg-meb.nl; hdiederik@planet.nl)



ISO 11238/19844 Substance Classification Overview (high-level)

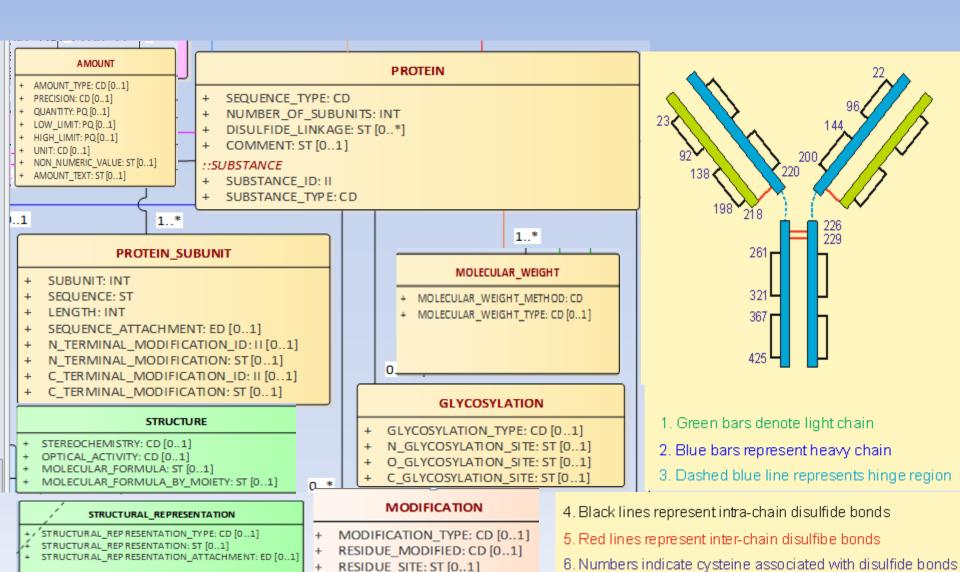


Grand S Protein Information model



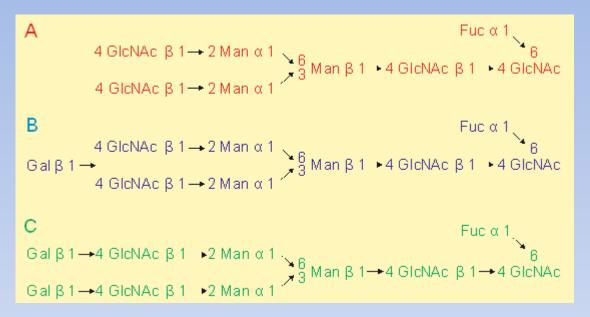


Characterising elements at the Substance info level for an IGG1

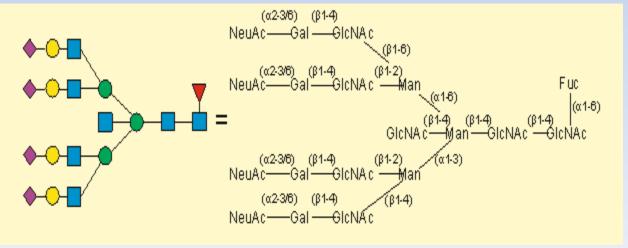


Predominant Glycan Information at the Substance level (A)

Characterising elements at the Specified Substance Group 1 info level: Glycan Information: Structural representation of common Glycans



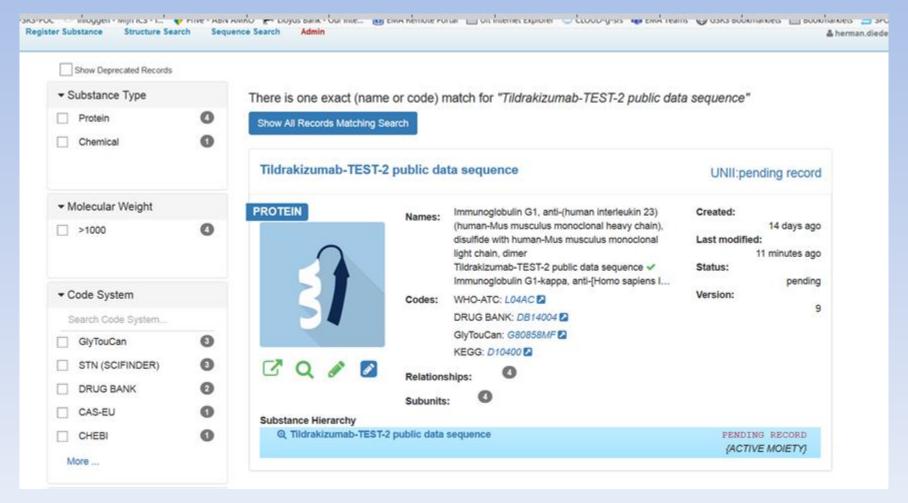
- N-Acetylneuraminic acid (NANA)
- 🔾 Galactose (Gall)
- N-Acetylglucosamine (GlcNAc)
- 🔵 Mannose (Man)
- 🛕 Fucose (Fuc)





Minimum fields based on Public data Example Tildrakizumab

 Record Overview prepared in NCATS 2.3.3. version Tildrakizumab-TEST-2-public data sequence





Overview and Names/ Codes

Definition information:

Public data Sources: INN List 108 WHO Drug Information, Vol. 26, No. 4, 2012 Proposed INN: List 108

Immunoglobulin G1-kappa, anti-[Homo sapiens IL23A (interleukin 23 alpha subunit (p19), IL-23A)], humanized monoclonal antibody; gamma1 heavy chain (1-446) [humanized VH (Homo sapiensIGHV1-18*01 (81.60%) -(IGHD)-IGHJ4*01)) [8.8.9] (1-116) -Homo sapiens IGHG1*01 (CH1 (117-214, hinge (215-229), CH2 (230-339), CH3 (340-444), CHS (445-446)) (117-446)], (219-214')-disulfide with kappa light chain (1'-214') [humanized V-KAPPA (Homo sapiensIGKV1-39*01 (85.30%) -IGKJ1*01) [6.3.9] (1'-107') -Homo sapiens IGKC*01 (108'-214')]; dimer (225-225":228-228")-bisdisulfide immunomodulator

Other Sources: WHO-ATC: <u>L04AC</u>; DRUG BANK: <u>DB14004</u>; GlyTouCan: <u>G80858MF</u>; KEGG: <u>D10400</u>

Web: https://newdrugapprovals.org/2018/06/01/tildrakizumab-asmn/



Detailed Overview

Tildrakizumab-TEST-2 public data sequence

UNII:pending record

PROTEIN



Names: Immunoglobulin G1, anti-(human interleukin 23) (...

Tildrakizumab-TEST-2 public data sequence ✓ Immunoglobulin G1-kappa, anti-[Homo sapiens IL23A (interleukin 23 alpha subunit (p19), IL-23A)], humanized monoclonal antibody; gamma1 heavy chain (1-446) [humanized VH (Homo sapiensIGHV1-18*01 (81.60%) -(IGHD)-IGHJ4*01)) [8.8.9] (1-116) -Homo sapiens IGHG1*01 (CH1 (117-214, hinge (215-229), CH2 (230-339), CH3 (340-444), CHS (445-446)) (117-446)], (219-214')-disulfide with kappa light chain (1'-214') [humanized V-KAPPA (Homo sapiensIGKV1-39*01 (85.30%) -IGKJ1*01) [6.3.9] (1'-107') -Homo sapiensIGKC*01 (108'-214')]; dimer (225-225":228-228")-bisdisulfide

Created:

14 days ago

Last modified:

15 minutes ago

Status:

pending

Version:

9

Codes: WHO-ATC: L04AC 2

DRUG BANK: DB14004 2

GlyTouCan: G80858MF 2

KEGG: D10400 2

immunomodulator

Relationships:





Walk through of the Addition Public Information, besides the INN Definitional information (1)



Tildrakizuwab-aswu

Heavy

chain:

QVQLVQSGAEVKKPGASVKVSCKASGYIFITYWMTWVRQAF

EWMGQIFPASGSADYNEKFEGRVTMTTDTSTSTAYMELRSL

TAVYYCARGGGGFAYWGQGTLVTVSSASTKGPSVFPLAPSS

GGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS

WHO-ATC Code:

ATC/DDD Index

https://www.whocc.no/atc_ddd_in
dex/?code=L04AC

WHO Collaborating Centre for Drug Statistics Methodology

News

ATC/DDD Index

Immunoglobulin G1, anti-(human interleukin 23) (human-Mus musculus monoclonal heavy chain), disulfide with human-Mus musculus monoclonal light chain, dimer

CAS 1326244-10-3, **BLA 761067**

Tildrakizumab (SCH 900222/MK-3222)

ILUMYA; MK-3222; SCH-900222; SUNPG 1622; SUNPG 1622 I; SUNPG 1623

I; SUNPG 1623 II; SUNPG 1623 III; SUNPG 1623 IV; SUNPG1623;

Tildrakizumab-asmn

DRUG BANK https://www.drugbank.ca/drugs/DB14004

Company Sun Pharmaceuticals

Approval Status FDA Approved March 2018 FOR Psoriasis, plaque

Treatments plaque psoriasis

Protein chemical formulaC₆₄₂₆H₉₉₁₈N₁₆₉₈O₂₀₀₀S₄₆

Protein average weight144400.0 DaSequences

L ANTINEOPLASTIC AND IMMUNOMODULATING AGENTS
L04 IMMUNOSUPPRESSANTS
L04AC Interleukin inhibitors



Walk through of the Addition Public Information, besides the INN Definitional information (2)

https://www.kegg.jp/dbget-bin/www bget?dr:D10400

KEGG	DRUG: Tildrakizumab	Help
Entry	D10400 Drug	
Name	Tildrakizumab (USAN); Tildrakizumab (genetical recombination) (JAN); Tildrakizumab-asmn; Ilumya (TN)	
Product	ILUMYA (Sun Pharmaceutical Industries)	
Formula	C6426H9918N169802000S46	
Exact mass	144345.3728	
Mol weight	144434.6737	
Sequence Type	Peptide	
Class	Other DG02019 Interleukin inhibitor	
Remark	ATC code: L04AC17 Product: D10400 <us></us>	
Efficacy	Antipsoriatic, Immunosuppressant, Anti-IL-23 antibody	
Disease	Plaque psoriasis [DS:H01656]	
Comment	Monoclonal antibody Treatment of immunologically mediated inflammatory disorders	
Target	IL23A [HSA:51561] [KO:K05426]	
Pathway	hsa04060 Cytokine-cytokine receptor interaction	
Brite	Anatomical Therapeutic Chemical (ATC) classification [BR:br08303] L ANTINEOPLASTIC AND IMMUNOMODULATING AGENTS L04 IMMUNOSUPPRESSANTS L04AC Interleukin inhibitors L04AC17 Tildrakizumab D10400 Tildrakizumab (USAN) <us> USP drug classification [BR:br08302] Immunological Agents Immunomodulators Tildrakizumab-asmn D10400 Tildrakizumab (USAN)</us>	

GIN AS

Walk through of the Addition Public Information, besides the INN Definitional information (3)

https://www.genome.jp/dbget-bin/www_bget?hsa:51561+D06556+D09214+D09588+D1



interleukin 23 subunit alpha Information Interleukin inhibitor

https://www.genome.jp/dbget-bin/www bget?hsa:51561+D06556+D09214+D09588+D1

Search for details!!

Sequences = Target

>Tildrakizumab Seguence

gcccatggagcagcaaccctgagtccctaa

MLGSRAVMLLLLLPWTAQGRAVPGGSSPAWTQCQQLSQKLCTLAWSAHPLVGHMDLREEG DEETTNDVPHIQCGDGCDPQGLRDNSQFCLQRIHQGLIFYEKLLGSDIFTGEPSLLPDSP VGQLHASLLGLSQLLQPEGHHWETQQIPSLSPSQPWQRLLLRFKILRSLQAFVAVAARVF AHGAATLSP

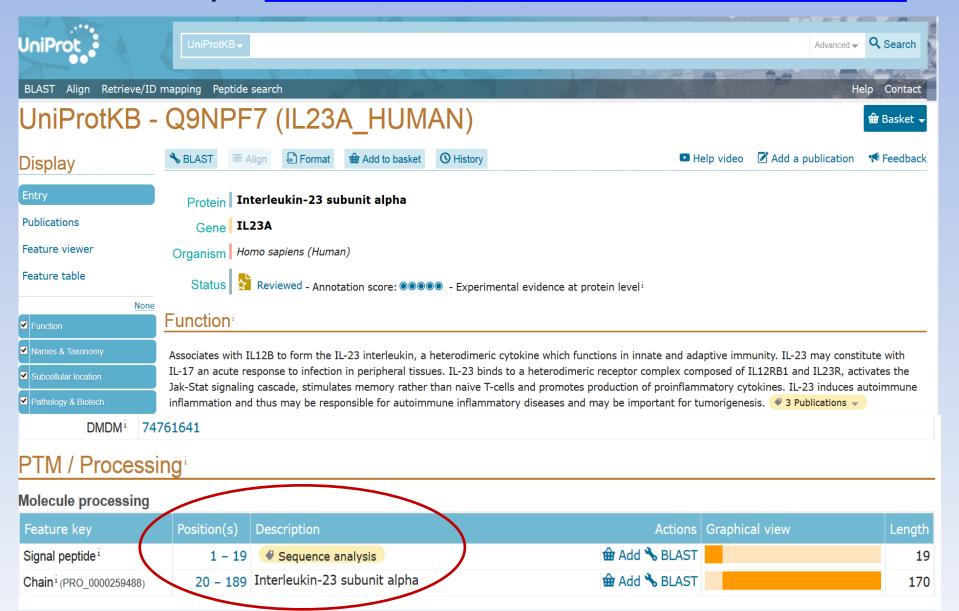
AA seq DB search MLGSRAVMLLLLLPWTAQGRAVPGGSSPAWTQCQQLSQKLCTLAWSAHPLVGHMDLREEG DEETTNDVPHIQCGDGCDPQGLRDNSQFCLQRIHQGLIFYEKLLGSDIFTGEPSLLPDSP VGQLHASLLGLSQLLQPEGHHWETQQIPSLSPSQPWQRLLLRFKILRSLQAFVAVAARVF AHGAATLSP NT seq 570 nt NT seq 0 atgctggggagcagagctgtaatgctgctgttgctgctgccctggacagctcagggcaga gctgtgcctgggggagcagcagccctgcctggactagtgccagcagctttcacagaagctc tgcacactggcctggagtgcacatccactagtgggacacatggatctaagagaagagga gatgaagagagctacaaatgatgttccccatatccagtgtggagatggctgtgacccccaa

ggactcagggacaacagtcagttctgcttgcaaaggatccaccagggtctgatttttat
gagaagctgctaggatcggatattttcacagggggagccttctctgctccctgatagccct
gtgggccagcttcatgcctccctactgggcctcagccaactcctgcagcctgagggtcac
cactgggagactcagcagattccaagcctcagtcccagccatggcagcgtctcctt
ctccgcttcaaaatccttcgcagcctccaggcctttgtggctgtagccgcccgggtcttt



Walk through of the Addition Public Information, besides the INN Definitional information (4)

https://www.uniprot.org/uniprot/Q9NPF7#ptm_processing





MAB, Predominant Glycans

- Characterising elements at the Specified Substance Group 1 info level:
 - 1) Structural representation of Glycans in the GlyTouCan data base Accession Number
 - 2) Display of GlyTouCan Accession Number G80858MF Fragment, IUPAC Extended Name and Calculated Monoisotopic Mass (1462.54)

https://glytoucan.org/Structures/Glycans/G80858MF

G80858MF



Accession number	G80858MF
Created Date	2014-07-27

Accession number: G80858MF Calculated Monoisotopic Mass: 1462.5444

IUPAC Extended:

beta-D-GIcpNAc-(1->2)-alpha-D-Manp-(1->3)[beta-D-GIcpNAc-(1->2)-alpha-D-Manp-(1->6)]-beta-D-Manp-(1->4)-beta-D-GIcpNAc-(1->4)[alpha-L-Fucp-(1->6)]-beta-D-GIcpNAc(1->



Signature Fields

- What is to be considered as Minimal Definitional Information (=Signature fields) for a protein enclosed in the complete Record, which is a subset (view) of the complete record.
- This information could be used to be the basic information on which a Global ID is created



Definitional Facets selected on the Left side

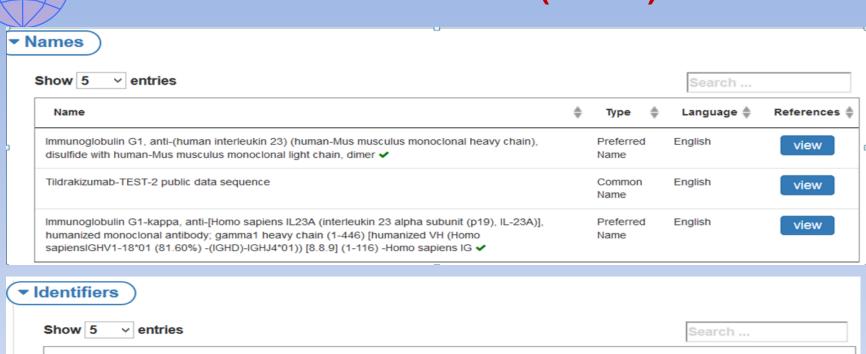
Tildrakizumab-TE	S	▼ Overview	
pending record		Substance Class	Protein
Overview		Protein Type	MONOCLONAL ANTIBODY
Names	3	Protein Sub Type	IGG1
Identifiers	5	Sequence Origin	HUMANIZED MOUSE
Subunits	4	Sequence Type	COMPLETE
	16	Record UNII	pending record
		Record Status	pending
Glycosylation	2	Record Version	9 🔻
Relationships	2	Show Definitional References ▼	
Active Moiety	2		
Modifications	3		
Characteristic Attributes	5 1	▼ Names	

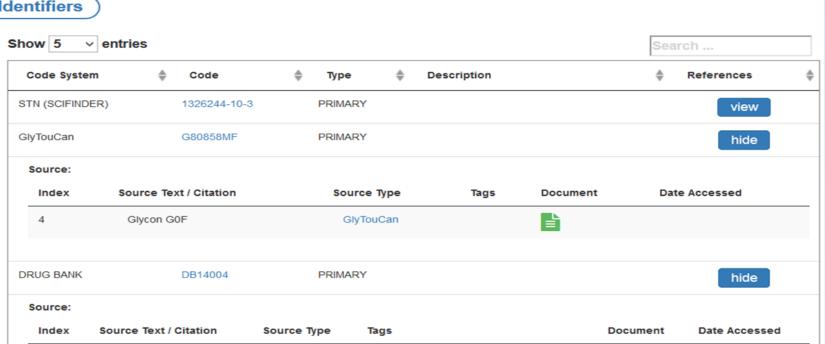
Overall Info: Subst. Class/ Protein Type/ Protein Subtype/ Sequence Origin/ Sequence Type

Facet information: Names, ID/ Codes, Subunits, Disulfide Links, Glycosylation, Modifications, Minimum relationship as Active Moiety/ Basis of Strength/
Reference Strength and Some properties like Molecular Weight (Molecular Formula)



Name & Identifiers (Codes)







Subunit Sequence and disulfide bonds

Heavy Chain (Partial representation)



Light Chain (Partial representation)

Subunit 3				
10	20	30	40	50
DIQMTQSPSS 60	LSASVGDRVT	ITCRTSENIY 80	SYLAWYQQKP	GKAPKLLIYN 100
AKTLAEGVPS	RFSGSGSGTD 120	FTLTISSLQP 130	EDFATYYCQH 140	HYGIPFTFGQ 150
GTKVEIKRTV 160	AAPSVFIFPP 170	SDEQLKSGTA 180	SVVCLLNNFY 190	P R E A K V Q W K V

7	Disulfide Links	
	From	То
	1_22	1_96
	1_143	1_199
	1_260	1_320
	1_366	1_424
	1_219	3_214

Partial representation of the Disulfide bonds



Glycosylation & Modification

Glycosylation Type and Side

ycosylation		
Glycosylation Type	MAMMALIAN	
Glycosylation Link Type		Site
N		1_296
N		2_296

Modification Type and Side

Modification Type	Location Site	Location Type	Residue Modified	Extent	Modification Name	Modification ID
AMINO_ACID_REMOVAL	[1_446] [2_446]	C-TERMINUS			LYSINE	K3Z4F929H6
N-GLYCAN FORMATION		RESIDUE_SPECIFIC	asparagine	Amount:	Glycan- G80858MF-	3a617623
				, anounce		
Degree of subs	stitutio	n is not Pub	lic	DEGREE OF SUBSTITUTION	A2G0F	
Degree of subs	stitutic	on is not Pub	lic			

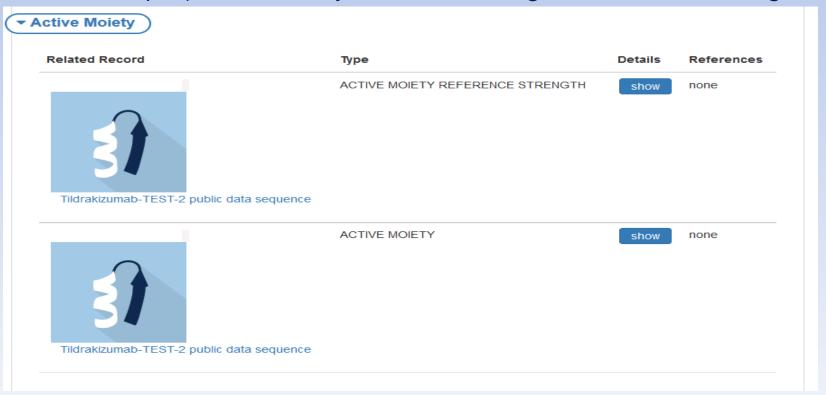


Properties & Relationships

Characteristic Attribute (Property like Mol. Weight)



Relationships (Active Moiety/ Basis or strength/ Reference strength





Other Relationships

Glycan -> Parent; Target -> Inhibitor and References

