



#### **JEDI Program/JDI01**

#### **Compound Screening for SARS-CoV-2 Proteins Using MST/Dianthus**

nsp12

nanoDSF / MST/ TRIC measurements

May 27, 2021



#### **Status**



#### nanoDSF:

- nsp12 shows good thermal stability (Tm of ~47°C).
- Reducing agents and detergents have no significant impact on protein thermal stability, and the protein tolerates up to 3% DMSO.
- However, none of the tool cpds (GS-443902 trisodium = active Remdesivir metabolite), Ribavirin (ICN-1229) and Favipiravir (MCE)) nor ATP show significant effects on protein thermal stability.

#### TRIC (Dianthus):

- Nsp12 was successfully labelled with RED-Tris-NTA dye.
- However, none of the tool cpds (GS-443902 trisodium = active Remdesivir metabolite), Ribavirin (ICN-1229) and Favipiravir (MCE)) show binding to RED-Tris-NTA labelled nsp12.

#### Labelled MST:

- For comparison, experiments were also run on the NT.115 instrument and are in line with TRIC and nanoDSF results.
- ➤ Nsp12 alone is insufficient for binding of ATP or tool compounds. Thus, a TRIC/MST assay cannot be established with nsp12.

### nanoDSF

nsp12 (DVT1, PC13929-1)

### nanoDSF Assay Conditions



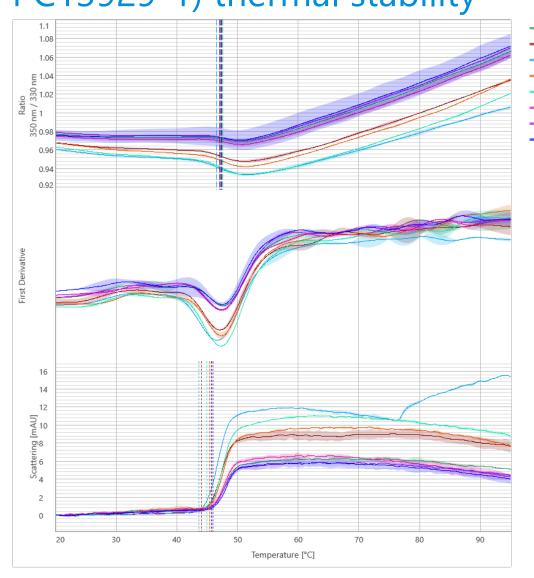
Protein	0.1 mg/ml nsp12 (DVT1, PC13929-1)
Assay Buffer	8 Buffers (see table)
Instrument	Prometheus NT.48
Capillary type	nanoDSF Standard Capillaries
Measurement parameters	LED Power: 50 % Temperature ramp: 2°C/min

	Conditions
Buffer 1	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2</sub>
Buffer 2	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2</sub> , <b>0.005 % Tween20</b>
Buffer 3	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2</sub> , <b>0.05 % Tween20</b>
Buffer 4	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2</sub> , <b>0.05 % Pluronic</b>
Buffer 5	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2</sub> , <b>0.1 % Pluronic</b>
Buffer 6	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2</sub> , <b>1 mM TCEP</b>
Buffer 7	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2</sub> , <b>2.5 mM DTT</b>
Buffer 8	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2</sub> , <b>2.5 mM GSH</b>



# Effect of reducing agents and detergents on nsp12 (DVT1, PC13929-1) thermal stability





Condition	Ø T <sub>m</sub> [°C] <sup>1</sup>	s.d. [°C]	Ø T <sub>agg</sub> [°C] <sup>1</sup>	s.d. [°C]	Analysis mode
Buffer 1	47.6	0.2	45.8	0.1	ratio
Buffer 2	47.1	0.3	44.1	0.1	ratio
Buffer 3	46.6	0.0	43.7	0.0	ratio
Buffer 4	47.2	0.1	45.4	0.1	ratio
Buffer 5	47.2	0.1	45.0	0.1	ratio
Buffer 6	47.3	0.2	45.6	0.1	ratio
Buffer 7	47.2	0.1	45.3	0.6	ratio
Buffer 8	47.3	0.1	45.9	0.2	ratio

<sup>&</sup>lt;sup>1</sup> determined in duplicate

 No significant impact of detergents or reducing agents on protein thermal stability was observed



### nanoDSF Assay Conditions

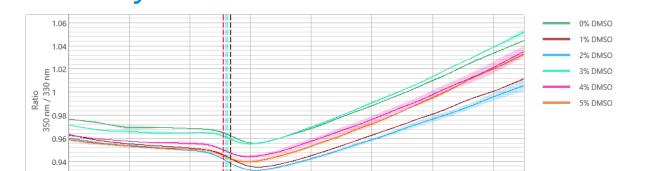


Protein	0.1 mg/ml nsp12 (DVT1, PC13929-1)
Assay Buffer	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl2,2.5 mM DTT, 0.005 % Tween20 DMSO: 0 - 5 %
Instrument	Prometheus NT.48
Capillary type	nanoDSF Standard Capillaries
Measurement parameters	LED Power: 50 % Temperature ramp: 2°C/min



# Impact of DMSO on nsp12 (DVT1, PC13929-1) thermal stability





Temperature [°C]

DMSO	T <sub>m</sub> [°C]	s.d. [°C]	ΔT <sub>m</sub> [°C] <sup>2</sup>	Ø T <sub>agg</sub> [°C] <sup>1</sup>	s.d. [°C]	Analysis mode
-	46.8	0.1	-	43.6	1.6	ratio
1%	46.6	0.2	- 0.2	43.6	0.3	ratio
2%	46.0	0.1	- 0.8	39.6	4.6	ratio
3%	46.3	0.1	- 0.5	43.7	0.2	ratio
4%	45.6	0.2	- 1.2	42.9	0.1	ratio
5%	45.5	0.2	- 1.3	42.2	0.4	ratio

- Increasing DMSO concentrations destabilize the protein.
- A concentration of 2.5 % was selected for further measurements.



95

### nanoDSF Assay Conditions



Protein	0.1 mg/ml nsp12 (DVT1, PC13929-1)
Assay Buffer	20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl2, 2.5 mM DTT, 0.005 % Tween20 DMSO: 2.5 %
compounds	ATP at 100 $\mu$ M and 500 $\mu$ M GS-443902 trisodium (JDI-880) at 10 $\mu$ M and 100 $\mu$ M Ribavirin (ICN-1229) (JDI-879) at 10 $\mu$ M and 100 $\mu$ M Favipiravir (MCE) (JDI-881) at 10 $\mu$ M and 100 $\mu$ M
Instrument	Prometheus NT.48
Capillary type	nanoDSF Standard Capillaries
Measurement parameters	LED Power: 50 % Temperature ramp: 2°C/min

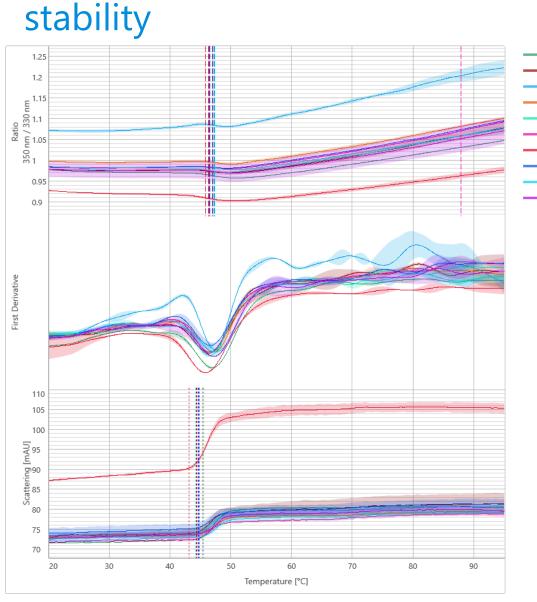


### Effect of compounds on nsp12 (DVT1, PC13929-1) thermal

500 µM ATP

10 μM Ribavirin (I 100 μM Favipiravi 10 μM Favipiravir





)2 trisodium	
2 trisodium	
(ICN-1229)	
CN-1229)	
r (MCE)	
(MCE)	
	100
	1 ( )( )

Titrant	Ø T <sub>m</sub> [°C] <sup>1</sup>	s.d. [°C]	ΔT <sub>m</sub> [°C] <sup>2</sup>	Ø T <sub>agg</sub> [°C]	s.d. [°C]	Analysis mode
Protein alone	46.9	0.1	-	44.7	0.2	ratio
2.5 % DMSO	46.4	0.1	- 0.5	44.3	0.5	ratio
500 μM ATP	47.3	0.1	0.4	44.8	0.5	ratio
100 μM ATP	47.1	0.2	0.2	45.3	0.3	ratio
100 μM GS-443902 trisodium	46.5	0.2	- 0.4	44.3	0.1	ratio
10 μM GS-443902 trisodium	46.5	0.2	- 0.4	44.5	0.0	ratio
100 μM Ribavirin (ICN-1229)	45.8	0.1	- 1.1	43.1	0.0	ratio
10 μM Ribavirin (ICN-1229)	46.6	0.4	- 0.3	44.7	0.1	ratio
100 μM Favipiravir (MCE)	47.4	0.0	0.5	45.6	0.2	ratio
10 μM Favipiravir (MCE)	47.1	0.1	0.2	44.7	0.5	ratio

<sup>&</sup>lt;sup>1</sup> determined in duplicate

 Neither ATP nor any of the tool compounds has a significant impact on protein thermal stability.



<sup>&</sup>lt;sup>2</sup> referenced to

#### nanoDSF Summary: cpds effects on nsp12 (DVT1, PC13929-1)



Effect of 3 compounds and ATP on the thermal stability of nsp12 (DVT1, PC13929-1) analyzed by nanoDSF

Target name	Titrant / Condition	Ø T <sub>m</sub> [°C]	s.d. [°C]	ΔT <sub>m</sub> [°C]	Ø T <sub>agg</sub> [°C]	s.d. [°C]	Analysis mode
	Protein alone	46.9	0.1	-	44.7	0.2	ratio
	2.5 % DMSO	46.4	0.1	- 0.5	44.3	0.5	ratio
	500 μM ATP	47.3	0.1	0.4	44.8	0.5	ratio
	100 μM ATP	47.1	0.2	0.2	45.3	0.3	ratio
nsp12	100 μM GS-443902 trisodium	46.5	0.2	- 0.4	44.3	0.1	ratio
	10 μM GS-443902 trisodium	46.5	0.2	- 0.4	44.5	0.0	ratio
	100 μM Ribavirin (ICN-1229)	45.8	0.1	- 1.1	43.1	0.0	ratio
	10 μM Ribavirin (ICN-1229)	46.6	0.4	- 0.3	44.7	0.1	ratio
	100 μM Favipiravir (MCE)	47.4	0.0	0.5	45.6	0.2	ratio
	10 μM Favipiravir (MCE)	47.1	0.1	0.2	44.7	0.5	ratio



## TRIC (Dianthus)

nsp12 (DVT1, PC13929-1)

### TRIC Assay Conditions

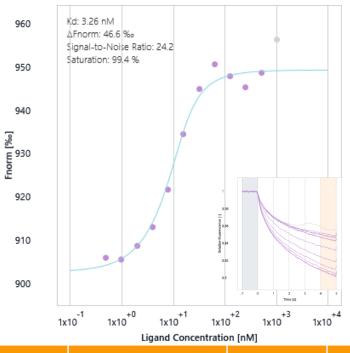


Fluor. Molecule	25 nM RED-tris-NTA 1 <sup>st</sup> gen. 25 nM RED-tris-NTA 2 <sup>nd</sup> gen.									
Fluorophore	RED-tris-NTA 1 <sup>st</sup> and 2 <sup>nd</sup> gen.									
Instrument	Dianthus NT.23PicoDuo									
Measurement parameter	LED Power: 11 % (nano detector)  TRIC settings: 1 - 5 - 1 (s) (initial fluorescence – MST on Singlicates	time – back-diffusion)								
Assay buffer	Buffer 1: 20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2,</sub> 2.5 mM DTT, 0.05% Tween20 Buffer 2: 20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2,</sub> 2.5 mM DTT, 0.5 mM ATP, 0.05% Tween20									
Titrant	nsp12 (DVT1, PC13929-1)	1000 – 0.49 nM (12 conc.)								

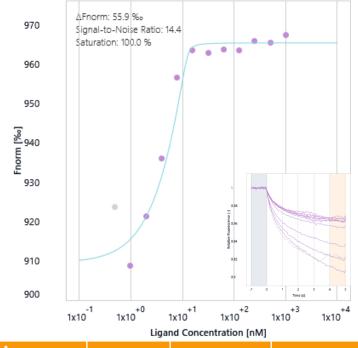


## RED-tris-NTA 1<sup>st</sup> and 2<sup>nd</sup> gen. vs. nsp12 (DVT1, PC13929-1) at Buffer 1





RED-tris-NTA 1st gen.



RED-tris-NTA 2<sup>nd</sup> gen.

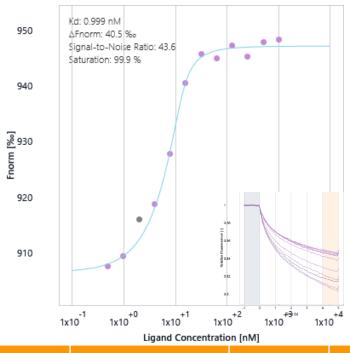
Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	Lower confidence [M]	Upper confidence [M]	ΔFnorm [‰]	Signal / Noise	TRIC On [s]	Comment
RED-tris-NTA 1st gen.	RED-tris-NTA 1st gen.	nsp12	3.3E-09	1.8E-09	5.9E-09	46.6	24.2	5	Buffer 1
RED-tris-NTA 2 <sup>nd</sup> gen.	RED-tris-NTA 2 <sup>nd</sup> gen.	nsp12	1.0E-10	1.1E-14	8.8E-07	55.9	14.4	5	Buffer 1

- RED-tris-NTA 1<sup>st</sup> gen. binds nsp12 with a determined K<sub>D</sub> of 3.3 nM.
- RED-tris-NTA 2<sup>nd</sup> gen. binds nsp12 with a determined K<sub>D</sub> of 0.1 nM.
- Both dyes can be used to non-covalently label nsp12.

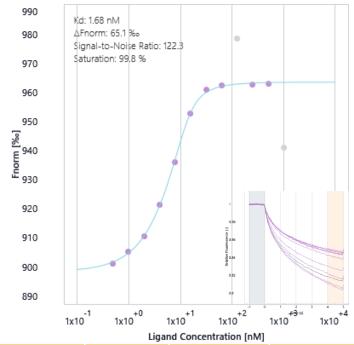


## RED-tris-NTA 1<sup>st</sup> and 2<sup>nd</sup> gen. vs. nsp12 (DVT1, PC13929-1) at Buffer 2





RED-tris-NTA 1st gen.



RED-tris-NTA 2<sup>nd</sup> gen.

Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	Lower confidence [M]	Upper confidence [M]	ΔFnorm [‰]	Signal / Noise	TRIC On [s]	Comment
RED-tris-NTA 1st gen.	RED-tris-NTA 1st gen.	nsp12	1.0E-10	-	-	40.5	43.6	5	Buffer 2
RED-tris-NTA 2 <sup>nd</sup> gen.	RED-tris-NTA 2 <sup>nd</sup> gen.	nsp12	1.7E-09	1.4E-09	2.0E-09	65.1	122.3	5	Buffer 2

- RED-tris-NTA 1<sup>st</sup> gen. binds nsp12 with a determined K<sub>D</sub> of 1.0 nM.
- RED-tris-NTA  $2^{nd}$  gen. binds nsp12 with a determined  $K_D$  of 1.7 nM.
- Both dyes can be used to non-covalently label nsp12.



### TRIC Assay Conditions

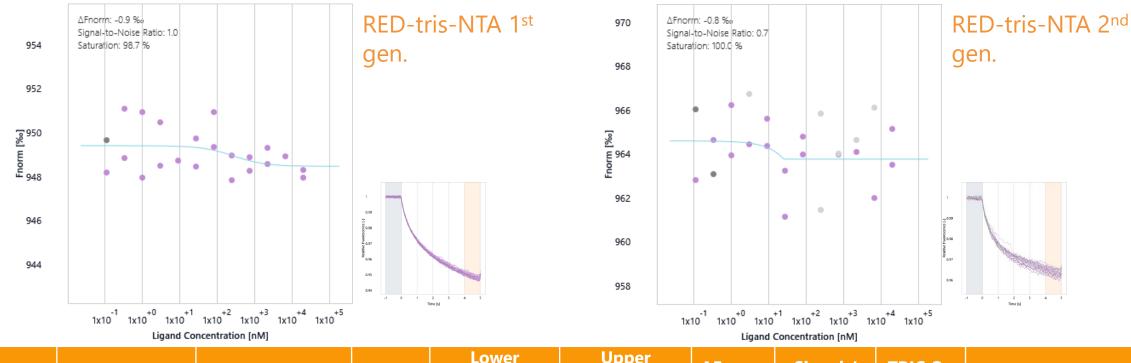


Fluor. Molecule	25 nM nsp12 (DVT1, PC13929-1)							
Fluorophore	RED-tris-NTA 1 <sup>st</sup> and 2 <sup>nd</sup> gen.							
Labelling conditions	25 nM protein / 12.5 nM dye Incubation time: 30 min Centrifugation: 10 min at 15000g							
Instrument	Dianthus NT.23PicoDuo							
Measurement parameter	LED Power: 23 % (nano detector)  TRIC settings: 1 - 5 - 1 (s) (initial fluorescence – MST on time – back-diffusion)  Duplicates							
Assay buffer	Buffer 1: 20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2,</sub> 2.5 mM DTT, 0.05% Tween20 Buffer 2: 20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2,</sub> 2.5 mM DTT, 0.5 mM ATP, 0.05% Tween20							
Titrant	GS-443902 trisodium (active DDI-880 $\mu$ M = 0.11 nM (12 conc.) $\mu$ M = 0.11 nM (12 conc.)							



## RED-tris-NTA 1<sup>st</sup> and 2<sup>nd</sup> gen. vs. nsp12 (DVT1, PC13929-1) in Buffer 1





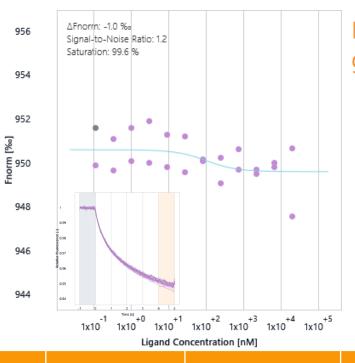
Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	Lower confidence [M]	Upper confidence [M]	ΔFnorm [‰]	Signal / Noise	TRIC On [s]	Comment
RED-tris-NTA 1st gen.	nsp12	GS-443902 trisodium	-	-	-	-	-	5	Buffer 1
RED-tris-NTA 2 <sup>nd</sup> gen.	nsp12	GS-443902 trisodium	-	-	-	-	-	5	Buffer 1

• RED-tris-NTA 1st and 2nd gen. labelled nsp12 does not bind GS-443902 trisodium in buffer 1.

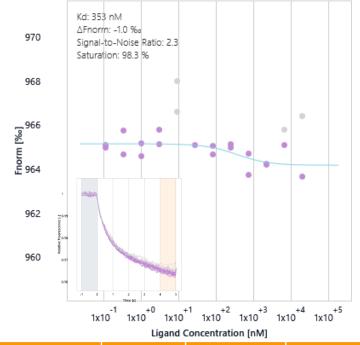


## RED-tris-NTA 1<sup>st</sup> and 2<sup>nd</sup> gen. vs. nsp12 (DVT1, PC13929-1) in Buffer 2





RED-tris-NTA 1st gen.



RED-tris-NTA 2<sup>nd</sup> gen.

Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	Lower confidence [M]	Upper confidence [M]	ΔFnorm [‰]	Signal / Noise	TRIC On [s]	Comment
RED-tris-NTA 1st gen.	nsp12	GS-443902 trisodium	-	-	-	-	=	5	Buffer 2
RED-tris-NTA 2 <sup>nd</sup> gen.	nsp12	GS-443902 trisodium	-	-	-	-	-	5	Buffer 2

• RED-tris-NTA 1st and 2nd gen. labelled nsp12 does not bind GS-443902 trisodium in buffer 2.



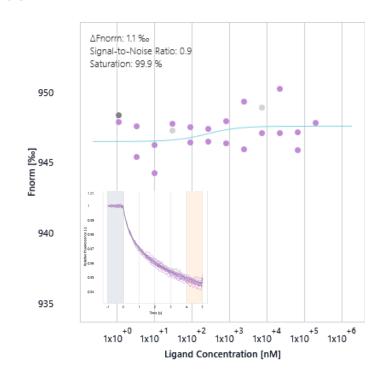
### TRIC Assay Conditions



Fluor. Molecule	25 nM nsp12 (DVT1, PC13929	9-1)							
Fluorophore	RED-tris-NTA 1 <sup>st</sup>								
Labelling conditions	25 nM protein / 12.5 nM dye Incubation time: 30 min Centrifugation: 10 min at 150	00g							
Instrument	Pianthus NT.23PicoDuo								
Measurement parameter									
Assay buffer	Buffer 1: 20 mM HEPES pH 7. DMSO: 2.5%	Buffer 1: 20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2,</sub> 2.5 mM DTT, 0.05% Tween20 DMSO: 2.5%							
Titrant	GS-443902 trisodium Ribavirin (ICN-1229) Favipiravir (MCE)	JDI-880 JDI-879 JDI-881	200 μM – 1.13 nM (12 conc.)						

## RED-tris-NTA 1<sup>st</sup> labelled nsp12 (DVT1, PC13929-1) vs. GS-443902 trisodium





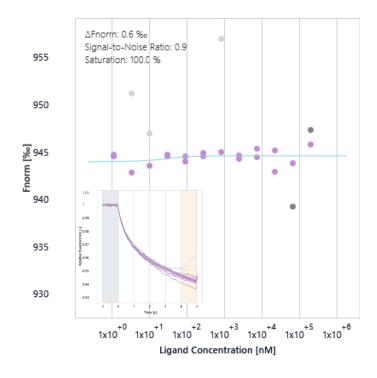
Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	Lower confidence [M]	Upper confidence [M]	ΔFnorm [‰]	Signal / Noise	TRIC On [s]	Comment
RED-tris-NTA 1st gen.	nsp12	GS-443902 trisodium	-	-	-	-	-	5	

RED-tris-NTA labelled nsp12 does not bind GS-443902 trisodium.



## RED-tris-NTA 1<sup>st</sup> labelled nsp12 (DVT1, PC13929-1) vs. Ribavirin (ICN-1229)





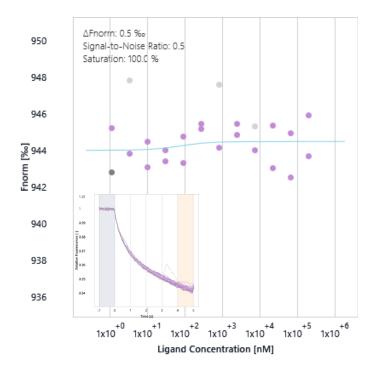
Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	Lower confidence [M]	Upper confidence [M]	ΔFnorm [‰]	Signal / Noise	TRIC On [s]	Comment
RED-tris-NTA 1st gen.	nsp12	Ribavirin (ICN-1229)	-	-	-	-	=	5	

• RED-tris-NTA labelled nsp12 does not bind Ribavirin (ICN-1229).



# RED-tris-NTA 1<sup>st</sup> labelled nsp12 (DVT1, PC13929-1) vs. Favipiravir (MCE)





Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	Lower confidence [M]	Upper confidence [M]	ΔFnorm [‰]	Signal / Noise	TRIC On [s]	Comment
RED-tris-NTA 1st gen.	nsp12	Favipiravir (MCE)	-	-	-	-	-	5	

RED-tris-NTA labelled nsp12 does not bind Favipiravir (MCE).



### Labelled MST

nsp12 (DVT1, PC13929-1)

### MST labelled assay conditions



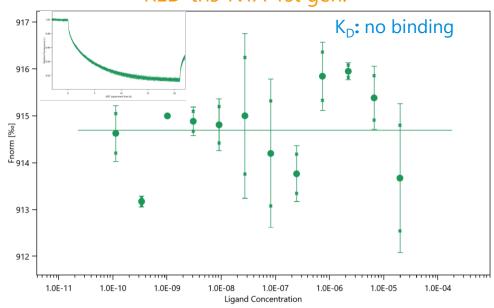
Fluor. Molecule	25 nM nsp12 (DVT1, PC13929-1)								
Fluorophore	RED-tris-NTA 1st and 2nd gen.								
	25 nM protein / 12.5 nM dye								
<b>Labelling conditions</b>	Incubation time: 30 min								
	Centrifugation: 10 min at 15000g								
Instrument	Monolith NT.115 (03)								
Capillary type	Monolith™ NT.115 Series MST Premium Coated Capillaries								
	LED Power: 80 %								
Measurement	MST Power: 40 %								
parameter	MST settings: 3 – 20 – 1 (s) (initial fluorescence – MST on time – back-diffusion)								
	Duplicate								
Access buffer	Buffer 1: 20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2,</sub> 2.5 mM DTT, 0.05% Tween20								
Assay buffer	Buffer 2: 20 mM HEPES pH 7.5, 150 mM NaCl, 1 mM MgCl <sub>2,</sub> 2.5 mM DTT, 0.5 mM ATP, 0.05% Tween20								
Titrant	GS-443902 trisodium JDI-880 20 μM – 0.11 nM (12 conc.)								



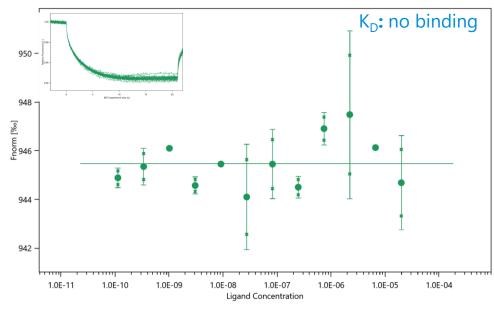
#### RED-tris-NTA 1<sup>st</sup> and 2<sup>nd</sup> gen. labelled nsp12 (DVT1, PC13929-1) vs. GS-443902 trisodium in buffer 1







#### RED-tris-NTA 2nd gen.



Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	K <sub>D</sub> Confidence [M]	ΔFnorm [‰]	Signal / Noise	MST on [s]	Comment
RED-tris-NTA 1 <sup>st</sup> gen.	nsp12	GS-443902 trisodium	-	-	-	-	20	Buffer 1
RED-tris-NTA 2 <sup>nd</sup> gen.	nsp12	GS-443902 trisodium	-	-	-	-	20	Buffer 1

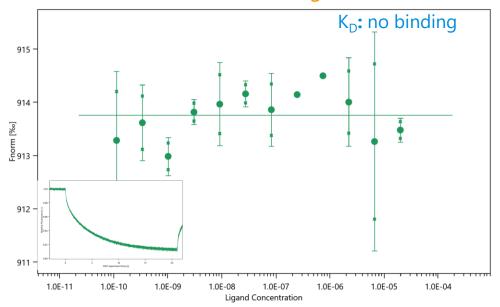
- RED-tris-NTA 1st gen. labeled nsp12 does not bind GS-443902 trisodium.
- RED-tris-NTA 2<sup>nd</sup> gen. labeled nsp12 does not bind GS-443902 trisodium.



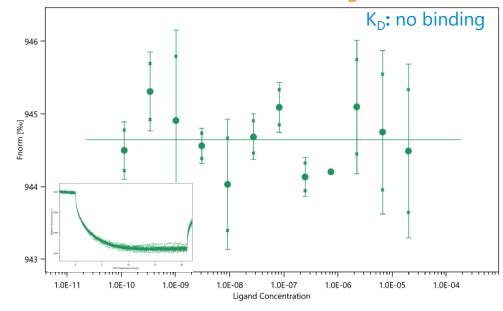
## RED-tris-NTA 1<sup>st</sup> and 2<sup>nd</sup> gen. labelled nsp12 (DVT1, PC13929-1) vs. GS-443902 trisodium in buffer 2



#### RED-tris-NTA 1st gen.



#### RED-tris-NTA 2nd gen.



Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	K <sub>D</sub> Confidence [M]	ΔFnorm [‰]	Signal / Noise	MST on [s]	Comment
RED-tris-NTA 1st gen.	nsp12	GS-443902 trisodium	-	-	-	-	20	Buffer 2
RED-tris-NTA 2 <sup>nd</sup> gen.	nsp12	GS-443902 trisodium	-	-	-	-	20	Buffer 2

- RED-tris-NTA 1<sup>st</sup> gen. labelled nsp12 does not bind GS-443902 trisodium.
- RED-tris-NTA 2<sup>nd</sup> gen. labelled nsp12 does not bind GS-443902 trisodium.



### MST labelled assay conditions

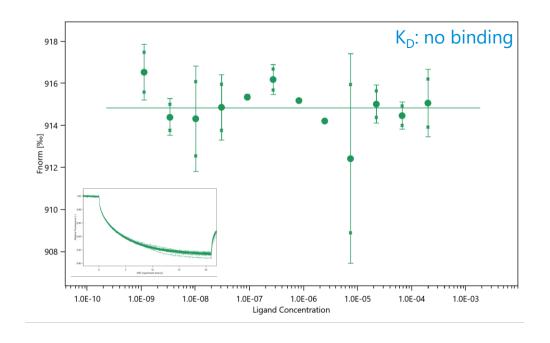


Fluor. Molecule	25 nM nsp12 (DVT1, PC13	929-1)							
Fluorophore	RED-tris-NTA 1st and 2nd g	jen.							
	25 nM protein / 12.5 nM c	dye							
<b>Labelling conditions</b>	Incubation time: 30 min								
	Centrifugation: 10 min at	Centrifugation: 10 min at 15000g							
Instrument	Monolith NT.115 (03)	Monolith NT.115 (03)							
Capillary type	Monolith™ NT.115 Series MST Premium Coated Capillaries								
	LED Power: 100 %								
Measurement	MST Power: 40 %								
parameter	MST settings: 3 – 20 – 1 (s	(initial fluorescence – I	MST on time – back-diffusion)						
	Duplicate								
Access builton	Buffer 1: 20 mM HEPES ph	1 7.5, 150 mM NaCl, 1 mM I	MgCl <sub>2,</sub> 2.5 mM DTT, 0.05% Tween20						
Assay buffer	DMSO: 2.5%								
	GS-443902 trisodium	JDI-880							
Titrant	Ribavirin (ICN-1229)	JDI-879	200 μM – 1.13 nM (12 conc.)						
	Favipiravir (MCE)	JDI-881							



## [RED-tris-NTA 1<sup>st</sup> labelled nsp12 (DVT1, PC13929-1) vs. GS-443902 trisodium





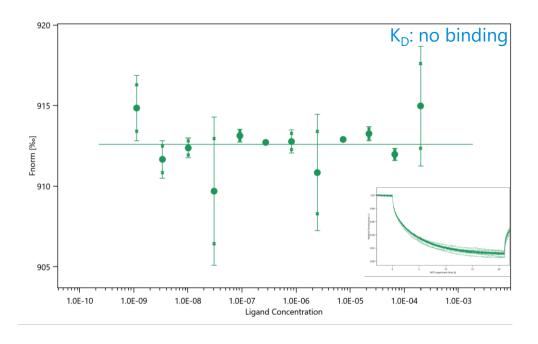
Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	K <sub>D</sub> Confidence [M]	ΔFnorm [‰]	Signal / Noise	MST on [s]	Comment
RED-tris-NTA 1st gen.	nsp12	GS-443902 trisodium	-	-	-	-	20	

• RED-tris-NTA labelled nsp12 does not bind GS-443902 trisodium.



## [RED-tris-NTA 1<sup>st</sup> labelled nsp12 (DVT1, PC13929-1) vs. Ribavirin (ICN-1229)





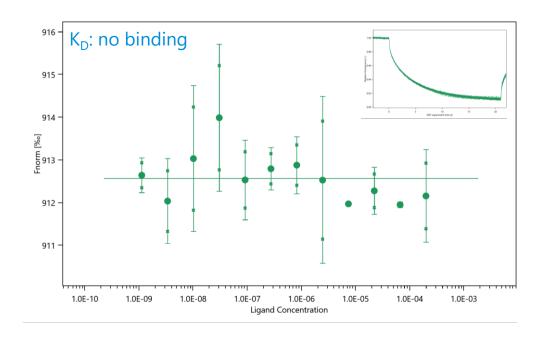
Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	K <sub>D</sub> Confidence [M]	ΔFnorm [‰]	Signal / Noise	MST on [s]	Comment
RED-tris-NTA 1st gen.	nsp12	Ribavirin (ICN-1229)	-	-	-	-	20	

• RED-tris-NTA labelled nsp12 does not bind Ribavirin (ICN-1229).



# RED-tris-NTA 1<sup>st</sup> labelled nsp12 (DVT1, PC13929-1) vs. Favipiravir (MCE)





Fluorophore	Fluor. Molecule	Titrant	K <sub>D</sub> [M]	K <sub>D</sub> Confidence [M]	ΔFnorm [‰]	Signal / Noise	MST on [s]	Comment
RED-tris-NTA 1st gen.	nsp12	Favipiravir (MCE)	-	-	=	-	20	

• RED-tris-NTA labelled nsp12 does not bind Favipiravir (MCE).



#### Next steps



• As neither nsp12 nor the RdRp complex allow biophysical characterization of compound interactions using TRIC (Dianthus), we recommend to stop the project







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