

Relationship of binding-site occupancy, transthyretin stabilisation, and disease modification in tafamidis treated transthyretin amyloid cardiomyopathy patients

Tess, D. A., Maurer, T. S., Li, Z., Bulawa, C., Fleming, J., & Moody, A. T. (2022). Relationship of binding-site occupancy, transthyretin stabilization and disease modification in patients with tafamidis-treated transthyretin amyloid cardiomyopathy. *Amyloid*, 1-12.

<https://doi.org/10.1080/13506129.2022.2145876>

Tafamidis* is a selective stabilizer of TTR

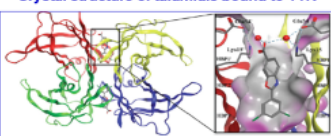
Tafamidis is a selective stabilizer of TTR and prevents the tetramer from further dissociating into monomers, misfolding, and causing amyloid fibril formation and accumulation in tissues and organs^{1,5}

Tafamidis is indicated for the treatment of wild-type and hereditary ATTR-CM in adults to reduce cardiovascular mortality and cardiovascular-related hospitalization⁴

The Phase 3 Tafamidis in Transthyretin Cardiomyopathy Clinical Trial (ATTR-CT) demonstrated the efficacy and safety of Tafamidis for patients with ATTR-CM¹

- Tafamidis binds in one or both of the two thyroxine binding pockets at the TTR dimer-dimer interface
- Tafamidis inhibits tetramer dissociation, resulting in stabilization and increased abundance of the tetramer²

Crystal structure of tafamidis bound to TTR



Left: 3D ribbon diagram depiction of the TTR tetramer with tafamidis bound. The four TTR monomers are individually colored. Right: Magnified image of tafamidis bound in one of the thyroxine T4 binding sites.
Image reproduced from Bulawa CE, et al. *Proc Natl Acad Sci USA* 2012²

*The FDA approved drug is other tafamidis meglumine (TTR 28 mg tafamidis meglumine capsules) only once daily (TR) tafamidis free acid (Taf) (one 60 mg tafamidis free acid capsule) only once daily

ATTR-CT: Tafamidis in Transthyretin Cardiomyopathy Clinical Trial. QD, every day; TTR, transthyretin.
1. Maurer TS, et al. *N Engl J Med* 2021;385(11):1057-67. 2. Bulawa CE, et al. *Proc Natl Acad Sci USA* 2012;109(4):1020-24. 3. Bulawa CE, et al. *Proc Natl Acad Sci USA* 2012;109(4):1020-24. 4. VINDICATE: VINDICATE
Prescribing Information. New York, NY: Pfizer Inc.; 2021. 5. Li Z, et al. *Can J Cardiol* 2021;37(1):1-11

Study Design

In this study, TTR stabilization was defined as the degree to which the tetramer dissociation rate decreased in human plasma following the 20 mg and the approved 80 mg once daily (QD) doses of tafamidis meglumine.¹

This definition was chosen because the change in TTR dissociation rate under therapeutic and physiologic conditions is the most clinically relevant measure of pharmacologic activity for this class.²

Methods for characterizing the binding of tafamidis meglumine to both TTR tetramer and human albumin¹

- Isothermal titration calorimetry (ITC)
- Subunit fraction exchange
- Albumin binding (equilibrium dialysis methods)

Clinical data from the Phase 3 ATTR-CT study was used in this analysis¹

TTR concentrations over time were simultaneously characterized as a function of inter-individual variability in TTR production rate, and also as a function of tafamidis meglumine concentration, to develop population TTR binding and concentration-time models

Data obtained from ITC, subunit fraction exchange, and albumin binding experiments were simultaneously modeled to estimate a common set of tafamidis meglumine binding dissociation constants to TTR (K_{d1} and K_{d2}) and albumin (K_{dA})

Population disease progression response-time models were developed using three disease progression measures:

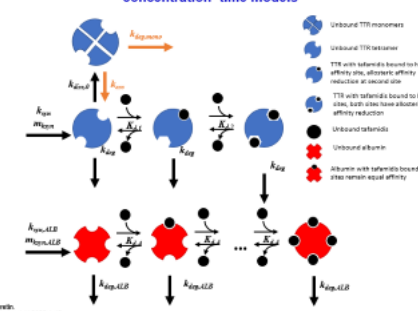
- N-terminal pro-B natriuretic peptide (NT-proBNP) plasma levels
- Kansas City Cardiomyopathy Questionnaire Overall Score (KCCQ-OS) score
- 6-minute walk test (6MWT) distance

6MWT, 6-minute walk test; ATTR-CT, Tafamidis in Transthyretin Cardiomyopathy Clinical Trial; KCCQ-OS, Kansas City Cardiomyopathy Questionnaire Overall Score; NT-proBNP, N-terminal pro-B type natriuretic peptide; QD, once daily; TTR, transthyretin.
1. Tess DA, et al. *Amyloid* 2022;1-12. 2. Bulawa CE, et al. *Am J Pathol* 2012;121(1):24-4

Results: Characterization of tafamidis meglumine binding to TTR and albumin in plasma

The final model structure used to characterize the relationship between tafamidis meglumine and TTR concentrations is shown in the figure

Schematic of plasma TTR and albumin concentration-time models



Use of these values provided accurate predictions of the concentration-dependent unbound fraction of tafamidis meglumine in human plasma and confidence in the TTR occupancies inferred by these parameters

Pfizer
Rare Diseases

TTR, transthyretin;
Tess DA, et al. *Amyloid* 2022;1-12

Results: Relationship between TTR occupancy, TTR stabilization, and clinical measures related to disease progression

Tafamidis meglumine fractional TTR occupancy

- Population average plasma tafamidis meglumine concentrations associated with 20 and 80 mg QD doses were 8.8 and 35 μ M, respectively
- As such, tafamidis meglumine doses are estimated to achieve population average site occupancy as follows:

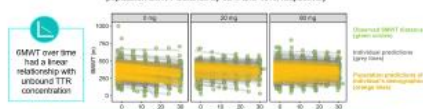
TTR occupancy		
	Single site	Double site
20 mg	73%	1%
80 mg	94%	9%

- Modeling, consistent with in vitro results and literature reports, found that the dissociation rate of TTR in the presence of singly bound tafamidis was negligible
- With a TTR single site occupancy of 94%, this results in tafamidis 80 mg achieving TTR stabilization that is close to the estimated maximum stabilization that would be achieved with 100% TTR occupancy

TTR, transthyretin; QD, every day

6MWT distance-time model

Tafamidis meglumine doses of 20 mg and 80 mg were estimated to inhibit the decline in average population 6MWT distance by 36% and 49%, respectively

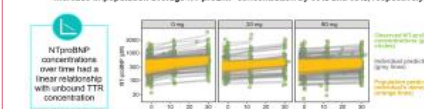


Maximal inhibition (at 100% TTR occupancy) was estimated to be only slightly greater (94%) than the average inhibition achieved with tafamidis meglumine at the 80 mg dose

6MWT, 6-minute walk test; QD, every day; TTR, transthyretin.
Tess DA, et al. *Amyloid* 2022;1-12

NT-proBNP concentration-time model

Tafamidis meglumine doses of 20 mg and 80 mg were estimated to inhibit the increase in population average NT-proBNP concentration by 39% and 53%, respectively

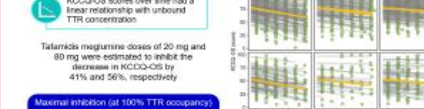


Maximal inhibition (at 100% TTR occupancy) was estimated to be only slightly greater (59%) than the average inhibition achieved with tafamidis meglumine at the 80 mg dose

NT-proBNP, N-terminal pro-B natriuretic peptide; QD, every day; TTR, transthyretin.
Tess DA, et al. *Amyloid* 2022;1-12

KCCQ-OS score-time model

Tafamidis meglumine doses of 20 mg and 80 mg were estimated to inhibit the decrease in KCCQ-OS score by 41% and 56%, respectively



Maximal inhibition (at 100% TTR occupancy) was estimated to be only slightly greater (61%) than the average inhibition achieved with tafamidis meglumine at the 80 mg dose

KCCQ-OS, Kansas City Cardiomyopathy Questionnaire Overall Score; QD, every day; TTR, transthyretin.
Tess DA, et al. *Amyloid* 2022;1-12

Conclusions

These results indicate that singly bound tafamidis provides near-complete TTR stabilization

The model demonstrates that the clinically approved dose of 80 mg tafamidis meglumine achieves a near-maximum (>90%) effect on both TTR stabilization and disease-relevant measures

These findings support the value of TTR stabilization as a clinically effective treatment for ATTR-CM

Pfizer
Rare Diseases

ATTR-CM, transthyretin amyloid cardiomyopathy; TTR, transthyretin;
Tess DA, et al. *Amyloid* 2022;1-12

Tafamidis* is a selective stabilizer of TTR



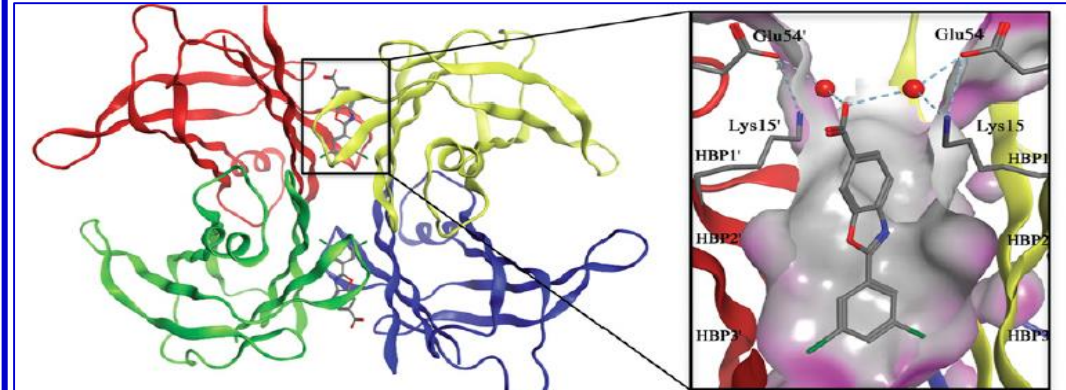
Tafamidis is a selective stabilizer of TTR and prevents the tetramer from further dissociating into monomers, misfolding, and causing amyloid fibril formation and accumulation in tissues and organs^{1,5}

Tafamidis is indicated for the treatment of wild-type and hereditary ATTR-CM in adults to reduce cardiovascular mortality and cardiovascular-related hospitalization⁴

The Phase 3 Tafamidis in Transthyretin Cardiomyopathy Clinical Trial (ATTR-ACT) demonstrated the efficacy and safety of Tafamidis for patients with ATTR-CM¹

- Tafamidis binds in one or both of the two thyroxine binding pockets at the TTR dimer-dimer interface
- **Tafamidis inhibits tetramer dissociation, resulting in stabilization and increased abundance of the tetramer²**

Crystal structure of tafamidis bound to TTR



Left: 3D ribbon diagram depiction of the TTR tetramer with tafamidis bound. The four TTR monomers are individually colored. Right: Magnified image of tafamidis bound in one of the thyroxine (T4) binding sites.

Image reproduced from Bulawa CE, et al. *Proc Natl Acad Sci USA* 2012³

*The FDA approved dosing is either tafamidis meglumine 80mg (four 20 mg tafamidis meglumine capsules) orally once daily OR tafamidis free acid 61mg (one 61mg tafamidis free acid capsule) orally once daily.

ATTR-ACT, Tafamidis in Transthyretin Cardiomyopathy Clinical Trial; QD, every day; TTR, transthyretin.

1. Maurer MS, et al. *N Engl J Med* 2018;379(11):1007–16; 2. Maurer MS, et al. *Circ Heart Fail* 2015;8(3):519–26; 3. Bulawa CE, et al. *Proc Natl Acad Sci USA* 2012;109(24):9629–34 4. VYNDAQEL® / VYNDAMAX® Prescribing Information. New York, NY: Pfizer Inc 5. Nativi-Nicolau J, et al. *Curr Opin Cardiol* 2018;33:571–9.



Transthyretin Cardiomyopathy (ATTR-CM) with Tafamidis Meglumine in the ATTR-ACT study



Current median survival expectations for untreated patients with ATTR-CM are **~2 to 5 years** from diagnosis.^{1–3} During this time, patients experience **declining health and quality of life (QoL)**.^{1,3}



ATTR-ACT was a Phase 3, international, randomized, double-blind, placebo-controlled evaluation of the safety and efficacy of tafamidis meglumine in patients with ATTR-CM.⁴



The primary outcomes showed that tafamidis meglumine **significantly reduced mortality and cardiovascular (CV)-related hospitalizations** compared with placebo over 30 months.⁴

Tafamidis was well tolerated, with a safety profile comparable to placebo.⁴

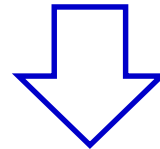


Patients treated with tafamidis demonstrated better overall scores in 6MWT distance, NT-proBNP concentration, KCCQ-OS, and PGA of overall health score compared to placebo despite both tafamidis and placebo treated patients showing progressively declining health at Month 30.^{4–7}

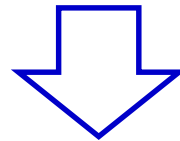


TTR Stabilization in patients with Tafamidis treated ATTR-CM

TTR stabilization has multiple meanings as per the current literature and is often associated with methods that do not measure the tetramer dissociation rate and/or include non-physiological conditions such as addition of denaturants



The receptor occupancy study by Tess et al leverages physiologically relevant measures of tafamidis binding to TTR tetramer and albumin in conjunction with therapeutically relevant patient data from ATTR-ACT to establish the relationship **between TTR occupancy, TTR stabilization, and clinical measures related to disease progression** in patients with ATTR-CM treated with tafamidis meglumine



The model-based characterization of these data is used to **assess the degree to which the clinically approved daily 80 mg dose of tafamidis meglumine achieves the full potential** for TTR kinetic stabilizers

Study Design



i In this study, TTR stabilization was defined as the degree to which the tetramer dissociation rate decreased in human plasma following the 20 mg and the approved 80 mg once daily (QD) doses of tafamidis meglumine.¹



This definition was chosen because the change in TTR dissociation rate under therapeutic and physiologic conditions is the most clinically relevant measure of pharmacologic activity for this class.²

Methods for characterizing the binding of tafamidis meglumine to both TTR tetramer and human albumin¹

- Isothermal titration calorimetry (ITC)
- Subunit fraction exchange
- Albumin binding (equilibrium dialysis methods)



Data obtained from ITC, subunit fraction exchange, and albumin binding experiments were simultaneously modelled to estimate a common set of **tafamidis meglumine binding dissociation constants to TTR ($K_{d,1}$ and $K_{d,2}$) and albumin ($K_{d,A}$)**

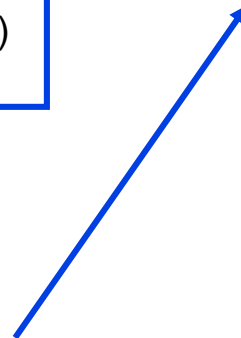
Clinical data from the Phase 3 ATTR-ACT study was used in this analysis¹

TTR concentrations over time were simultaneously characterized as a function of inter-individual variability in TTR production rate, and also as a function of tafamidis meglumine concentration, to develop **population TTR binding and concentration–time models**



Population disease progression response–time models were developed using three disease progression measures:

- N-terminal pro-B natriuretic peptide (NT-proBNP) plasma levels
- Kansas City Cardiomyopathy Questionnaire Overall Score (KCCQ-OS) score
- 6-minute walk test (6MWT) distance



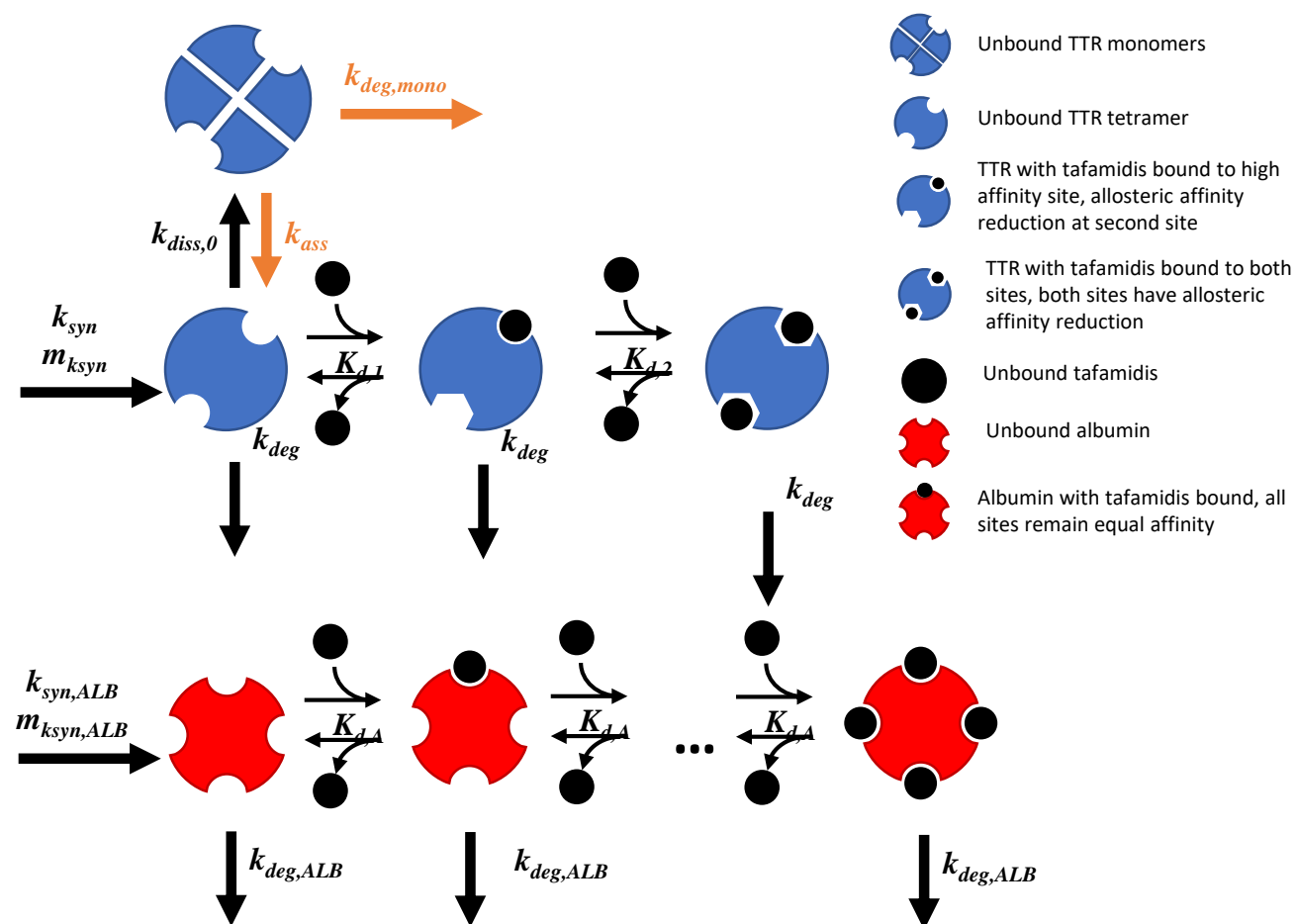


Results: Characterization of tafamidis meglumine binding to TTR and albumin in plasma

The final model structure used to characterize the relationship between tafamidis meglumine and TTR concentrations is shown in the figure

Use of these values provided **accurate predictions** of the concentration-dependent unbound fraction of tafamidis meglumine in human plasma and confidence in the TTR occupancies inferred by these parameters

Schematic of plasma TTR and albumin concentration-time models



TTR, transthyretin.
Tess DA, et al. *Amyloid* 2022;1-12.



Results: Characterization of tafamidis meglumine binding to TTR and albumin in plasma

- The two known tafamidis binding partners in human plasma are TTR and albumin
- In order to create a dynamic model of how TTR and tafamidis concentrations relate to TTR occupancy by tafamidis and unbound concentrations of TTR in human plasma:
 - The binding affinity estimates of tafamidis to TTR and albumin were calculated
 - Creation of population albumin and TTR concentration time models accounted for albumin and TTR concentrations changes over time

Binding affinity estimates

Modeling data from ITC, subunit fraction exchange, and equilibrium dialysis provided a reasonable characterization and precise binding affinity estimates:

- $K_{d,1}$ estimates of 5.08 nM (4.1-6.3)
- $K_{d,2}$ estimates of 203 nM (160-250)
- $K_{d,A}$ value of 3.53 μ M (3.0-4.2)
(95% confidence interval)

Population albumin and tafamidis concentration–time models

- The models used to estimate change in albumin or tafamidis concentrations used the rates in the table below
- The structure of these time-dependent aspects in the final model were determined based on the ability to provide a good fit to the data

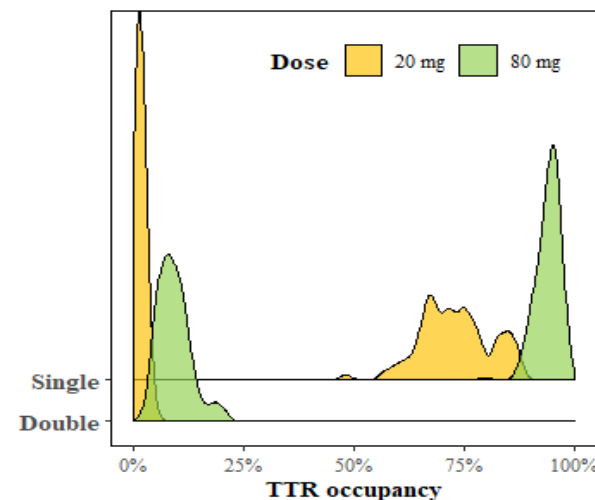
k_{syn}	TTR production rate
mk_{syn}	Linear slope of change of k_{syn}
$k_{\text{deg}, 0}$	Unbound TTR degradation rate
$k_{\text{deg}, \text{mono}}$	TTR monomer degradation rate
$k_{\text{diss}, 1}$	Single-bound tafamidis-bound TTR dissociation rates to monomer
$k_{\text{diss}, 2}$	Double-bound tafamidis-bound TTR dissociation rates to monomer
k_{ass}	Association rate of monomer back to tetramer
$k_{\text{syn}, \text{ALB}}$	Albumin production rate
$mk_{\text{syn}, \text{ALB}}$	Linear slope of change of $k_{\text{syn}, \text{ALB}}$
$k_{\text{deg}, \text{ALB}}$	Albumin fixed degradation rate



Tafamidis meglumine fractional TTR occupancy

- Population average plasma tafamidis meglumine concentrations associated with 20 and 80 mg QD doses were 8.8 and 35 μ M, respectively
- As such, tafamidis meglumine doses are estimated to achieve population average site occupancy as follows:**

	TTR occupancy	
	Single site	Dual site
20 mg	73%	1%
80 mg	94%	9%

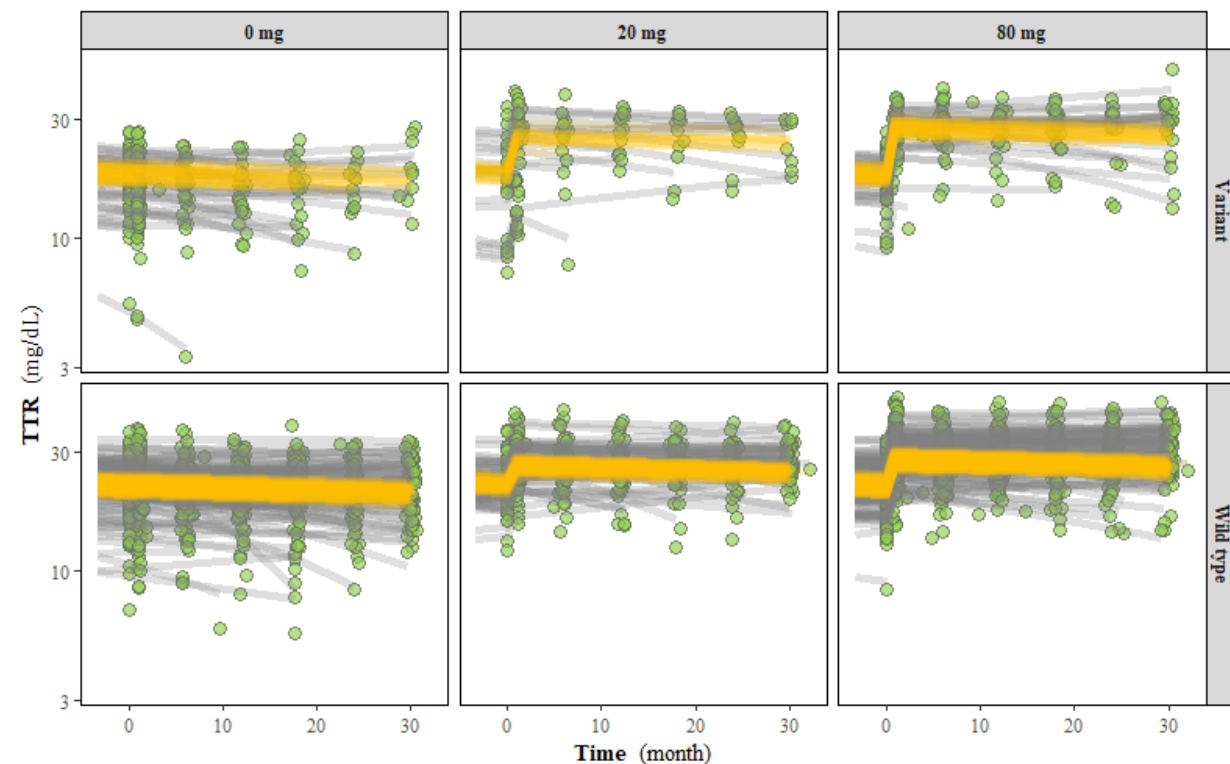


- Modelling, consistent with *in vitro* results and literature reports, found that the dissociation rate of TTR in the presence of singly bound tafamidis was negligible
- With a TTR single site occupancy of 94%, this results in tafamidis 80 mg achieving TTR stabilisation that is close to the estimated maximum stabilisation that would be achieved with 100% TTR occupancy



Population TTR binding and concentration–time model

- The model captured key characteristics of **placebo patients**:
 - **Lower initial TTR concentration in variant vs wild-type** captured by two-fold faster tetramer dissociation rate
 - **Decrease in TTR concentration over time**
- The **effect of Tafamidis on TTR concentration** was captured by a **decrease in the tetramer dissociation rate with a linear dependence on TTR occupancy**



Observed TTR concentrations (green circles)

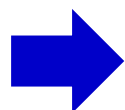
Individual predictions (grey lines)

Population predictions of individual's demographics (orange lines)



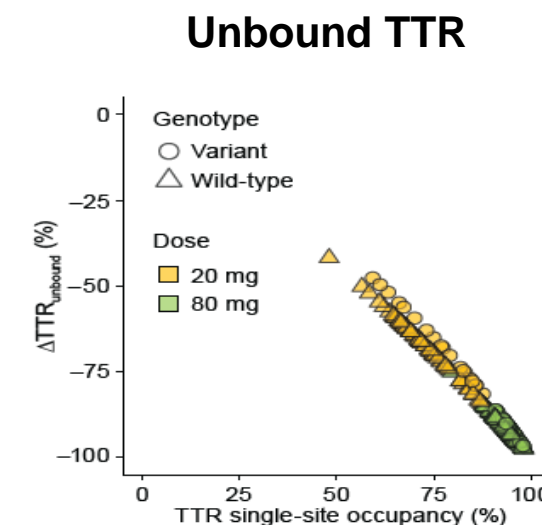
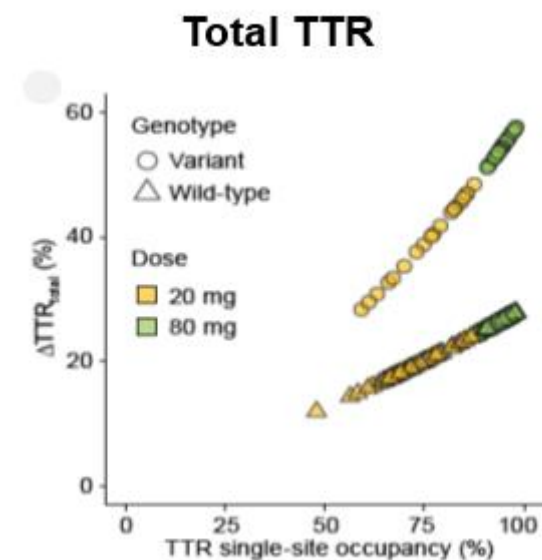
Effect of tafamidis meglumine on total and unbound TTR concentrations

- Tafamidis meglumine 20 and 80 mg increased **total TTR** concentrations by a population average of 19% and 26% in wild-type TTR genotype patients, and by 40% and 55% in variant TTR genotype patients, respectively



This is near what would be expected for **complete stabilization** of TTR (29% [wild-type] and 59% [variant TTR genotype]), which indicates that the approved dose of tafamidis meglumine provides a **degree of stabilization that is near the maximum achievable** (i.e. TTR dissociation <10%)

- Unbound TTR** decreased by an average of 67% and 92% with tafamidis meglumine doses of 20 and 80 mg, respectively



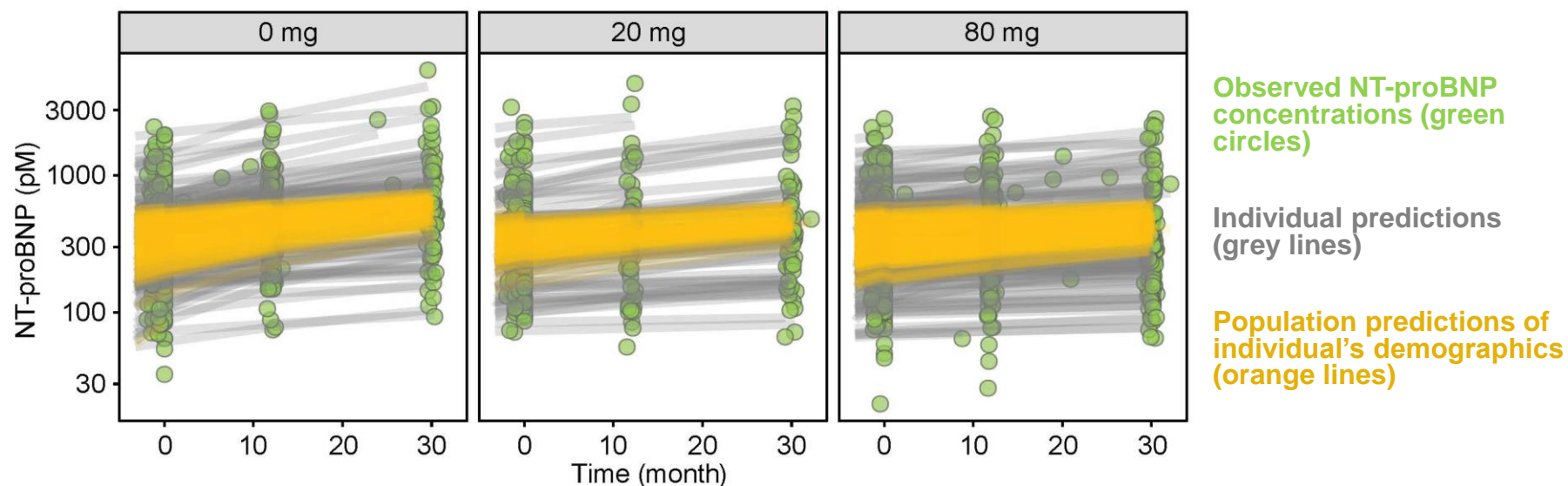


NT-proBNP concentration-time model

Tafamidis meglumine doses of 20 mg and 80 mg were estimated to inhibit the increase in population average NT-proBNP concentration by 39% and 53%, respectively



NTproBNP concentrations over time had a linear relationship with unbound TTR concentration



Maximal inhibition (at 100% TTR occupancy) was estimated to be **only slightly greater** (58%) than the average inhibition achieved with tafamidis meglumine at the 80 mg dose



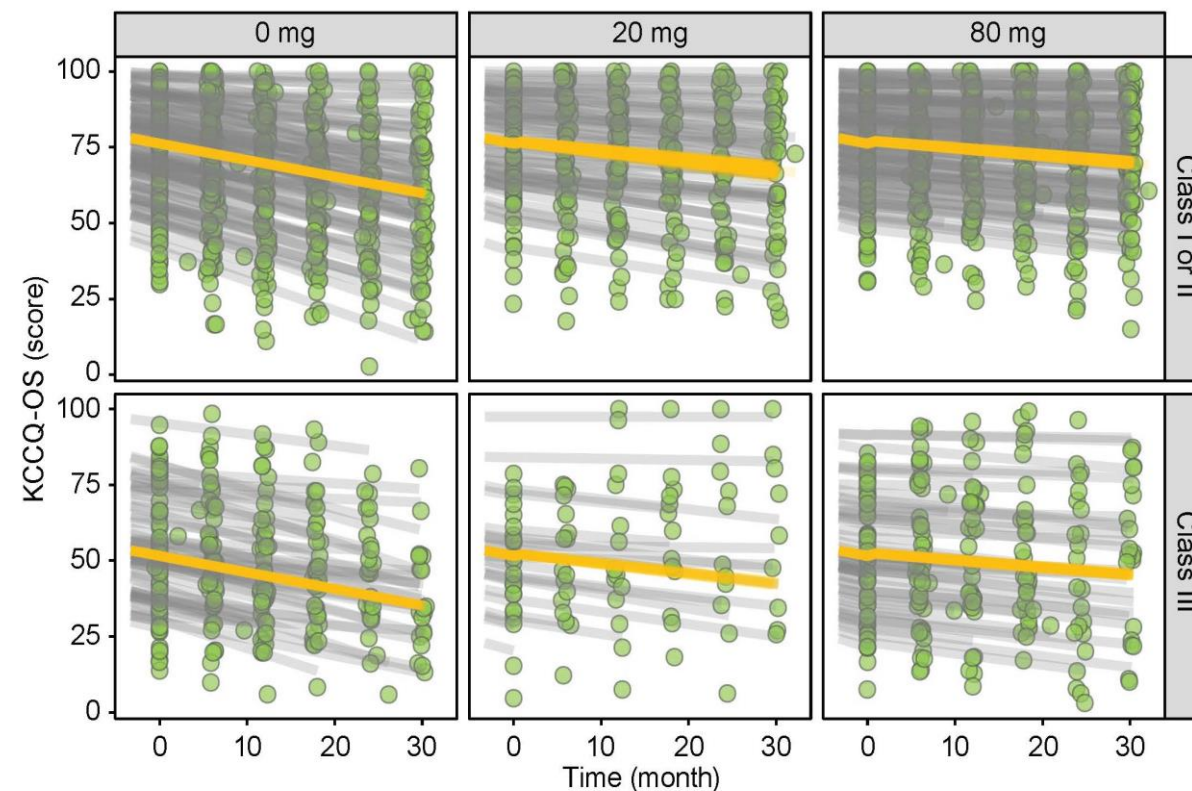
KCCQ-OS score-time model



KCCQ-OS scores over time had a linear relationship with unbound TTR concentration

Tafamidis meglumine doses of 20 mg and 80 mg were estimated to inhibit the decrease in KCCQ-OS by 41% and 56%, respectively

Maximal inhibition (at 100% TTR occupancy) was estimated to be **only slightly greater (61%)** than the average inhibition achieved with tafamidis meglumine at the 80 mg dose



Observed KCCQ scores (green circles)

Individual predictions (grey lines)

Population predictions of individual's demographics (orange lines)

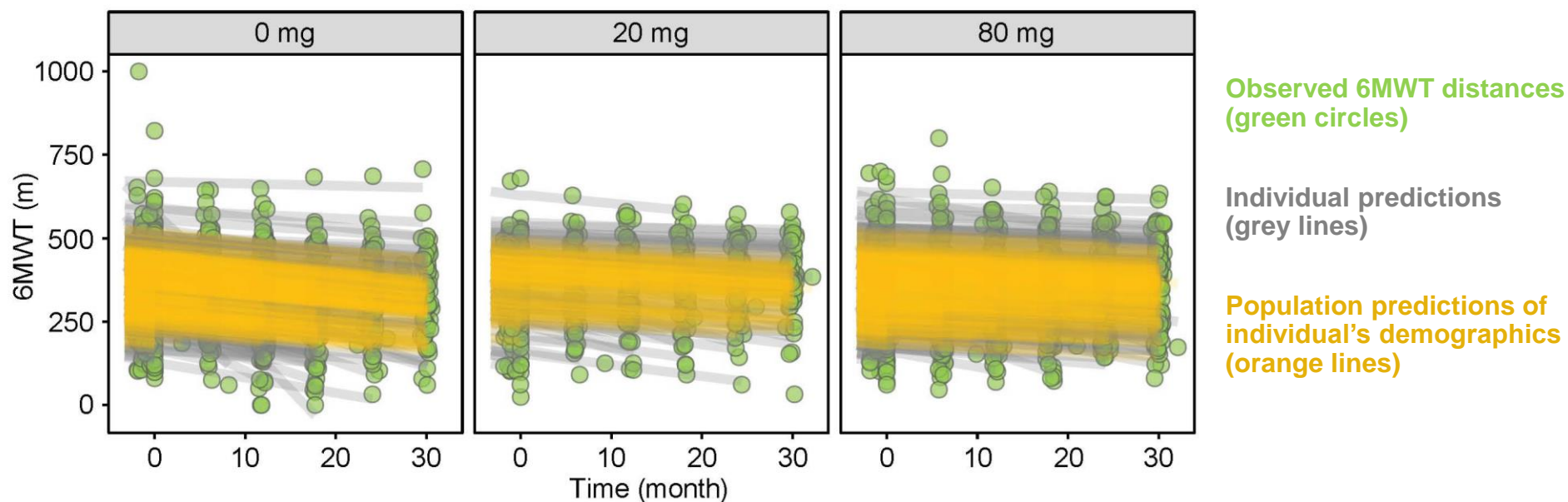


6MWT distance-time model

Tafamidis meglumine doses of 20 mg and 80 mg were estimated to inhibit the decline in average population 6MWT distance by 36% and 49%, respectively



6MWT over time had a linear relationship with unbound TTR concentration



Maximal inhibition (at 100% TTR occupancy) was estimated to be **only slightly greater (54%)** than the average inhibition achieved with tafamidis meglumine at the 80 mg dose



Conclusions

These results indicate that **singly bound tafamidis** provides **near-complete TTR stabilization**

The model demonstrates that the clinically approved dose of **80 mg tafamidis meglumine** achieves **a near- maximum (>90%) effect** on both TTR stabilization and disease-relevant measures

These findings support the value of TTR stabilization as a **clinically effective** treatment for ATTR-CM



Thank You

