

Subject: RE: GeneData Error for new VSIG4 sequences
Date: Thursday, November 16, 2023 at 2:39:27 PM Eastern Standard Time
From: Sausen, Christopher
To: Bard, Joel
CC: Jakobek, Ryan Andrew, Francis, Christopher Alan, Bitzas, Gabrielle, Kerns, Kelvin
Attachments: image002.png, image005.png, image006.png, image007.png

Hi Joel,

Thanks so much, that worked perfectly. Easy fix! 😊

Best,
Chris

Christopher Sausen
Senior Scientist

BioMedicine Design
Biotherapeutic Production

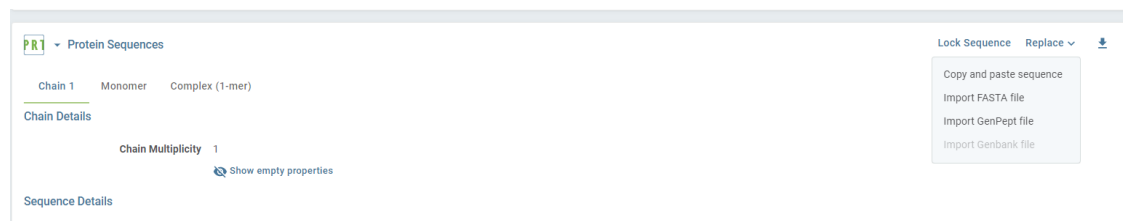
O: 1-978-247-1374 **M:** 1-218-851-7391
Christopher.sausen@pfizer.com




From: Bard, Joel <Joel.Bard@pfizer.com>
Sent: Thursday, November 16, 2023 1:31 PM
To: Sausen, Christopher <Christopher.Sausen@pfizer.com>
Cc: Jakobek, Ryan Andrew <Ryan.Jakobek@pfizer.com>; Francis, Christopher Alan <Christopher.Francis@pfizer.com>; Bitzas, Gabrielle <Gabrielle.Bitzas@pfizer.com>; Kerns, Kelvin <Kelvin.Kerns@pfizer.com>
Subject: RE: GeneData Error for new VSIG4 sequences

Hi Chris and Ryan-

The way to change complexation is to use the Replace button on the sequence tab on the TPP page and choose the Copy Paste option. That will let you edit the complexation. You won't need to do anything to the sequence, just change complexation and Finish.



 This Target Product already has 3 Actual Products attached.

Replace the Sequences for Target Product TPP-112645 (GBT-VSIG4-1044)

Sequences

Paste in Protein sequences for the Target Product.


Chain 1

Sequence

```

RPILVPEVSCTCPWKGDCNLPCTYDPLQGYTQVLVKWLVQRGSDPVTIFLRDSSGDHIQQA
KYQGRHLVSHKVPGDVSLQLSTLEMDDRRSHYTCVETWGTDPDGNQVVRDKITELRVQKHSS
KLLATATEAPTMTITYPLKATSTVKQSWDWTDMDGYLGETSAGPGASLPGGGGEPKSSDK
THTCPCPAPELLGGPSVFLFPKPKDITLMSIRTEPVTCVVVDVSHEDPEVKFNWYYVDGVE
VHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTIKAKGQPRE
PDVYTLPPSREEMTKNQVSLTCLVKGFYPSDIAVEVESNGQPENNYKTTTPVLDSDGSFFL
YSKLTVDKSRWQQGNVFCSCVMHEALHNNHYTKQSLSLSPGK
                    
```

Protein Sequence Properties

Complexation * 

Chain Multiplicity

Chain 1 * 1

Cancel
Finish

show that the complex is a 2-mer?

Thanks,
Chris

[Target Product TPP-112645 \(GBT-VSIG4-1044\) \(pfizer.com\)](#)

PRT

▼ Protein Sequences

Chain 1

Monomer

Complex (1-mer)

Details

Length

406 AA

Molecular Weight, MW

45,114.8 Da

Molar Extinction Coefficient

72,475 M⁻¹ cm⁻¹

Absorbance [280nm] of 1mg/mL

1.61

Isoelectric Point, pI

6.6

Calculation Method

pI from EMBOSS Pepstats version 6.5.7.0a. Mass calculated in GDBxT is oxidized, deglycosylated.

Sequence

1

RP

ILEVPESV

TCPWKGDCNL

PCTYDPLQGY

TQVLVKWL

RGSDPVTIFL

RDSSGDHIQQ

AKYQGR

LHVS

HKVP

GDVSL

81

LST

LEMDDRS

HYTCEVTWQT

PDGNQVVRDK

ITELRVQKHS

SKLLATATEA

PTTMTYPLKA

TSTVKQSWDW

TTDMDGYLGI

161

TSAG

PGASLP

GGGGEPKSSD

KTHTCPPCPA

PELLGGPSVF

LFPPKPKDTL

MISRTPEVTC

VVVDVSHEDP

EVKFNWYVDI

241

VEVH

NAKTKP

REEQYNSTYR

VVSVLTVLHQ

DWLNGKEYKC

KVSNKALPAP

IEKTISKAKG

QPREPQVYTL

PPSREEMTKI

321

QVSL

TCLVKG

FYP

SDIAVEW

ESNGQPENNY

KTPPVLDSD

GSFFLYSKLT

VDKSRWQQGN

VFSCSV

MHEA

LHNHYTQKSI

401

SL

SPGK

Show empty properties

The complex sequence has been generated in-silico by concatenating the monomer sequence 1 times (complexation number).

Christopher Sausen
Senior Scientist

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