



Water content, transition temperature and fragility influence protection and anhydrobiotic capacity

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ABSTRACT

Water is essential for metabolism and all life processes. Despite this, many organisms distributed across the kingdoms of life survive near-complete desiccation or anhydrobiosis. Increased intracellular viscosity, leading to the formation of a vitrified state is necessary, but not sufficient, for survival while dry. What properties of a vitrified system make it desiccation-tolerant or -sensitive are unknown. We have analyzed 18 different *in vitro* vitrified systems, composed of one of three protective disaccharides (trehalose, sucrose, or maltose) and glycerol, quantifying their enzyme-protective capacity and their material properties in a dry state. Protection conferred by mixtures containing maltose correlates strongly with increased water content, increased glass-transition temperature, and reduced glass former fragility, while the protection of glasses formed with sucrose correlates with increased glass transition temperature and the protection conferred by trehalose glasses correlates with reduced glass former fragility. Thus, *in vitro* different vitrified sugars confer protection through distinct material properties. Next, we examined the material properties of a dry desiccation tolerant and intolerant life stage from three different organisms. The dried desiccation tolerant life stage of all organisms had an increased glass transition temperature and reduced glass former fragility relative to its dried desiccation intolerant life stage. These results suggest in nature organismal desiccation tolerance relies on a combination of various material properties. This study advances our understanding of how protective and non-protective glasses differ in terms of material properties that promote anhydrobiosis. This knowledge presents avenues to develop novel stabilization technologies for pharmaceuticals that currently rely on the cold-chain.

Statement of significance: For the past three decades the anhydrobiosis field has lived with a paradox, while vitrification is necessary for survival in the dry state, it is not sufficient. Understanding what property(s) distinguishes a desiccation tolerant from an intolerant vitrified system and how anhydrobiotic organisms survive drying is one of the enduring mysteries of organismal physiology. Here we show *in vitro* the enzyme-protective capacity of different vitrifying sugars can be correlated with distinct material properties. However, *in vivo*, diverse desiccation tolerant organisms appear to combine these material properties to promote their survival in a dry state.

1. Introduction

Water is required for metabolism and so is often considered essential for life. However, a number of organisms, spread across every biological kingdom, are capable of surviving near-complete water loss through a process known as anhydrobiosis (Greek for “life without water”) [1]. As organisms dry, they face a number of physical and chemical changes to their cellular environment [1,2]. As water is lost, cellular constituents are concentrated, molecular crowding increases, pH and ionic concentrations change, and osmotic pressure increases [2]. These

physiochemical changes lead to detrimental perturbations such as protein unfolding, aggregation, and membrane leakage [2]. Importantly, drying is not an all-or-nothing process and these changes as well as the perturbation they induce, occur along a continuum, with some perturbations occurring earlier as an organism is dehydrating while others manifest later, once more substantial amounts of water have been lost [1–4]. How organisms survive desiccation is one of the enduring mysteries of organismal physiology.

Historically, anhydrobiosis has been thought to be mediated, at least in part, through the concentration of cellular constituents until these

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constituents solidify into a vitrified material (a glass). In this hypothesis, known as the ‘vitrification hypothesis,’ glasses slow physical and biochemical change, making them natural promoters of desiccation tolerance. Within the anhydrobiosis field, vitrification is considered a necessary process for desiccation tolerance [5–8].

However, a major shortcoming of the vitrification hypothesis has been known for decades, namely the observation that essentially every biological, or sufficiently heterogeneous, system will vitrify when dried, regardless of whether it is desiccation-tolerant or -sensitive [1,5–7]. This observation implies that while vitrification is necessary, it is not sufficient for desiccation tolerance and that there must be some property, or properties, that distinguishes a protective from a non-protective vitrified state [5,9]. The properties distinguishing a desiccation-protective glass from a non-protective glass are not currently fully understood.

Previous studies have identified that small additions of glycerol changes the enzyme-protective capacity of trehalose [10]. However, the material properties of these mixtures and how they correspond with changes in the level of protection have not been investigated. To address this gap in knowledge, we test the hypothesis that the enzyme-protective capacity of disaccharide-glycerol mixtures during desiccation correlates with their material properties. These properties include water content, glass transition temperature, and glass former fragility.

Water content is a property of vitrified materials that has been implicated in survival during extreme desiccation [4,11,12]. Water content refers to the mass percent of water in a desiccated sample. This can be measured by taking the starting mass of a desiccated sample and dividing by the mass of the sample after heating to a temperature sufficient to evaporate residual water. While hydrated, water molecules within a cell are able to solvate and then stabilize sensitive intracellular components. By retaining more water, it has been proposed that a vitrified material could prevent damage to sensitive intracellular components by maintaining hydration shells around them [3,4]. Additionally, residual water is implicated in several other proposed mechanisms of desiccation tolerance such as water entrapment [13–17], preferential exclusion [16,18], and the anchorage hypothesis [19–22].

Glass transition temperature (T_g) is the temperature at which a vitrified solid transitions from a glassy to a rubbery state [23–25]. Increases or decreases to the T_g of a vitrified material occur through the inclusion of an additive [26]. Increasing the glass transition temperature of a vitrified material has been observed to increase the shelf-life of sensitive proteins in a dry state [27] and is implicated in being essential for survival during desiccation at low relative humidity [3]. At the organismal level, it has been demonstrated that many anhydrobiotic organisms survive heating up to, but not beyond, their T_g [6,28]. This suggests that anhydrobiotic organisms rely on being in a vitrified state and the production of small molecules which increase T_g may be an effective strategy for increasing desiccation tolerance, or at least for increasing thermal tolerance while desiccated.

Finally, glass former fragility has been implicated as a key property of vitrified materials that promotes desiccation tolerance [9]. This property distinguishes strong glass forming materials, whose viscosity increases steadily well before the liquid-to-solid transition, from fragile glass forming materials, whose viscosity increases slowly at first but then rises abruptly at the onset of vitrification [9,23,29]. It should be noted that in this context, glass fragility/strength does not refer to the brittleness of a vitrified material, but rather to how the viscosity of the material changes as it approaches the point of vitrification. It is hypothesized that strong glass forming materials confer more protection during desiccation than their more fragile counterparts [4,9,23,29]. This hypothesis relies on the logic that a fragile glass forming material will not produce a sufficiently viscous state to slow down or prevent perturbation such as protein unfolding and aggregation until it is too late. Conversely, a strong glass forming material will increase in viscosity and provide protection along the continuum of drying.

To empirically test which, if any, of these three material properties correlate with desiccation tolerance, we first used a panel of simple

reductive systems, each composed of two mediators of desiccation tolerance - a disaccharide including maltose [30,31], sucrose [1,5,32], or trehalose [6,7,28,33–35] and the polyol glycerol [36–38]. Comparing the measured material properties (water content, T_g , and glass former fragility) of our disaccharide-glycerol glasses with their *in vitro* enzyme-protective capacities (Fig. 1a), we find that there is not a strict pattern in terms of the correlation of material properties to protection that all disaccharides follow. Instead, it appears that each disaccharide has a particular material property that is best correlated with its enzyme-protective capacity.

We extend this analysis into three organismal systems, each of which has a well-characterized desiccation tolerant and intolerant life stage(s). We find that while the enzyme-protective capacity of our *in vitro* systems tends to correlate with a single material property, our *in vivo* studies showed both increased glass transition temperature and reduced glass former fragility are hallmarks of desiccation tolerance. This suggests that anhydrobionts modulate multiple material properties to promote organismal desiccation tolerance.

The extension of our mechanistic understanding of the principles underlying desiccation-tolerance provides insights into how organisms can cope with changing, often extreme, environments. These findings also help to address the decades-long paradox that some glasses are protective, and others not, by providing insights into material properties that promote the protective capacity of different vitrifying mixtures. Our work identifies increased glass transition temperature and glass former fragility as major considerations in the development of technologies for the dry preservation of pharmaceuticals and the engineering of crops that are better able to withstand climate change and extreme weather.

2. Materials and methods

2.1. Disaccharide-glycerol mixtures

D-Maltose monohydrate was sourced from Caisson Labs (M004-500 GM). D-Sucrose was sourced from Sigma-Aldrich (S0389-500 G). D-Trehalose dihydrate was sourced from VWR (VWRB3599-1 KG). Glycerol was sourced from Biobasic (GB0232). Mixtures of each disaccharide and glycerol were made in 25 mM Tris at pH 7.0. Individual masses of each component were formulated (weight by weight) to additively produce mixtures of 10 g/L.

2.2. Sample desiccation

Samples were desiccated using a speedvac (Savant SpeedVac SC110 with a Thermo OFP400 vacuum pump) for 16 h. Prior to desiccation, 1 mL aliquot samples were dispensed into individual plastic weigh-boats (for aqueous samples) and at least 200 mg of organism samples were similarly loaded into individual plastic weigh-boats. The greater surface area of the weigh-boat, as opposed to Eppendorf tubes, allows for even desiccation of the entire sample, which reduces noise on the DSC. After the 16 h desiccation, DSC samples were transferred to pre-massed pairs of DSC aluminum hermetic pan and aluminum hermetic lids (TA 900793.901 and 901684.901, respectively) while TGA samples were transferred to pre-tared platinum crucibles (TA 957207.904), and XRD samples were kept in the desiccation weigh-boats. DSC sample masses were determined after the sample was sealed within the pan and lid.

2.3. Single crystal X-ray diffractometry

Powder diffraction patterns for the samples were measured at 20 °C on a Bruker SMART APEX II CCD area detector system equipped with a graphite monochromator and a Mo K fine-focus sealed tube operated at 1.2 kW power. The dried samples were rolled into a ball of approximate diameter 0.5 mm, mounted on a goniometer head using a glass fiber, and centered using the APEX2 software. The detector was placed at 6.12 cm during the data collection. Four frames of data were collected at four

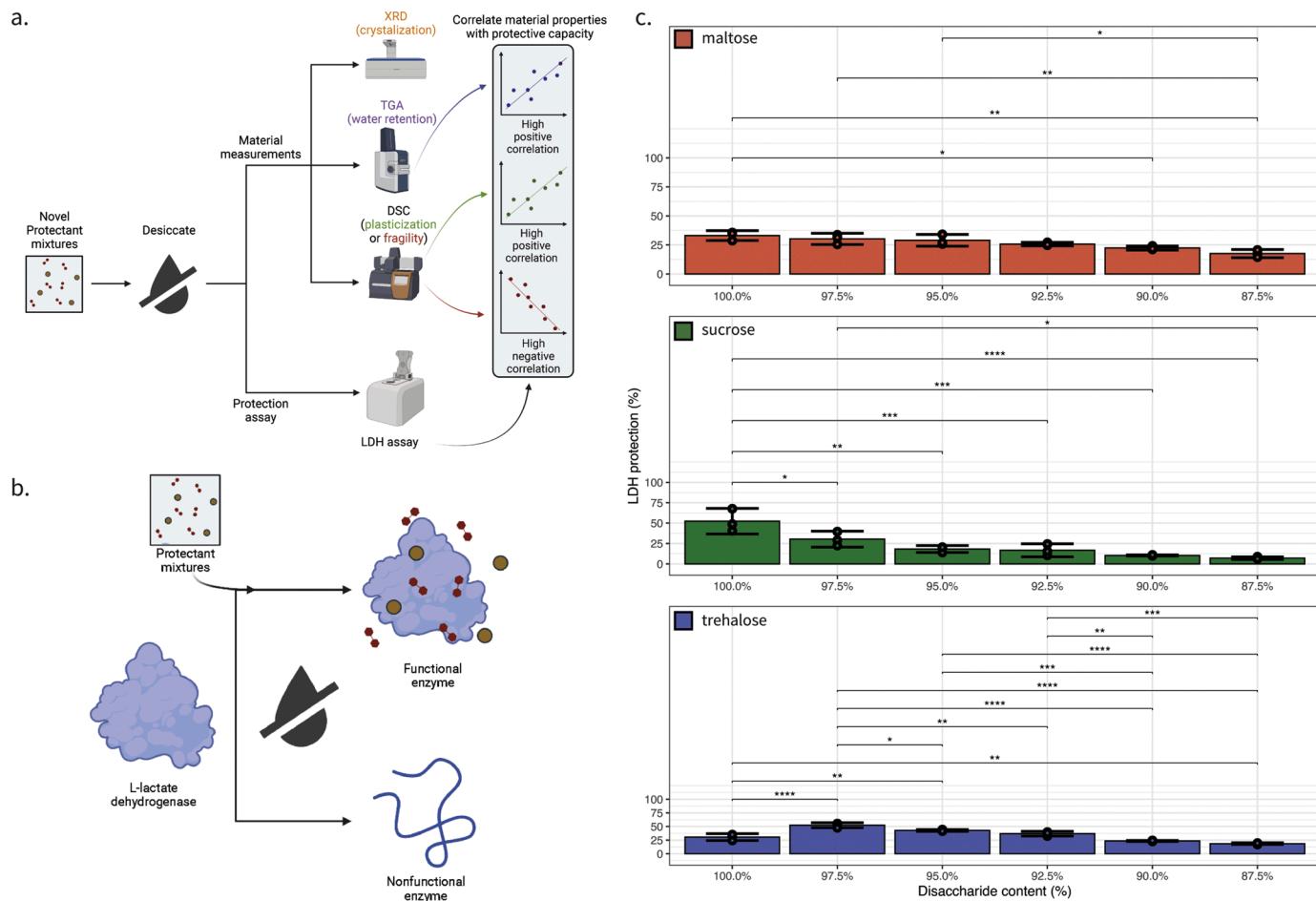


Fig. 1. Vitrified disaccharide-glycerol mixtures display variable enzyme-protective properties. a) Schematic overview of workflow to produce correlation values between material properties and enzyme protection effect. b) Schematic of LDH assay. c) Lactate dehydrogenase (LDH) enzyme protection values for disaccharide-glycerol mixtures organized by disaccharide. Statistics calculated using a one-way ANOVA and Tukey post-hoc test: p-value ≤ 0.05 (*) p-value ≤ 0.01 (**), p-value ≤ 0.001 (***), pairwise relationships not shown are not significant, error bars represent 95% CI.

different sets of angles with a scan width of 0.2° and an exposure time of 3 min per frame. The frames were integrated using the APEX2 program. The measured powder diffraction images were integrated, and the data were plotted in the 5 to 50° 2 θ . All analysis was performed using the APEX3 Software Suite V2017.3-0, Bruker AXS Inc.: Madison, WI, 2017.

2.4. Lactate dehydrogenase (LDH) enzyme protection assay

LDH assays were performed using a combination of methodologies described in Goyal et al. [49] and Boothby et al. [7]. Briefly, stock solutions of 25 mM Tris HCl (pH 7.0), 100 mM sodium phosphate (pH 6.0), and 2 mM pyruvate prepared in bulk and stored at room temperature prior to assay. In addition, 10 mM NADH was also prepared prior to the assay and then stored at 4°C . L-Lactate Dehydrogenase (LDH) was sourced from Sigma (SKU #10127230001) and is supplied in ammonium sulfate at a pH of approximately 7. Prior to assay, LDH was diluted to a working concentration of 1 g/L. Experimental disaccharide-glycerol mixtures were formulated with LDH at a 1:10 (LDH:disaccharide-glycerol) ratio. Enough solution was prepared for three test excipient replicates and three control replicates each with a total volume of 50 μL . Each experimental and control mixture were aliquoted into a 1.5 mL microcentrifuge tube. The test mixtures were then vacuum desiccated for 16 hours with controls kept refrigerated at 4°C . After vacuum desiccation, control and test excipient mixtures were brought to 250 μL total volume. Absorbance readings at 340 nm wavelength were taken every two seconds for 60 s with a quartz cuvette on a Thermo

Scientific NanoDrop One^c (Thermo Scientific 840274200) spectrophotometer. A 100 mM sodium phosphate and 2 mM pyruvate solution was used as a blank. For control samples and experimental samples, 10 μL of control mixture or 10 μL of experimental sample mixture were combined with 10 μL of NADH and 980 μL of the 100 mM sodium phosphate, 2 mM pyruvate solution and then were measured.

This same procedure was used to measure the A340 of the experimental and control mixtures. A340 was plotted as a function of time and the slope of the linear portion of this plot calculated. A ratio of experimental over control slope, multiplied by one hundred, was taken to produce the percent protection of the experimental mixture. All raw LDH protection data is available in Supplemental File 8 while calculation of percent protection is available in Supplemental File 1.

2.5. Thermogravimetric analysis

Samples were run on a TA TGA5500 instrument in 100 μL platinum crucibles (TA 952018.906). Crucibles were tared prior to each run and prior to sample loading. Crucibles were loaded with between 5 mg and 10 mg of sample mixture. Each sample was heated from 30°C to 220°C at a 10°C per minute ramp. All TGA data and thermograms are available in Supplemental File 2 and Supplemental File 3.

Determination of water loss was conducted using TA's Trios software (TA Instruments TRIOS version #5.0.0.44608). Thermograms were used to calculate starting masses of samples and the mass of samples at the plateau that occurs after $\sim 100^\circ\text{C}$ but before the thermal denaturation at

~200 °C. The Trios software “Smart Analysis” tool was used to identify the inflection point between these two mass loss events.

2.6. Differential scanning calorimetry

Samples were run on a TA DSC2500 instrument with Trios software (TA Instruments TRIOS version #5.0.0.44608). Analysis of DSC output was performed using Trios software. The heating run consisted of being loaded at ambient temperatures of 20 °C, then cooled at a rate of 13.2 °C per minute ramp to -10 °C where the temperature was equilibrated at -10 °C and then heated using a 10 °C per minute ramp to 220 °C. All raw DSC measurements are available in Supplemental File 4 and Supplemental File 5.

2.7. Fragility (m-index) calculation

Trios software (TA Instruments TRIOS version #5.0.0.44608) provided by TA Instruments was used to perform analysis of the DSC data. Calculations of glass former fragility (m-index) were performed based on equations 10 and 14 proposed by Crowley and Zografi [39]. On a thermogram with a completed heating ramp to 220 °C, the degradation peak, melt peak, and glass transition were identified. The Trios software built-in Onset and Endset analysis was used to determine the Glass transition onset and offset (endset). The software-identified glass transition onset and offset was used to calculate the m-index using Crowley and Zografi's equations 10 where m is the alternative fragility parameter, ΔE_{T_g} is the activation enthalpy of structural relaxation at T_g , R is the gas constant, T_g is the experimental glass transition temperature onset, and 14 where ΔE_η is the activation enthalpy for viscosity, R is the gas constant, T_g is the experimental glass transition onset temperature, T_g^{off} is the experimental glass transition offset temperature, and constant is an empirical constant of 5 [39]. A mean of each set of replicates was obtained. All fragility calculations are available in Supplemental File 6.

$$m = (\Delta E_{T_g}) / ((\ln 10) R T_g) \quad (1)$$

$$constant = (\Delta E_\eta / R) \left(1 / T_g - 1 / T_g^{off} \right) \quad (2)$$

2.8. Selected organisms for *in vivo* assays

Artemia franciscana, *Caenorhabditis elegans*, and *Saccharomyces cerevisiae* were all selected based on availability of access to both desiccation sensitive and desiccation tolerant life stages.

Artemia franciscana adults (#1) and cysts (#11) were acquired from the Northeast Brine Shrimp, LLC. *A. franciscana* adults were separated from culture media by tube-top filtration using a pluriSelect 200 µm pluriStrainer (#43-50200-03) prior to desiccation.

Caenorhabditis elegans non-dauered *daf-2* strains were cultured in S Medium [40] at 16 °C and fed with *E. coli* OP50. Dauer pre-conditioned *daf-2* strain worms were cultured identically to non-dauer worms but when the initial aliquot of *E. coli* OP50 were consumed an additional five days of incubation at 25 °C was given to starve the culture and induce dauer arrest. Dauer worms were then placed on non-spotted agar plates in a 25 °C and 95% relative humidity atmosphere for 3 additional days to accomplish pre-conditioning. Both non-dauer and pre-conditioned dauer *C. elegans* sample sets were separated from media and residual OP50 food stock by tube-top filtration using a pluriSelect 200 µm pluriStrainer (#43-50200-03) prior to desiccation.

Saccharomyces cerevisiae strain BY4742 were cultured aerobically in YPD media [41] at 30 °C. After nine hours samples were taken representing the logarithmic growth stage yeast. The remaining culture was allowed to grow for an additional five days to ensure confluent growth in the stationary phase. Both logarithmic and stationary phase samples were pelleted by centrifugation, media was decanted, and a water wash

of cell pellets was performed prior to desiccation.

2.9. Statistical methods

One-way ANOVA and Tukey post-hoc test was used for all pairwise comparisons. A p-value of less than 0.05 being one level of statistical significance (*), less than 0.01 being two levels of statistical significance (**), and less than 0.001 being three levels of statistical significance (***)). All error values represent 95% confidence intervals.

Student's T-Test was used to compare statistical differences between desiccation tolerant and desiccation sensitive life stages. A p-value of less than 0.05 being one level of statistical significance (*). All error values represent 95% confidence intervals.

3. Key resources table

Reagent or resource	Source	Identifier
D-Maltose monohydrate	Caisson	M004-500GM
D-Sucrose	Sigma-Aldrich	S0389-500G
D-Trehalose dihydrate	VWR	VWRB3599-1KG
Glycerol	Biobasic	GB0232
Tris HCl	US Biological	1185-53-1
Sodium pyruvate	TCI	P0582
NADH, Disodium Salt	Millipore Sigma	481913
L-Lactate Dehydrogenase	Sigma	SKU #10127230001
100 µL platinum crucibles	TA instruments	TA 952018.906
DSC aluminum hermetic pan	TA instruments	TA 900793.901
DSC aluminum hermetic lids	TA instruments	TA 901684.901
200 µm pluriStrainer	pluriSelect	#43-50200-03
<i>A. franciscana</i> adults	Northeast Brine Shrimp LLC	#1
<i>A. franciscana</i> cysts	Northeast Brine Shrimp LLC	#11
<i>C. elegans</i> <i>daf-2</i>	Caenorhabditis Genetics Center	CB1370
<i>E. coli</i> OP50	Caenorhabditis Genetics Center	OP50
<i>S. cerevisiae</i> BY4742	ATCC	201389

4. Results

4.1. Disaccharide-glycerol mixtures vitrify when dried

To begin to address which material properties correlate with enzyme-protective capacity in a vitreous state, we generated 18 different glass forming mixtures composed of one of three disaccharides (maltose, sucrose, or trehalose) and varying amounts of glycerol [10, 42]. Previous studies have identified that small additions of glycerol change the enzyme-protective capacity of trehalose. However, the material properties of these mixtures and how they correlate with changes in the level of protection have not been investigated. Furthermore, it is not known to what extent additions of glycerol will influence the material and enzyme-protective properties of other disaccharides or how these two properties are linked [10].

Maltose is a reducing disaccharide consisting of two glucose molecules joined by an α (1→4) bond [43]. Sucrose is a non-reducing disaccharide formed by the glycosidic linkage between C1 of a glucose molecule to the C2 of a fructose molecule [44]. Trehalose is a non-reducing disaccharide formed through the (1-1) glycosidic linkage of two glucose molecules [45]. While trehalose and sucrose are better recognized mediators of desiccation tolerance, all three of these disaccharides are known to accumulate in a number of anhydrobiotic organisms during drying and in many cases they are known to be essential for surviving desiccation [6,28,30,31,33–35]. Maltose, though rarely seen as a desiccation-protectant in nature, is accumulated in some resurrection plants [6,28,30,31,33–35]. Furthermore, maltose has been shown to confer, short- but not long-term, desiccation tolerance in yeast,

likely due to its propensity to be reduced [34]. Glycerol is a polyol that has been implicated in the tolerance of several stresses ranging from drying to freezing [36,38].

Mixtures containing 100%, 97.5%, 95%, 92.5%, 90%, and 87.5% of a single disaccharide (maltose, sucrose, or trehalose), combined with glycerol (weight % by weight % with glycerol) were created. Mixtures were dried overnight in a vacuum desiccator for 16 hours to produce glasses.

To ensure that these mixtures vitrified, rather than crystallized, each sample's powder diffraction pattern was observed by powder X-ray diffraction (XRD) using Mo K α radiation. The integrated plots of our 18 vitrifying samples contain a nearly identical broad absorption, and do not reveal any sharp peaks (Fig. S1a and S1b). The absence of sharp peaks indicates that the samples are glassy with no crystallinity [46,47]. Powder diffraction data was also measured for a desiccated sample of D-(+)-glucose, which is known to crystalize when dried. In contrast to the XRD patterns measured for the disaccharide-glycerol samples, the diffraction pattern for glucose exhibited a large number of distinct, closely spaced peaks due to the crystalline nature of this sample (Fig. S1a and S1b). These results indicate that maltose, sucrose, or trehalose by themselves or in conjunction with varying degrees of glycerol vitrify when dried under the drying regime used here (see Methods).

4.2. Disaccharide-glycerol mixtures have varying levels of enzyme-protection during desiccation

To address the question of which property(s) of a vitrified solid correlate with enzyme-protective capacity during desiccation, we assessed the ability of our 18 disaccharide-glycerol mixtures to protect the enzyme lactate dehydrogenase (LDH) during desiccation [7,48,49] (Fig. 1b). Previous reports have shown that LDH is sensitive to desiccation and that drying, and rehydration of this enzyme results in ~95–99% loss in functionality [49] (Fig. 1b). Taking the ratio of the enzymatic activity of rehydrated *versus* control LDH, we observed that the protection of desiccated LDH varied significantly between different disaccharide-glycerol mixtures (Fig. 1c).

For maltose-glycerol mixtures, the highest level of protection was observed for our 100% maltose sample, while the lowest level of protection was conferred by the 87.5% maltose sample. No significant difference in protection was observed until the percentage of maltose in mixtures reached 92.5%, after which protection steadily decreased (Fig. 1c, top panel).

Similar to maltose-glycerol mixtures, sucrose-glycerol mixtures showed the highest level of protection at 100% sucrose, while the lowest level of protection was conferred by the 87.5% sucrose sample. However, unlike maltose samples, sucrose mixtures rapidly lost enzyme-protective capacity with statistically significant decreases in protection being observed upon the first (2.5%) addition of glycerol (Fig. 1b, middle panel).

Trehalose-glycerol mixtures differed from both maltose and sucrose mixtures in that the highest level of protection was achieved at 97.5% (Fig. 1c, bottom panel). Additionally, while mixing maltose or sucrose with glycerol resulted in decreases in protection, additions of glycerol to trehalose were found to have a nonmonotonic relationship with protection (Fig. 1c, bottom panel).

These results demonstrate that different disaccharide and glycerol mixtures provide varying levels of protection to LDH during desiccation and rehydration, with trehalose responding in a non-monotonic fashion, maltose decreasing in enzyme-protective capacity in a linear fashion, and protection conferred by sucrose decreasing exponentially as a function of glycerol content.

4.3. Water content correlates with the enzyme-protective capacity of maltose-glycerol, but not trehalose-glycerol or sucrose-glycerol, mixtures

To begin to assess which properties of a vitrified system correlate

with protection, we first assessed whether or not water content could account for these differences. To assess whether the differences in protection observed in our vitrified samples correspond to the amount of water they retain, we tested each of our mixtures using thermogravimetric analysis (TGA) (Fig. 2a and b). Water contents of our 18 dry mixtures ranged from 10.52 to 0.81% (Fig. 2c). Pure dry trehalose contained approximately 8% water content, while sucrose and maltose contained less and approximately the same water content (5–6%) (Fig. 2c). In maltose-glycerol mixtures the amount of retained water decreased with each addition of glycerol (Fig. 2c). In contrast, water content did not vary significantly in any sucrose-glycerol mixture. The trehalose-glycerol mixtures only demonstrate statistically significant decreases in water content after the addition of 5% glycerol (Fig. 2c).

After observing distinct water retentive behaviors in dry disaccharide-glycerol mixtures, we assessed the relationship between water content and protection. For each sugar, there was a positive trend between enzyme-protection and increasing water content, however the correlation between these properties varied significantly between sugars. For maltose mixtures, this correlative analysis produced an R^2 value of 0.96 (p-value = 0.00052) (Figs. 2d and S5a), an R^2 value of 0.13 (p-value = 0.48) for sucrose mixtures (Figs. 2d and S5b), and R^2 value of 0.28 (p-value = 0.28) for trehalose mixtures (Figs. 2d and S5c). These results indicate that for maltose-glycerol systems the amount of retained water is a good indicator of the enzyme-protective capacity in the dry state. However, the amount of water in mixtures made of dry trehalose or sucrose and glycerol is a poor indicator of enzyme-protective capacity (Fig. 2d).

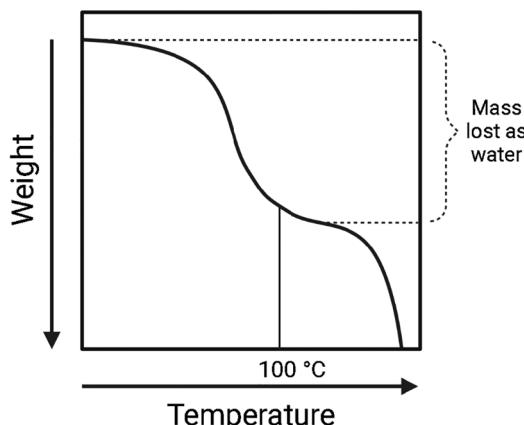
It should be acknowledged that precise determinations of water content using TGA can be complicated by an overlap between the offset of water evaporation and the onset of deterioration of a material. We found that for maltose and sucrose samples there was no such overlap, however in some cases trehalose samples showed an overlap requiring more refined methods of analysis. For example, when measuring the water content of some trehalose-glycerol mixtures, the step transitions calculated were numerous and ambiguously bordered the step transitions attributed to deterioration of a sample (Fig. S7b). In these cases, we made use of the derivative weight loss to differentiate between water loss and sample deterioration (Fig. S7a and S7b). This may result in less accurate determinations of water content for very complex thermograms. However, the behavior of most samples, especially the maltose- and sucrose-glycerol mixtures, were much more straightforward and allowed for simple differentiation of the water-loss and deterioration processes (Fig. S7a). Adding validity to our determination of water contents, even in complex samples, was the alignment of glass transition temperatures and water contents from previously published work [14].

4.4. An increase in the glass transition temperature correlates with enzyme protection conferred by maltose-glycerol and to a lesser extent sucrose-glycerol and trehalose-glycerol mixtures in the dry state

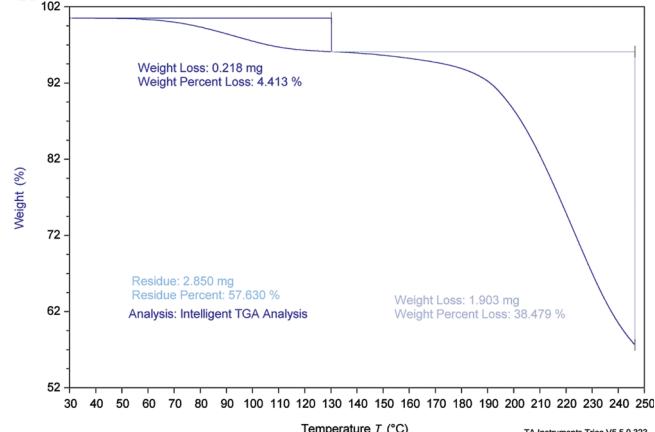
After observing that water retention is a poor indicator of the enzyme-protective capacity of trehalose- and sucrose-glycerol glasses, we wondered if glass transition temperature (T_g) might be a property that correlates with the stabilizing effects of these sugars. Previous work has established that even small amounts of additives, such as glycerol, can lead to decreased or increased T_g of a vitrifying material (Fig. 3a and b) [50–55]. With this in mind, we were curious if different additions of glycerol to our disaccharides would serve to increase or decrease the T_g resulting glasses, and whether or not these changes in T_g correlates with the enzyme-protective capacity of our mixtures.

The glass transition temperature onset and offset, the temperatures that the material starts and stops undergoing a change from a glassy state to a rubbery state, were assessed using differential scanning calorimetry (DSC), and the glass transition midpoint temperature calculated by taking the mean of the onset and offset temperature (Supplemental File 6). Figs. 3c, S3a, and S3b show the average T_g onset, offset, and

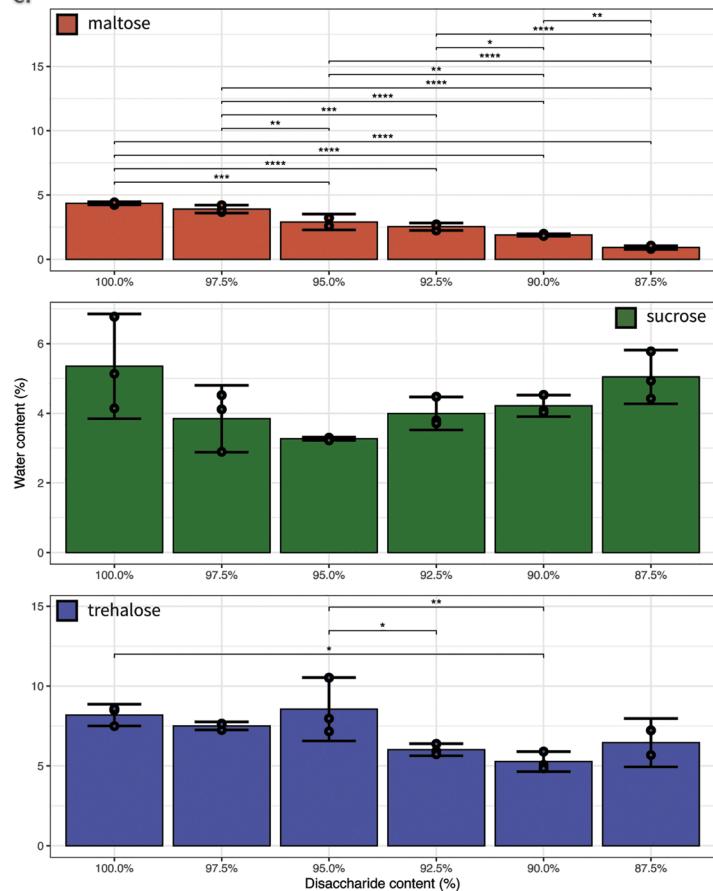
a.



b.



c.



d.

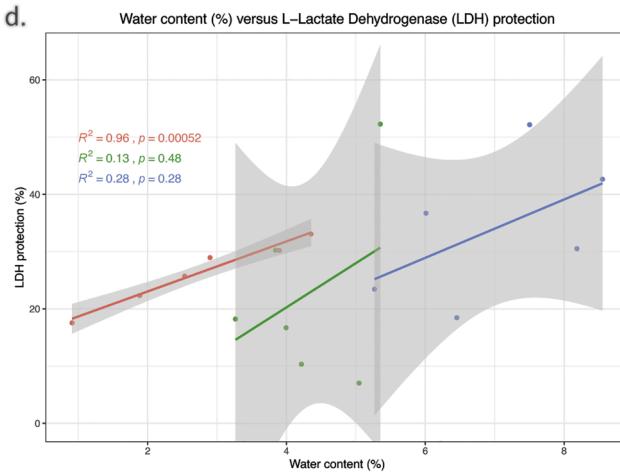


Fig. 2. Water content correlates with the enzyme-protective capacity of maltose-glycerol but not sucrose- or trehalose-glycerol glasses. a) Schematic representation of an idealized TGA thermogram highlighting region of mass lost as water. b) Example TGA thermogram of a 100% trehalose-glycerol mixture. c) Water content values for disaccharide-glycerol mixtures organized by disaccharide. d) Correlation plot of water content versus protection for all disaccharide-glycerol mixtures (red = maltose, green = sucrose, blue = trehalose). Statistics calculated using a one-way ANOVA and Tukey post-hoc test: p-value ≤ 0.05 (*), p-value ≤ 0.01 (**), p-value ≤ 0.001 (***), pairwise relationships not shown are not significant, error bars represent 95% CI.

midpoint for each of our mixtures. Here one can see that when considering the glass transition midpoint even small additions of glycerol, starting at 2.5%, act to decrease the T_g for trehalose, while decreasing the T_g of maltose began at 7.5% glycerol and a decrease in sucrose is only observed with an even larger (12.5%) addition of glycerol (Fig. 3c). For all disaccharide-glycerol mixtures, glass transition midpoint temperatures are positively correlated with enzyme protective capacity, but only the maltose-glycerol mixtures demonstrated a significant correlation ($p = 0.0063$) (Fig. 3d).

However, when considering the onset or offset glass transition temperature for each set of disaccharide-glycerol mixtures, we saw different behaviors appear. For example, when observing changes to the glass transition onset temperature for the maltose-glycerol mixtures, even after the addition of 12.5% glycerol there was no significant change (Fig. S3a). As opposed to maltose, sucrose saw a significant decrease in glass transition onset temperature after the addition of 12.5% glycerol (Fig. S3a). Finally, the trehalose-glycerol mixtures show statistically significant changes in the glass transition onset temperature beginning

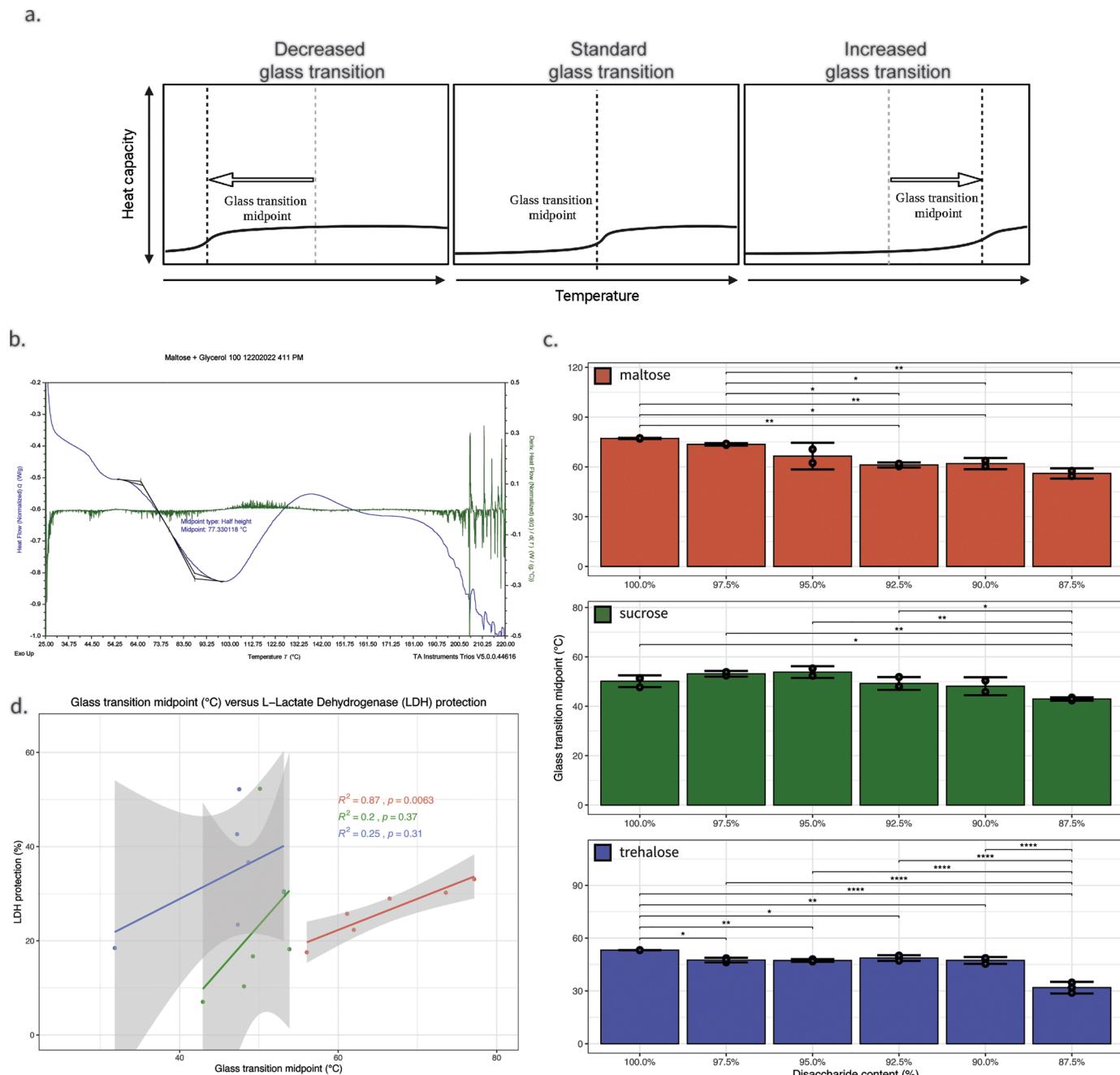


Fig. 3. Anti-plasticization of maltose- and sucrose-, but not trehalose-glycerol glasses correlate with enzyme protection. a) Schematic representation of an idealized DSC thermogram illustrating a decrease or increase in glass transition midpoint. b) Example DSC thermogram of a 100% trehalose-glycerol mixture. c) Glass transition midpoint values for disaccharide-glycerol mixtures organized by disaccharide. d) Correlation plot of glass transition midpoint versus protection for all disaccharide-glycerol mixtures (red = maltose, green = sucrose, blue = trehalose). Statistics calculated using a one-way ANOVA and Tukey post-hoc test: p-value ≤ 0.05 (*) p-value ≤ 0.01 (**), p-value ≤ 0.001 (***), pairwise relationships not shown are not significant, error bars represent 95% CI.

with the first addition of glycerol (Fig. S3a). The trehalose-glycerol mixtures also showed a statistically significant decrease after the addition of 12.5% glycerol (Fig. S3a).

When observing the offset of glass transition temperature for each of the disaccharide-glycerol mixtures, we observed that the maltose-glycerol and sucrose-glycerol mixtures demonstrated similar behavior to the midpoint observations. Specifically, starting at 5% for maltose and 12.5% for sucrose, glycerol acts to decrease the glass transition offset temperature (Fig. S3c). On the other hand, trehalose does not show a statistically significant decrease in the glass transition offset until after the addition of 10% glycerol (Fig. S3c).

Next, we evaluated the relationship between enzyme-protective

capacity and the onset, offset, and midpoint T_g for each mixture (Figs. 3d, S4, S5d, S5e, and S5f). This correlative analysis produced positive correlations with R^2 values ranging from 0.87 to 0.81 for maltose (Figs. 3d, S4, S5d), positive correlations with R^2 values ranging from 0.38 to 0.1 for sucrose (Figs. 3d, S4, S5e), and positive correlations for trehalose the R^2 values ranged from 0.36 to 0.1 (Figs. 3d, S4, and S5f). We observe that the enzyme-protective capacity of the maltose-glycerol mixtures are influenced significantly by variation in their T_g and that the enzyme-protective capacity of the sucrose- and trehalose-glycerol mixtures are moderately influenced by variation in their T_g (Fig. 3d).

Finally, we examined the relationship between the enzyme-

protective effect of our sugar mixtures and the difference between the temperature at which LDH is protected ($T_{exp} = 22^\circ\text{C}$) and their T_g . We reasoned that for a mixture with a T_g close to T_{exp} protection might be lowered due to the mixture undergoing a relaxation.

Consistent with this reasoning, for all disaccharide-glycerol mixtures enzyme-protection and T_{exp} - T_g had a negative correlation (S6). However, only the maltose mixtures approached, but was not, a significant correlation between enzyme-protection and T_{exp} - T_g (S6). This indicates that at the temperature used in this study, relaxation of dry sugar mixtures due to a similarity in T_{exp} and T_g is not a significant influence on enzyme-protection.

Our results demonstrate that generally, protection positively

correlates with increasing T_g . However, this correlation is clearly stronger for some sugar-glasses (e.g., maltose) compared to others (e.g., sucrose).

4.5. Vitrified maltose-glycerol and trehalose-glycerol mixtures demonstrate a reduction in glass former fragility that is correlated with enzyme-protection

Next, we empirically determined the glass former fragility (m-index) [25,39] of our disaccharide-glycerol mixtures (Fig. 4a). Moynihan et. al. described the use of thermal methods to explore the relationship between the width of a glass transition and the activation enthalpy for

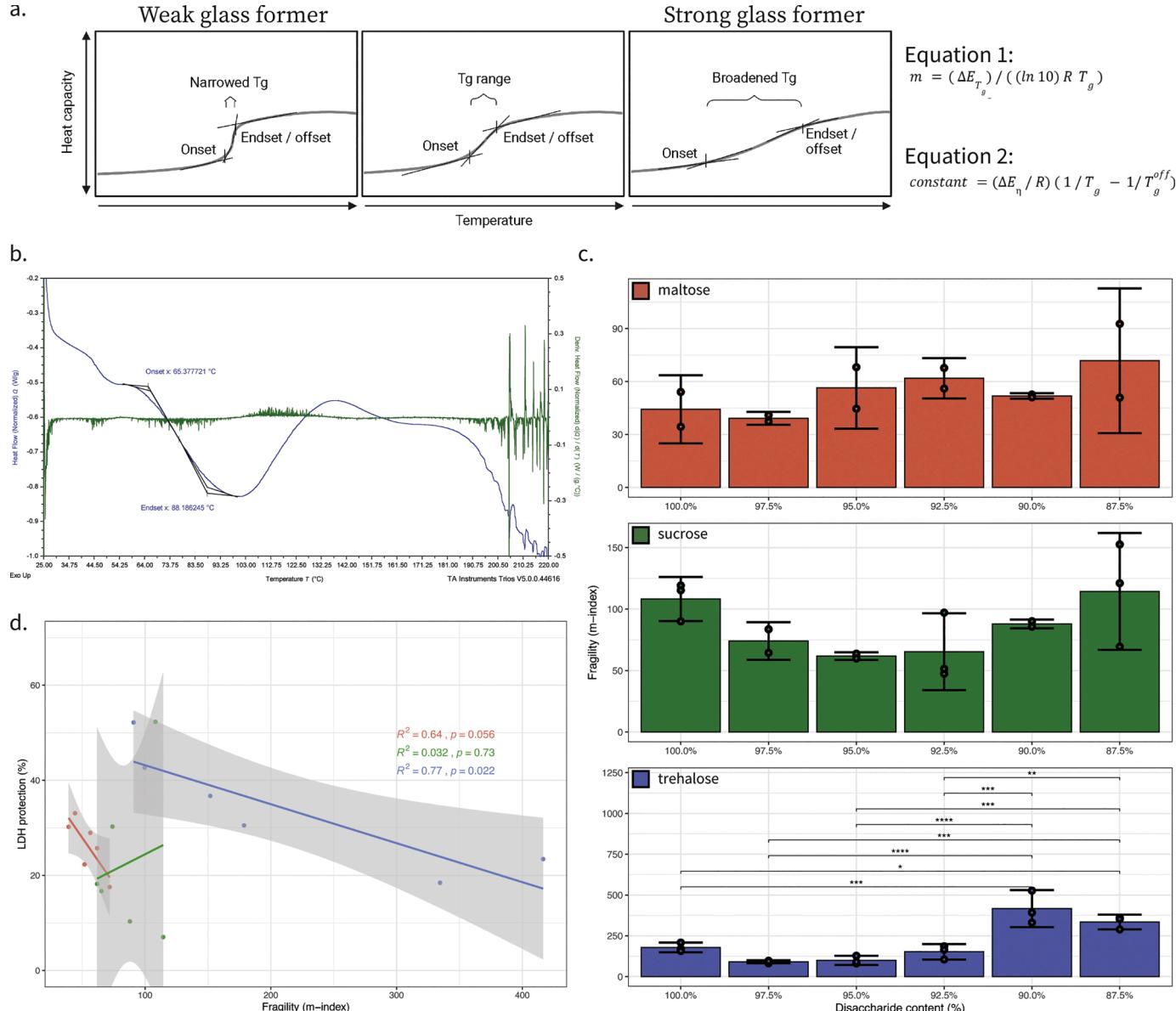


Fig. 4. Reduced glass forming fragility of maltose- and trehalose-, but not sucrose-glycerol glasses correlate with enzyme protection. a) Schematic representation of idealized DSC thermograms illustrating a narrowing or broadening of the glass transition and the two equations used to calculate m-index. Eq. (1) where m is the alternative fragility parameter, ΔE_{T_g} is the activation enthalpy of structural relaxation at T_g , R is the gas constant, T_g is the experimental glass transition temperature onset, and 2 where ΔE_η is the activation enthalpy for viscosity, R is the gas constant, T_g is the experimental glass transition onset temperature, T_g^{off} is the experimental glass transition offset temperature, and constant is an empirical constant of 5. Eqs. (1) and (2) in our manuscript are derived from Eqs. (10) and 14, respectively from [39]. Please note that Eq. (14) is equivalent to Eq. (7) from Moynihan et. al. [57]. b.) Example thermogram of 100% trehalose with glass transition onset, glass transition offset, and 1st derivative with respect to temperature (green) shown. c) Glass former fragility (m-index) values for disaccharide-glycerol mixtures organized by disaccharide. d) Correlation plot of glass former fragility (m-index) versus protection for all disaccharide-glycerol mixtures (red = maltose, green = sucrose, blue = trehalose). Statistics calculated using a one-way ANOVA and Tukey post-hoc test: p -value ≤ 0.05 (*), p -value ≤ 0.01 (**), p -value ≤ 0.001 (***) pairwise relationships not shown are not significant, error bars represent 95% CI.

viscosity [56]. Crowley and Zografi later applied the assumption that the activation enthalpy for viscosity is equivalent to the activation enthalpy of structural relaxation at T_g [39]. Utilizing this method glass former fragility (m-index) was calculated from DSC thermogram outputs (Fig. 4a and b) [39]. The fragility of glass forming solutions composed from pure disaccharides was lowest for maltose, followed by sucrose, and finally trehalose (Fig. 4c). For maltose samples, glass former fragility steadily increased with each addition of glycerol but did not vary significantly (Fig. 4c). However, for both sucrose and trehalose small additions of glycerol (up to 5 and 2.5%, respectively) decreased glass former fragility (Fig. 4c). However, greater additions of glycerol to both trehalose and sucrose increased glass former fragility (Fig. 4c). For sucrose, variations in glass former fragility were not significant. On the other hand, after the addition of 10% glycerol, trehalose saw significant increases in glass former fragility (Fig. 4c).

Next, we assessed the relationship between the fragility of our glass forming mixtures and protection of LDH. For maltose and trehalose, the trend between enzyme-protection and glass former fragility was negative, while for sucrose this trend was slightly positive. As observed previously, the strength of these trends varied dramatically between sugars, as this correlative analysis produced an R^2 value of 0.64 for maltose (Figs. 4d and S5g), an R^2 value of 0.032 for sucrose (Figs. 4d and S5h), and R^2 value of 0.77 for trehalose (Figs. 4d and S5i). This indicates that for maltose and trehalose, observed variation in glass former fragility can explain some of the enzyme-protective capacity of the desiccated mixtures (Fig. 4d).

It was observed that the glass forming fragility of 100% sucrose was potentially behaving differently from mixtures containing glycerol additions. It was supposed that this datapoint might be 'dragging' the relationship between the glass former fragility of our glass forming mixtures and protection of LDH upward into a positive correlation. A correlation was calculated excluding the pure (100%) disaccharide samples from the mixtures (97.5–87.5%). This exclusion of the pure disaccharide samples greatly improved the correlation value of the sucrose mixtures, from an R^2 value of 0.032 to an R^2 value of 0.41 (Fig. S8c and d). Interestingly, the correlation for the trehalose-glycerol mixtures was also improved by excluding the 100% disaccharide samples, from an R^2 value of 0.77 to an R^2 value of 0.83 (Fig. S8e and S8f). However, the correlation for the maltose-glycerol mixtures was made worse, from an R^2 value of 0.64 to an R^2 value of 0.57 (Fig. S8a and b). These results suggest that while pure disaccharides may behave in substantively different ways than disaccharide-glycerol mixtures, this is not always the case.

4.6. Increased T_g and decreased glass former fragility are characteristics of anhydrobiotic life-stages

We wondered if the correlation between material properties, such as minimal glass former fragility or anti-plasticization, and enzyme-protection that we observed *in vitro* appeared to carry over to organismal systems. Here we considered reduced glass former fragility and anti-plasticization, but not water retention, since water content has previously been observed to influence glass-like properties [26,58–60]. This influence is observed in our samples, where for example in maltose-glycerol mixtures, we see a very strong correlation ($R^2 = 0.989$) between water content and increased T_g (Fig. S2a).

We measured the T_g and m-indexes of three desiccation tolerant organisms when in a desiccation sensitive life stage or a desiccation tolerant life stage. The three desiccation tolerant organisms we selected for this study were: the brine shrimp *Artemia franciscana*, the nematode worm *Caenorhabditis elegans*, and yeast *Saccharomyces cerevisiae* (Fig. 5a). Each of the selected organisms is known to accumulate protective disaccharides and polyols during drying [34,36,61–63].

For *A. franciscana*, there was a significant increase in T_g offset between the adult (desiccation-sensitive life stage) and cyst (desiccation-tolerant life stage) (Fig. 5b). For *S. cerevisiae*, we saw a statistical

increase between the exponential phase (desiccation sensitive life stage) and stationary phase (desiccation tolerant life stage) when measuring T_g offset (Fig. 5b). And finally, for *C. elegans*, we observed complex thermograms containing more than one obvious glass transition. Specifically, we observed two glass transitions on the thermograms of non-dauer worms and three glass transitions on the thermograms of dauer pre-conditioned worms. The two glass transitions from non-dauer worms were paired with the first two glass transitions from the dauer pre-conditioned worms based on similarities in glass transition temperature. The trend of desiccation tolerant life stages possessing a significantly higher T_g offset in comparison to the desiccation sensitive life stage continues across all three ranges of potential glass transitions for the *C. elegans* non-dauer and dauer pre-conditioned worms (Fig. 5b).

Thus, for all three of the desiccation tolerant organisms tested we observe significantly increased T_g in the anhydrobiotic state, indicating that increased T_g is a hallmark of some desiccation-tolerant organismal systems.

Next, we evaluated our three selected organisms' life stages for changes in m-index (glass former fragility). For *A. franciscana*, we saw significant decrease between the adult (desiccation sensitive life stage) and cyst (desiccation tolerant life stage) when measuring m-index (Fig. 5c). For *S. cerevisiae*, we also saw a significant decrease between the exponential phase (desiccation-sensitive life stage) and stationary phase (desiccation-tolerant life stage) when measuring m-index (Fig. 5c). For *C. elegans*, we saw a significant decrease in glass former fragility between the non-dauer (desiccation sensitive life stage) and dauer pre-conditioned (desiccation tolerant life stage) across all three glass transitions.

Taken together, these results indicate that while *in vitro* simple mixtures of protectants vary widely in what properties correlate with protection in the vitrified state, *in vivo* both increased glass transition temperature and decreased glass former fragility are good indicators of survival in the dry state.

5. Discussion and conclusions

Since its conception, the vitrification hypothesis has provided a compelling possible explanation as to how anhydrobiotic organisms preserve their cells and cellular components during desiccation [1,5–7]. However, while vitrification is considered necessary for desiccation tolerance it is not sufficient [5,9]. This implies that there is some property of a glassy material that makes it more, or less, protective.

Here we have quantified the enzyme-protective capacity and material properties of 18 different vitrified systems, each composed of one of three different disaccharides (maltose, sucrose, and trehalose) formulated with varying amounts of glycerol (0–12.5%). We find that both enzyme-protective capacity and material properties of disaccharide-glycerol mixtures are modulated differently depending on the disaccharide used. Consistent with this, water retention (maltose), increased T_g (sucrose), and reduced glass fragility (trehalose) each correlated best with protection for a particular disaccharide. Interestingly, reduced glass former fragility and increased glass transition (anti-plasticization) is observed in desiccation-tolerant life stages of diverse organisms. Thus, individual protective properties observed for reductive enzyme systems appear to be used in combination *in vivo*.

5.1. Water retention, mechanisms of protection involving water, and water's effects on glass transition temperature and glass former fragility

Our results demonstrate that different disaccharide-glycerol mixtures contain different quantities of water. Furthermore, the relationship between water content and protection also varies between disaccharide-glycerol mixtures. Maltose-glycerol mixtures display a significant positive correlation between water content and enzyme protection, while the sucrose-glycerol and trehalose-glycerol mixtures show only mildly positive correlations ($R^2 = 0.13$ and 0.28, respectively).

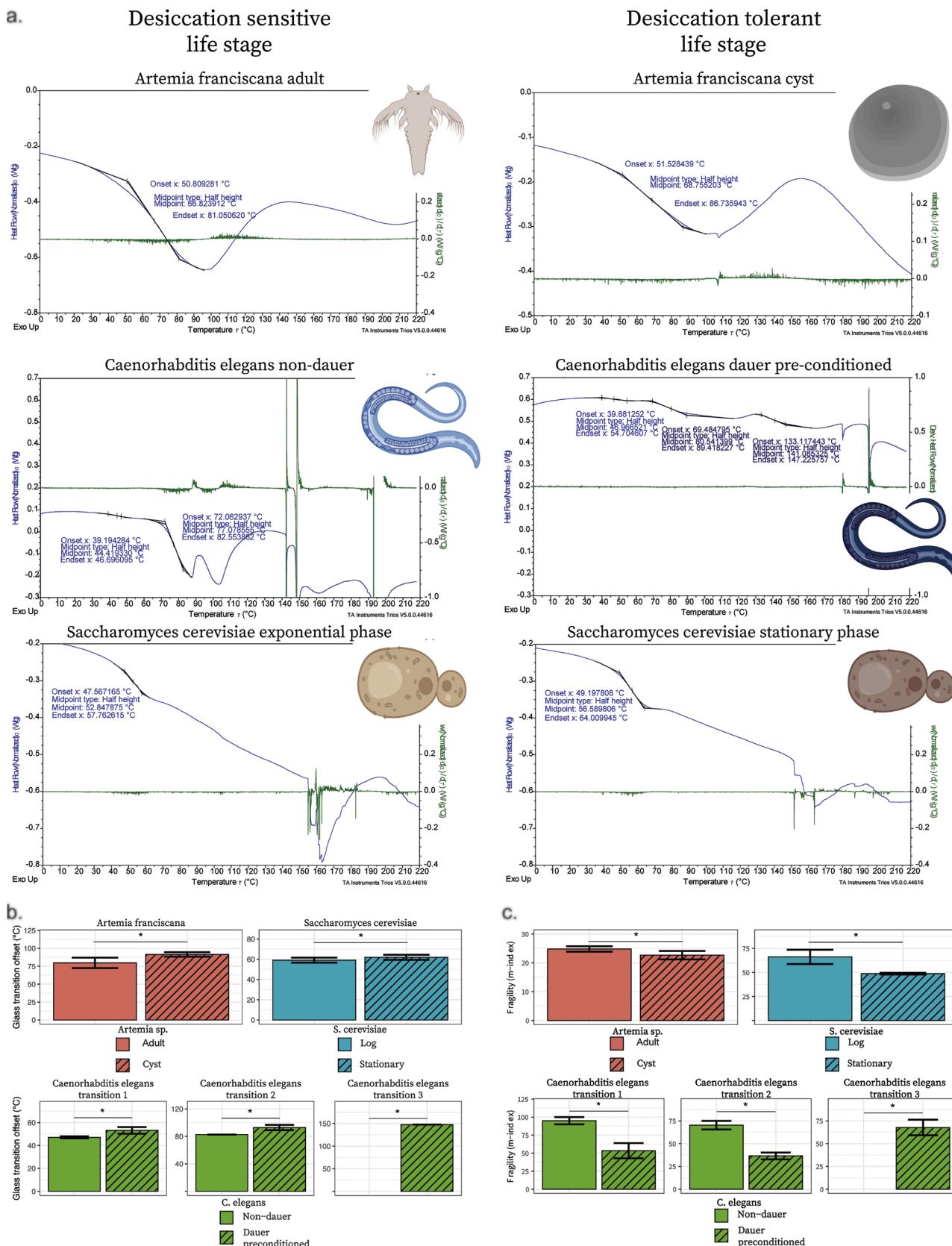


Fig. 5. *In vivo* glass transition temperature is increased, and glass former fragility reduced in desiccation-tolerant *versus* sensitive life stages of diverse anhydrobiotic organisms. (a) Selected organisms for *in vivo* assays with example DSC thermogram outputs. (b) Glass transition offset values for selected desiccation tolerant organisms organized by organism and life stage. (c) Glass former fragility (m-index) values for selected desiccation tolerant organisms organized by organism and life stage. Statistics calculated using T-test: p-value ≤ 0.05 (*), pairwise relationships not shown are not significant, error bars represent 95% CI.

The water content of a glassy material could affect enzyme-protection through several mechanisms. Loss of water during dehydration can lead to a loss of important, stabilizing hydrogen bonds, which help to maintain protein folding. While theories, such as the water replacement hypothesis, which propose mechanisms by which this loss of this hydrogen bond network can be dealt with, others such as the water entrapment [13–17], preferential exclusion/hydration hypothesis [16,18], or anchorage hypothesis [19–22] offer up mechanisms by which residual water can be utilized to provide protection even at low levels.

The water entrapment hypothesis posits that a protectant which has a strong affinity for water but is preferentially excluded from client molecules could help to coordinate small amounts of residual water into proximity with desiccation-sensitive material (e.g., proteins, membrane, etc.). The effect would be to entrap and increase the local concentration of water around these desiccation-sensitive molecules which could help to maintain the hydrogen bond network needed for integrity. Conversely, the preferential exclusion hypothesis posits that a protectant which preferentially interacts with itself, to the exclusion of both water and client molecules, could in effect act as a space filling molecule. In this capacity, the protectant would reduce the overall accessible volume within the cell increasing the effective concentration of water and forcing water molecules into proximity with desiccation-sensitive molecules. Finally, the anchorage hypothesis posits that client molecules interact with the water-protectant matrix, and this interaction reduces the likelihood of protein unfolding since unfolding would have to lead to a reordering of the water-protectant matrix.

Beyond water serving directly in the stabilization of biomolecules via the formation of a hydrogen bond network, water can also serve as an important plasticizing agent of biological and hydrophilic materials. This means that increasing water content in a vitrified material typically leads to a decrease in T_g , which is also considered to reduce protection. However, while this is generally true, there have been reports of *bona fide* anti-plasticization effects of water [59]. Here we observe that water content of maltose-, sucrose-, and trehalose-glycerol glasses has varying degrees of a strong plasticizing effect on the glass transition. The only negative correlation between water content and decreased T_g was observed for the sucrose-glycerol mixtures ($R^2 = 0.44$, Fig. S2c). In trehalose-glycerol mixtures, there was essentially no correlation between water content and decreased T_g ($R^2 = 0.088$, Fig. S2e). Finally, in maltose-glycerol mixtures, rather than seeing water correlate with decreased T_g , surprisingly we observe a strong correlation between water content and increased T_g ($R^2 = 0.94$, Fig. S2a). This again showcases how each of the disaccharides, when in a desiccated disaccharide-glycerol mixture, displays disparate changes in material properties.

When instead considering the relationship between water content and glass former fragility, we again observe differing results by disaccharide-glycerol mixture. Here we see that the desiccated maltose-glycerol mixtures have a strong negatively correlated relationship ($R^2 = 0.74$, Fig. S2b), sucrose has a strong positively correlated relationship ($R^2 = 0.88$, Fig. S2d), and trehalose has a weak positively correlated relationship ($R^2 = 0.29$, Fig. S2f).

These results might lead one to believe that at the water contents examined here (< 11%) water retention might be a potential predictor of enzyme protection capacity. However, water content also seems to influence other material properties of the vitrified system in a non-stereotyped fashion. For example, increasing water content in maltose-glycerol mixtures strongly correlates with reduced glass former fragility, while in sucrose and trehalose-glycerol mixtures increasing water content increases glass former fragility. Thus, we conclude that water content itself is not a good predictor of desiccation tolerance nor of other properties of a vitrified system.

5.2. The enzyme-protective capacity of sucrose-glycerol mixtures was most influenced by increases and decreases in glass transition temperature

Glass transition temperature (T_g) is the temperature at which a hard glassy material will begin to transition into a rubbery solid [64]. Fig. 3a is a schematic illustration of how increases or decreases in the T_g are captured and visualized on a thermogram. The relationship of increases or decreases in T_g with protection varied between different disaccharide-glycerol mixtures but was most important for the sucrose-glycerol mixtures. While we observed that only maltose-glycerol mixtures had a strong relationship between increased T_g and protection (Fig. 3d), in the sucrose-glycerol mixtures we see that addition of glycerol results in an increase in the T_g when added up to 5%, but then further addition of glycerol causes a significant decrease in the T_g . Furthermore, the influence of glycerol on the T_g of sucrose-glycerol mixtures correlates weakly with protection but is the highest correlation with respect to enzyme-protection (Figs. 2d, 3d, and 4d). Finally, in the trehalose-glycerol mixtures we see at first a significant decrease coinciding with the first addition of glycerol (2.5%) and then no significant variation in the glass transition midpoint until the addition of significantly more (12.25%) glycerol where it decreases the T_g , and the small changes in T_g that are observed show no correlation with protection.

Thus, while increases in T_g are predictive of the protection conferred by maltose-glycerol mixtures, and to a lesser extent sucrose-glycerol, this predictive capacity does not extend to all vitrified systems.

5.3. The difference between the experimental temperature of enzyme protection and glass transition temperature did not correlate with protection

The closer a glassy protectant is to its T_g , the more molecular motion should be introduced, which could result in a loss of protective capacity. However, we observed that the difference in the experimental temperature at which LDH assays were conducted and the $T_{exp}-T_g$ of a protective glass did not correlate significantly with enzyme-protective capacity (S6). This does not mean that as a protective glass approaches its T_g that it does not lose enzyme-protective capacity, since it is possible that the experimental temperatures used here were still sufficiently lower than T_g to negatively impact enzyme-protection. Further studies where the experimental temperature is brought much closer to T_g could be insightful in this regard.

5.4. The relationship between T_g and protection varies dramatically depending on whether onset, midpoint, or offset glass transition temperatures are considered

When referring to the glass transition or the material properties based on the glass transition, the standard measure is to use the glass transition midpoint. This value is inherently influenced by both the glass transition onset and offset temperatures (Figs. 3c, S3a, and S3b) [65]. The glass transition offset temperature is representative of the point at which the ‘glassy’ nature of a vitreous system is finally overcome (Fig. S3b). By contrast the glass transition onset is only representative of the start of the transition of a ‘glassy’ state to a rubber-like solid (Fig. S3a).

In evaluating the relationship between T_g and protection, we observed a dramatic variation in this correlation depending on whether we used the onset, midpoint, or offset glass transition temperature (Fig. S4). Here we have reported midpoint T_g as is convention, but also have included correlations between protection and onset/offset T_g (Fig. S4).

5.5. Disaccharide-glycerol mixtures with similar concentrations produce fragilities of differing glass former fragility (m-index)

The fragility of different glass former mixtures varied by disaccharide. Only maltose and trehalose showed evidence of a relationship between glass former fragility and enzyme protection capacity. Interestingly, when comparing different disaccharide-glycerol systems that provide similar enzyme-protective capacity, those mixtures did not necessarily produce glasses with similar glass former fragility (m-index) measurements. For example, when comparing the 97.5% maltose, 97.5% sucrose, and 92.5% trehalose disaccharide-glycerol mixtures that provide approximately 30% enzyme protection during desiccation (30.21%, 30.26, and 36.72% respectively), those mixtures produced had a glass former fragility (m-index) of 39.15, 74.01, and 151.81 respectively (Table 1). In addition, when comparing the most enzyme-protective mixtures for each disaccharide-glycerol mixture, 100% maltose, 100% sucrose and 97.5% trehalose, only trehalose produced the lowest glass former fragility (m-index) measurement. This indicates that each disaccharide-glycerol system produced glass former fragility patterns that are only comparable within that system and not between different disaccharide-glycerol systems. Again, just as with water retention and shifts in the T_g , we see that the addition of glycerol induces different degrees of glass former fragility and that this property is a good indicator of protection for some sugar glasses, but not for others.

5.6. Both change in T_g and glass former fragility are hallmarks of organismal desiccation tolerance

The results from our *in vitro* experiments, while novel on their own, beg the question of whether these findings apply to whole desiccation

tolerant organisms. In particular, among the organisms studied, *A. franciscana*, *C. elegans*, and *S. cerevisiae*, trehalose is a major carbohydrate which is accumulated during desiccation and functionally thought to drive desiccation tolerance [4,33–36,63,66,67]. In *A. franciscana* up to 15% of the dry weight of the cysts is attributed to trehalose accumulation [66,67]. In *C. elegans* dauer larva that undergo pre-conditioning for four days will have accumulated up to approximately 375 µg of trehalose per mg of total protein [63,68,69]. Finally, in *S. cerevisiae* stationary phase yeast can accumulate up to 15% of the total dry cell mass as trehalose [33,34]. Of course, these organisms also accumulate a number of other disaccharides, diverse metabolites and proteins during drying, all of which likely combine to contribute to glassy properties.

Examining these three different organisms in both a desiccation-tolerant and -sensitive life stage we observe that both increased T_g and reduced m-index (glass former fragility) are hallmarks of successful anhydrobiosis. Specifically, when considering changes in T_g , we see that for each organism, *A. franciscana*, *C. elegans*, and *S. cerevisiae*, there is a statistically significant increase in T_g when comparing the desiccation-sensitive life stage to the desiccation-tolerant life stage. When considering the impact of glass former fragility, we yet again see that for each organism, *A. franciscana*, *C. elegans*, and *S. cerevisiae*, there is a statistically significant decrease in glass former fragility when comparing the desiccation-tolerant life stage to the desiccation-sensitive life stage.

Fig. 6 provides an overview of what we envision is occurring to enable changes in the T_g (Fig. 6a) and glass former fragility (Fig. 6b and c) to be protective during desiccation. In Fig. 6a we depict a glass with a decreased T_g as one with relatively few and/or weak bonds, which do not effectively slow down molecular motions leading to the destabilization and/or aggregation of embedded clients over time. In contrast, a

Table 1

In vitro measurement results organized by disaccharide-glycerol content and by material property, error values represent 95% CI.

Disaccharide	Disaccharide content (% weight)	Glycerol content (% weight)	LDH protection (%)	Water content (%)	Tg onset (°C)	Tg offset (°C)	Tg midpoint (°C)	Calculated m-index	TgΔ (Texp - Tg)
Maltose	100.0	0.0	33.08 +/- (4.29)	4.35 +/- (0.12)	67.91 +/- (4.96)	86.50 +/- (3.31)	77.16 +/- (0.34)	44.25 +/- (19.30)	-55.16 +/- (0.34)
	97.5	2.5	30.21 +/- (4.77)	3.90 +/- (0.31)	63.98 +/- (1.22)	83.83 +/- (0.66)	73.58 +/- (0.71)	39.15 +/- 3.62	-51.58 +/- (0.71)
	95.0	5.0	28.95 +/- (5.08)	2.90 +/- (0.62)	60.03 +/- (11.24)	73.99 +/- (5.75)	66.50 +/- (8.06)	56.36 +/- (23.05)	-44.50 +/- (8.06)
	92.5	7.5	25.73 +/- (1.45)	2.53 +/- (0.29)	54.58 +/- (2.25)	66.62 +/- (0.03)	61.13 +/- (1.53)	61.85 +/- (11.44)	-39.13 +/- (1.53)
	90.0	10.0	22.32 +/- (1.84)	1.89 +/- (0.09)	55.83 +/- (2.62)	70.22 +/- (3.18)	61.98 +/- (3.40)	51.83 +/- 1.57	-39.98 +/- (3.40)
	87.5	12.5	17.56 +/- (3.51)	0.91 +/- (0.15)	51.44 +/- (6.65)	62.53 +/- (0.33)	56.06 +/- (3.11)	71.77 +/- (40.95)	-34.06 +/- (3.11)
Sucrose	100.0	0.0	52.28 +/- (15.69)	5.35 +/- (1.5)	46.80 +/- (2.09)	53.46 +/- (2.79)	50.13 +/- (2.38)	108.17 +/- (17.91)	-28.13 +/- (2.38)
	97.5	2.5	30.26 +/- (9.79)	3.84 +/- (0.96)	48.21 +/- (0.08)	58.10 +/- (0.08)	53.16 +/- (2.19)	74.01 +/- (15.33)	-31.16 +/- (1.39)
	95.0	5.0	18.22 +/- (4.21)	3.27 +/- (0.05)	47.99 +/- (2.63)	59.70 +/- (2.11)	53.84 +/- (2.37)	61.75 +/- 3.13	-31.84 +/- (2.91)
	92.5	7.5	16.70 +/- (7.99)	3.99 +/- (0.47)	43.19 +/- (1.97)	55.30 +/- (4.59)	49.25 +/- (2.58)	65.30 +/- (31.21)	-27.25 +/- (2.58)
	90.0	10.0	10.33 +/- (0.69)	4.22 +/- (0.31)	44.10 +/- (3.76)	52.14 +/- (3.52)	48.12 +/- (3.64)	87.90 +/- 3.58	-26.12 +/- (4.46)
	87.5	12.5	7.03 +/- (1.68)	5.05 +/- (0.77)	39.56 +/- (1.89)	46.32 +/- (1.65)	42.94 +/- (0.67)	114.33 +/- (47.52)	-20.94 +/- (0.67)
Trehalose	100.0	0.0	30.52 +/- (6.24)	8.18 +/- (0.68)	51.27 +/- (0.23)	55.32 +/- (0.58)	53.14 +/- (0.03)	178.62 +/- (30.04)	-31.14 +/- (0.03)
	97.5	2.5	52.17 +/- (4.66)	7.50 +/- (0.25)	43.14 +/- (2.35)	50.93 +/- (1.60)	47.52 +/- (1.34)	90.79 +/- 9.26	-25.52 +/- (1.34)
	95.0	5.0	42.65 +/- (1.67)	8.55 +/- (1.99)	43.89 +/- (0.3)	51.24 +/- (1.91)	47.25 +/- (0.74)	99.58 +/- (27.97)	-25.25 +/- (0.74)
	92.5	7.5	36.72 +/- (4.34)	6.01 +/- (0.38)	46.23 +/- (1.44)	51.15 +/- (1.86)	48.67 +/- (1.64)	151.81 +/- (47.27)	-26.67 +/- (1.64)
	90.0	10.0	23.44 +/- (1.05)	5.27 +/- (0.63)	46.78 +/- (2.01)	48.52 +/- (1.97)	47.33 +/- (1.95)	416.47 +/- (113.38)	-25.33 +/- (1.95)
	87.5	12.5	18.46 +/- (1.57)	6.45 +/- (1.52)	30.85 +/- (3.27)	32.85 +/- (3.32)	31.83 +/- (3.31)	334.52 +/- (44.82)	-9.83 +/- (3.31)

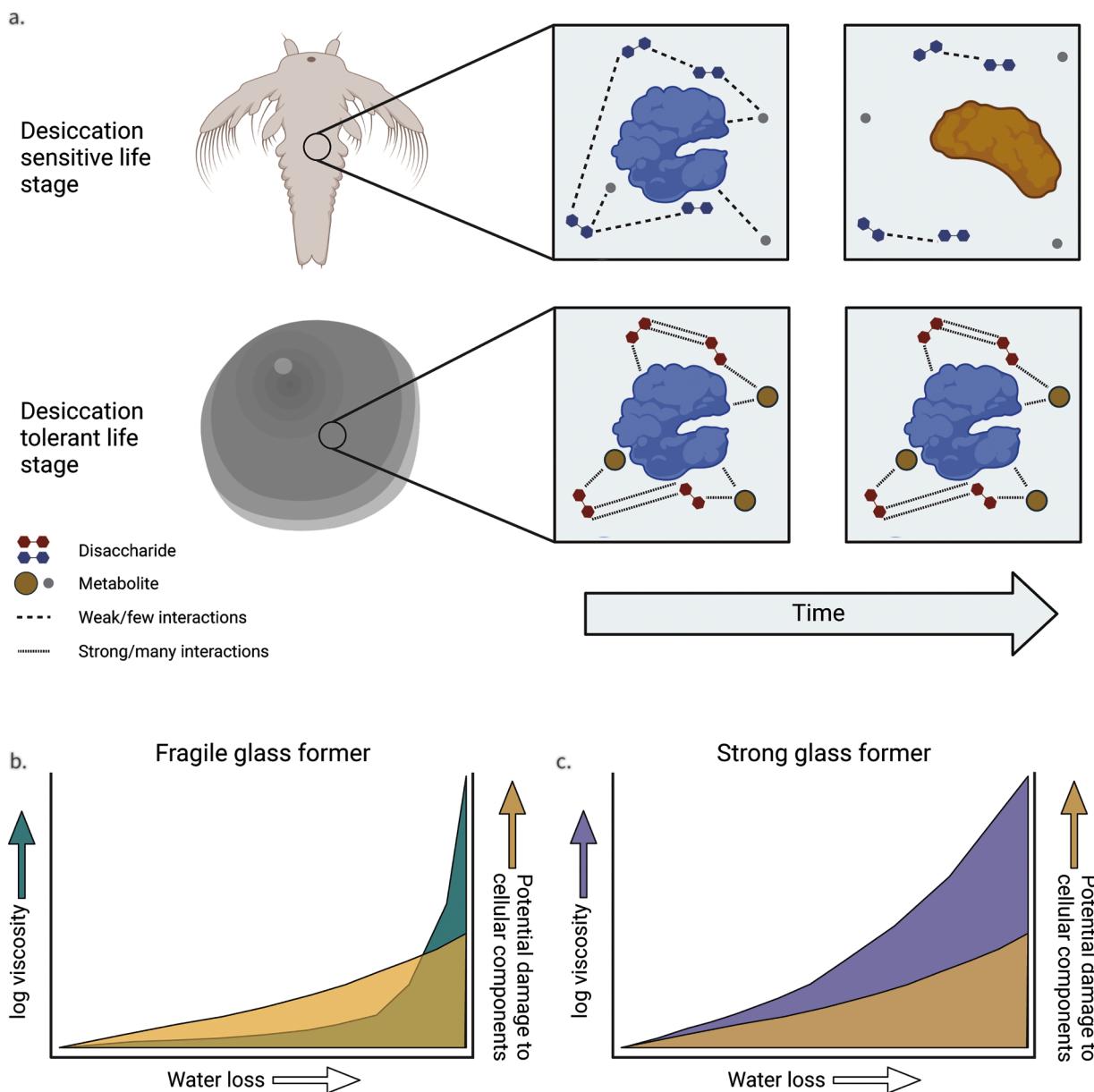


Fig. 6. Model of protection conferred by increased glass transition temperature and reduced glass former fragility. (a) Schematic representation of the potential mechanism of protection conferred by increasing glass transition temperature. (b) Schematic representation of the potential mechanism underlying how damage is accrued in fragile glass former mixtures. (c) Schematic of the potential mechanism underlying how damage is prevented in strong glass former systems.

glass with an increased T_g is one with increased and/or stronger bonds leading to reduction in molecular motion and an increase in stability of a cell over time. This is in line with previous suggestions that inducing a super viscous (glassy) state is stabilizing because to unfold a protein would need to displace the embedding media [5,14]. In this light, increased T_g would be protective as materials will become more fluid as the temperature they are stored at approaches T_g .

Fig. 6b illustrates what we envision is happening when a glass forming mixture forms a weak glass. As water is lost, the weak glass forming ability of the mixture does not become viscous (green) soon enough to prevent drying induced damage (tan). Specifically, a weak glass former material will only begin gaining sufficient viscosity to confer protection after significant water loss. Conversely, Fig. 6c illustrates how a strong glass forming mixture starts to gain viscosity (purple) much earlier in the desiccation process compared to weaker glass forming materials. This steady increase in viscosity allows for slowing detrimental perturbations (tan) that manifest early during drying. One

such perturbation might be the unfolding and aggregation of proteins, while another might be the fusion of membranes. Furthermore, previous studies have shown that volume loss during drying is associated with loss of cellular viability. Fragile glass formers, owing to their general lack of viscosity, might allow for more rapid cell shrinkage during drying. In contrast, strong glass formers might provide resistance to cell shrinkage via the induction of a super-viscous state early during the drying process [70–73]. This model implies to some degree that it is the drying process, in addition to being in a dry state, that must be protected against.

Previous studies pertaining to desiccation tolerant organisms have established that these organisms use a variety of cellular metabolites to survive the process of desiccation, anhydrobiosis, and even rehydration [34,63,66–69,74,75]. In particular, it should be noted that the production of these cellular metabolites can vary significantly between different desiccation tolerant organisms [4,30,76–78]. When considering the organisms investigated within this study, *A. franciscana*, *C.*

elegans, and *S. cerevisiae*, each organism produces a variety of metabolites consisting of proteins and carbohydrates which are necessary for surviving desiccation [4,33,34,63,66–69,74,75]. However, as seen in other desiccation tolerant organisms, such as rotifers, plants, and cyanobacteria, an array of disaccharides may be used instead of trehalose [79,80]. While all three organisms studied here express only trehalose, it has been demonstrated that the use of externally added disaccharide, such as sucrose or maltose, can improve an organisms' ability to survive desiccation [81]. For instance, Tapia et. al. demonstrated that uptake of maltose by way of media was able to provide desiccation protection [34]. In fact, it has been noted that sucrose, maltose, and trehalose are known to be expressed in different desiccation tolerant plant species [30,76,77,82]. In addition to disaccharides, many desiccation tolerant organisms also produce an assortment of intrinsically disordered proteins (IDPs). In desiccation tolerant organisms, some of these IDPs are expressed in large numbers to survive desiccation stresses, while others are expressed in much smaller numbers [4,7,83,84]. Nguyen et. al. demonstrated that an IDP (CAHS D) and trehalose can work synergistically to provide greater protection to desiccation stress than either could individually. This finding in particular is encouraging as it supports the concept we propose: that desiccation tolerant organisms can take advantage of different methods of desiccation protection to better survive the full continuum of stresses during desiccation [83].

While previous studies have examined the glass former fragility of seeds in relationship to their desiccation tolerance, to our knowledge this is the first study examining changes in T_g and glass former fragility in animal and fungal systems [3,9,85–88]. These comparative organismal studies show a stark contrast to our *in vitro* data, in that rather than a single material property correlating with protection, it appears that in living anhydrobiotic systems both increased T_g and reduced glass former fragility are generally increased. This may be due to the nature of the *in vitro* systems being simple, or less complex, in their interactions while the *in vivo* studies are by their nature much more complex, both in their material makeup and interactions.

These results hint at the fact that living systems likely make use of multiple mediators of desiccation tolerance to produce protective glasses. Indeed, an emerging paradigm in the anhydrobiosis field is that beyond disaccharides, other molecules, such as intrinsically disordered proteins play vital roles in preserving biological function in the solid state.

Our study advances our understanding of what properties of a vitrified system promote desiccation tolerance and the phenomenon of anhydrobiosis both *in vitro* and *in vivo*. A deeper understanding of natural desiccation tolerance promises to provide avenues for pursuing real world applications such as biobanking of seeds and tissues, stabilization of pharmaceuticals, and the generation of stress tolerant crops.

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CRediT authorship contribution statement

John F. Ramirez: Conceptualization, Data curation, Formal

analysis, Investigation, Methodology, Visualization, Writing – original draft, Writing – review & editing. **U.G.V.S.S. Kumara:** Investigation, Writing – review & editing, Data curation, Methodology. **Navamoney Arulsamy:** Investigation, Writing – review & editing, Writing – original draft. **Thomas C. Boothby:** Conceptualization, Funding acquisition, Project administration, Supervision, Writing – original draft, Writing – review & editing, Methodology, Visualization.

Declaration of competing interest

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: Thomas Boothby reports financial support was provided by National Science Foundation. If there are other authors, they declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Data availability

All raw data and scripts used in this paper are available at https://figshare.com/articles/dataset/Ramirez_et_al_Supplementary_files_Data_and_Scripts_24964338.

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Supplementary materials

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References

- [1] J.H. Crowe, F.A. Hoekstra, L.M. Crowe, Anhydrobiosis, *Annu. Rev. Physiol.* 54 (1992) 579–599.
- [2] C. Hesgrave, T.C. Boothby, The biology of tardigrade disordered proteins in extreme stress tolerance, *Cell Commun. Signal.* 18 (2020) 178.
- [3] D. Ballesteros, H.W. Pritchard, C. Walters, Dry architecture: towards the understanding of the variation of longevity in desiccation-tolerant germplasm, *Seed Sci. Res.* 30 (2020) 142–155.
- [4] J.D. Hibshman, J.S. Clegg, B. Goldstein, Mechanisms of desiccation tolerance: themes and variations in brine shrimp, roundworms, and tardigrades, *Front. Physiol.* 11 (2020) 592016.
- [5] J.H. Crowe, J.F. Carpenter, L.M. Crowe, The role of vitrification in anhydrobiosis, *Annu. Rev. Physiol.* 60 (1998) 73–103.
- [6] M. Sakurai, T. Furuki, K.-I. Akao, D. Tanaka, Y. Nakahara, T. Kikawada, M. Watanabe, T. Okuda, Vitrification is essential for anhydrobiosis in an African chironomid, *Polypedilum vanderplanki*, *Proc. Natl. Acad. Sci. U. S. A.* 105 (2008) 5093–5098.
- [7] T.C. Boothby, H. Tapia, A.H. Brozena, S. Piszkiewicz, A.E. Smith, I. Giovannini, L. Rebecchi, G.J. Pielak, D. Koshland, B. Goldstein, Tardigrades use intrinsically disordered proteins to survive desiccation, *Mol. Cell* 65 (2017) 975–984, e5.
- [8] T.C. Boothby, Water content influences the vitrified properties of CAHS proteins, *Mol. Cell* 81 (2021) 411–413.
- [9] D. Ballesteros, C. Walters, Solid-state biology and seed longevity: a mechanical analysis of glasses in pea and soybean embryonic axes, *Front. Plant Sci.* 10 (2019) 920.
- [10] L. Weng, G.D. Elliott, Local minimum in fragility for trehalose/glycerol mixtures: implications for biopharmaceutical stabilization, *J. Phys. Chem. B* 119 (2015) 6820–6827.
- [11] J.C. Wright, Desiccation tolerance and water-retentive mechanisms in tardigrades, *J. Exp. Biol.* 142 (1989) 267–292.

[12] F.A. Hoekstra, E.A. Golovina, J. Buitink, Mechanisms of plant desiccation tolerance, *Trends Plant Sci.* 6 (2001) 431–438.

[13] P.S. Belton, A.M. Gil, IR and Raman spectroscopic studies of the interaction of trehalose with hen egg white lysozyme, *Biopolymers* 34 (1994) 957–961.

[14] G.I. Olgenblum, L. Sapir, D. Harries, Properties of aqueous trehalose mixtures: glass transition and hydrogen bonding, *J. Chem. Theory Comput.* 16 (2020) 1249–1262.

[15] R. Politi, D. Harries, Enthalpically driven peptide stabilization by protective osmolytes, *Chem. Commun.* 46 (2010) 6449–6451.

[16] S.N. Timasheff, Protein hydration, thermodynamic binding, and preferential hydration, *Biochemistry* 41 (2002) 13473–13482.

[17] T.Y. Lin, S.N. Timasheff, On the role of surface tension in the stabilization of globular proteins, *Protein Sci.* 5 (1996) 372–381.

[18] S. Shimizu, D.J. Smith, Preferential hydration and the exclusion of cosolvents from protein surfaces, *J. Chem. Phys.* 121 (2004) 1148–1154.

[19] G. Bellavia, S. Giuffrida, G. Cottone, A. Cupane, L. Cordone, Protein thermal denaturation and matrix glass transition in different protein-trehalose-water systems, *J. Phys. Chem. B* 115 (2011) 6340–6346.

[20] F. Francia, M. Dezi, A. Mallardi, G. Palazzo, L. Cordone, G. Venturoli, Protein–Matrix coupling/uncoupling in “dry” systems of photosynthetic reaction center embedded in trehalose/sucrose: the origin of trehalose peculiarity, *J. Am. Chem. Soc.* 130 (2008) 10240–10246.

[21] A. Lerbret, F. Affouard, Molecular packing, hydrogen bonding, and fast dynamics in lysozyme/trehalose/glycerol and trehalose/glycerol glasses at low hydration, *J. Phys. Chem. B* 121 (2017) 9437–9451.

[22] J.C. Lee, S.N. Timasheff, The stabilization of proteins by sucrose, *J. Biol. Chem.* 256 (1981) 7193–7201.

[23] C.A. Angell, The old problems of glass and the glass transition, and the many new twists, *Proc. Natl. Acad. Sci. U. S. A.* 92 (1995) 6675–6682.

[24] C.A. Angell, Entropy, fragility, “landscapes”, and the glass transition, in: *Complex Behaviour of Glassy Systems*, Springer, Berlin Heidelberg, 1997, pp. 1–21.

[25] K. Ito, C.T. Moynihan, C.A. Angell, Thermodynamic determination of fragility in liquids and a fragile-to-strong liquid transition in water, *Nature* 398 (1999) 492–495.

[26] E.H. Immergut, H.F. Mark, Principles of plasticization, in: *Plasticization and Plasticizer Processes*, American Chemical Society, 1965, pp. 1–26.

[27] A. Anopchenko, T. Psurek, D. VanderHart, J.F. Douglas, J. Obrzut, Dielectric study of the antiplasticization of trehalose by glycerol, *Phys. Rev. E* 74 (2006) 031501. *Stat. Nonlin. Soft Matter Phys.*

[28] T.C. Boothby, Mechanisms and evolution of resistance to environmental extremes in animals, *EvoDevo* 10 (2019) 30.

[29] C.A. Angell, Entropy, fragility, “landscapes”, and the glass transition. Complex Behaviour of Glassy Systems, *Proceedings of the XIV Sitges Conference Sitges*, Barcelona, Spain, 1996, pp. 1–21.

[30] C. Dinakar, D. Bartels, Desiccation tolerance in resurrection plants: new insights from transcriptome, proteome and metabolome analysis, *Front. Plant Sci.* 4 (2013) 482.

[31] T.S. Gechev, M. Benina, T. Obata, T. Tohge, N. Sujeeeth, I. Minkov, J. Hille, M.-R. Temanni, A.S. Marriott, E. Bergström, J. Thomas-Oates, C. Antonio, B. Mueller-Roeber, J.H.M. Schippers, A.R. Fernie, V. Toneva, Molecular mechanisms of desiccation tolerance in the resurrection glacial relic *Haberlea rhodopensis*, *Cell. Mol. Life Sci.* 70 (2013) 689–709.

[32] L.M. Crowe, Lessons from nature: the role of sugars in anhydrobiosis, *Comp. Biochem. Physiol. A Mol. Integr. Physiol.* 131 (2002) 505–513.

[33] H. Tapia, D.E. Koshland, Trehalose is a versatile and long-lived chaperone for desiccation tolerance, *Curr. Biol.* 24 (2014) 2758–2766.

[34] H. Tapia, L. Young, D. Fox, C.R. Bertoza, D. Koshland, Increasing intracellular trehalose is sufficient to confer desiccation tolerance to *Saccharomyces cerevisiae*, *Proc. Natl. Acad. Sci. U. S. A.* 112 (2015) 6122–6127.

[35] S.H. Loomis, K.A.C. Madin, Anhydrobiosis in nematodes: biosynthesis of trehalose, *J. Exp. Zool.* 211 (3) (1980) 311–320.

[36] C. Womersley, L. Smith, Anhydrobiosis in nematodes—I The role of glycerol myoinositol and trehalose during desiccation, *Comp. Biochem. Physiol. B Comp. Biochem.* 70 (1981) 579–586.

[37] C. Womersley, Biochemical and physiological aspects of anhydrobiosis, *Comp. Biochem. Physiol. B: Comp. Biochem.* 70 (1981) 669–678.

[38] M. Robert Michaud, J.B. Benoit, G. Lopez-Martinez, M.A. Elnitsky, R.E. Lee, D. L. Denlinger, Metabolomics reveals unique and shared metabolic changes in response to heat shock, freezing and desiccation in the Antarctic midge, *Belgica antarctica*, *J. Insect Physiol.* 54 (2008) 645–655.

[39] K.J. Crowley, G. Zografi, The use of thermal methods for predicting glass-former fragility, *Thermochim. Acta* 380 (2001) 79–93.

[40] J.A. Lewis, J.T. Fleming, Basic culture methods, *Methods Cell Biol.* 48 (1995) 3–29.

[41] YPD media, *Cold Spring Harbor Protocols*, 2010, YPD media, 2010. *Protoc.db. rec12315*.

[42] T.E. Dirama, G.A. Carri, A.P. Sokolov, Role of hydrogen bonds in the fast dynamics of binary glasses of trehalose and glycerol: a molecular dynamics simulation study, *J. Chem. Phys.* 122 (2005) 114505.

[43] PubChem Compound Summary for CID 6255, Maltose, National Center for Biotechnology Information (Accessed 2023) <https://pubchem.ncbi.nlm.nih.gov/compound/Maltose>.

[44] PubChem Compound Summary for CID 5988, Sucrose, National Center for Biotechnology Information (Accessed 2023) <https://pubchem.ncbi.nlm.nih.gov/compound/Sucrose>.

[45] PubChem Compound Summary for CID 7427, Trehalose, National Center for Biotechnology Information (Accessed 2023) <https://pubchem.ncbi.nlm.nih.gov/compound/Trehalose>.

[46] A. Chauhan, Powder XRD technique and its applications in science and technology, *J. Anal. Bioanal. Tech.* 5 (2014).

[47] I.C. Madsen, N.V.Y. Scarlett, A. Kern, Description and survey of methodologies for the determination of amorphous content via X-ray powder diffraction, *Z. Krist.* 226 (2011) 944–955.

[48] S. Piszkiewicz, K.H. Gunn, O. Warmuth, A. Propst, A. Mehta, K.H. Nguyen, E. Kuhlman, A.J. Guseman, S.S. Stadtmiller, T.C. Boothby, S.B. Neher, G.J. Piekar, Protecting activity of desiccated enzymes, *Protein Sci.* 28 (2019) 941–951.

[49] K. Goyal, L.J. Walton, A. Tunnacliffe, LEA proteins prevent protein aggregation due to water stress, *Biochem. J.* 388 (2005) 151–157.

[50] M. Rizzuto, A. Mugica, M. Zubitur, D. Caretti, A.J. Müller, Plasticization and anti-plasticization effects caused by poly(lactide- ran -caprolactone) addition to double crystalline poly(l -lactide)/poly(ϵ -caprolactone) blends, *CrystEngComm* 18 (2016) 2014–2023.

[51] E. Luk, A.J. Sandoval, A. Cova, A.J. Müller, Anti-plasticization of cassava starch by complexing fatty acids, *Carbohydr. Polym.* 98 (2013) 659–664.

[52] Y. Chen, T. Tang, C. Ayrancı, Moisture-induced anti-plasticization of polylactic acid: experiments and modeling, *J. Appl. Polym. Sci.* 139 (2022) 52369.

[53] Y. Figueiroa, M. Guevara, A. Pérez, A. Cova, A.J. Sandoval, A.J. Müller, Effect of sugar addition on glass transition temperatures of cassava starch with low to intermediate moisture contents, *Carbohydr. Polym.* 146 (2016) 231–237.

[54] Y. Liu, A.K. Roy, A.A. Jones, P.T. Inglefield, P. Ogden, An NMR study of plasticization and antiplasticization of a polymeric glass, *Macromolecules* 23 (1990) 968–977.

[55] J. Tarique, S.M. Sapuan, A. Khalina, Effect of glycerol plasticizer loading on the physical, mechanical, thermal, and barrier properties of arrowroot (*Maranta arundinacea*) starch biopolymers, *Sci. Rep.* 11 (2021) 13900.

[56] C.T. Moynihan, Correlation between the width of the glass transition region and the temperature dependence of the viscosity of high-tg glasses, *J. Am. Ceram. Soc.* 76 (1993) 1081–1087.

[57] C.T. Moynihan, S.-K. Lee, M. Tatsumisago, T. Minami, Estimation of activation energies for structural relaxation and viscous flow from DTA and DSC experiments, *Thermochim. Acta* 280–281 (1996) 153–162.

[58] Y.I. Matveev, V.Y. Grinberg, V.B. Tolstoguzov, The plasticizing effect of water on proteins, polysaccharides and their mixtures Glassy state of biopolymers, food and seeds, *Food Hydrocollo.* 14 (2000) 425–437.

[59] G.N. Ruiz, M. Romanini, A. Hauptmann, T. Loerting, E. Shalaev, J.L. Tamarit, L. C. Pardo, R. Macovec, Genuine antiplasticizing effect of water on a glass-former drug, *Sci. Rep.* 7 (2017) 7470.

[60] I. Combarro Palacios, C. Olsson, C.S. Kamma-Lorger, J. Swenson, S. Cerveny, Motions of water and solutes-Slaving versus plasticization phenomena, *J. Chem. Phys.* 150 (2019) 124902.

[61] D. Calahan, M. Dunham, C. DeSevo, D.E. Koshland, Genetic analysis of desiccation tolerance in *Saccharomyces cerevisiae*, *Genetics* 189 (2011) 507–519.

[62] B. Janis, C. Belott, M.A. Menze, Role of Intrinsic disorder in animal desiccation tolerance, *Proteomics* 18 (2018) e1800067.

[63] C. Erkut, S. Penkov, H. Khesbak, D. Vorkel, J.M. Verbavatz, K. Fahmy, T. V. Kurzchalia, Trehalose renders the dauer larva of *Caenorhabditis elegans* resistant to extreme desiccation, *Curr. Biol.* 21 (2011) 1331–1336.

[64] J.C. Dyre, Colloquium: The glass transition and elastic models of glass-forming liquids, *Rev. Mod. Phys.* 78 (2006) 953–972.

[65] G. Roudaut, D. Simatos, D. Champion, E. Contreras-Lopez, M.L. Meste, Molecular mobility around the glass transition temperature: a mini review, *Innov. Food Sci. Emerg. Technol.* 5 (2004) 127–134.

[66] T.H. MacRae, Stress tolerance during diapause and quiescence of the brine shrimp, *Artemia*, *Cell Stress Chaperones* 21 (2016) 9–18.

[67] J.S. Clegg, The origin of trehalose and its significance during the formation of encysted dormant embryos of *artemia salina*, *Comp. Biochem. Physiol.* 14 (1965) 135–143.

[68] C. Erkut, A. Vasilj, S. Boland, B. Habermann, A. Shevchenko, T.V. Kurzchalia, Molecular strategies of the *Caenorhabditis elegans* dauer larva to survive extreme desiccation, *PLoS One* 8 (2013) e82473.

[69] C. Erkut, S. Penkov, K. Fahmy, T.V. Kurzchalia, How worms survive desiccation: trehalose pro water, *Worm* 1 (2012) 61–65.

[70] E. Anibal Disalvo, S.A. Simon, *Permeability and Stability of Lipid Bilayers*, CRC Press, 2017.

[71] H.T. Meryman, Freezing injury and its prevention in living cells, *Annu. Rev. Biophys. Bioeng.* 3 (1974) 341–363.

[72] J. Nadarajan, C. Walters, H.W. Pritchard, D. Ballesteros, L. Colville, Seed longevity—The evolution of knowledge and a conceptual framework, *Plants* 12 (2023).

[73] C. Walters, Orthodoxy, recalcitrance and in-between: describing variation in seed storage characteristics using threshold responses to water loss, *Planta* 242 (2015) 397–406.

[74] J. Francois, J.L. Parrou, Reserve carbohydrates metabolism in the yeast *Saccharomyces cerevisiae*, *FEMS Microbiol. Rev.* 25 (2001) 125–145.

[75] A. Zecic, I. Dhondt, B.P. Braeckman, Accumulation of glycogen and upregulation of LEA-1 in *C. elegans* daf-2(e1370) support stress resistance, not longevity, *Cells* 11 (2022).

[76] H.J. Dace, A.E. Adetunji, J.P. Moore, J.M. Farrant, H.W. Hilhorst, A review of the role of metabolites in vegetative desiccation tolerance of angiosperms, *Curr. Opin. Plant Biol.* 75 (2023) 102410.

[77] X. Xu, S. Legay, K. Sergeant, S. Zorzan, C.C. Leclercq, S. Charton, V. Giarola, X. Liu, D. Challabathula, J. Renault, J.-F. Hausman, D. Bartels, G. Guerrero, Molecular insights into plant desiccation tolerance: transcriptomics, proteomics and targeted metabolite profiling in *Craterostigma plantagineum*, *Plant J.* 107 (2021) 377–398.

[78] Q. Ren, R. Brenner, T.C. Boothby, Z. Zhang, Membrane and lipid metabolism plays an important role in desiccation resistance in the yeast *Saccharomyces cerevisiae*, *BMC Microbiol.* 20 (2020) 338.

[79] M. Potts, Desiccation tolerance: a simple process? *Trends Microbiol.* 9 (2001) 553–559.

[80] J. Lapinski, A. Tunnacliffe, Anhydrobiosis without trehalose in bdelloid rotifers, *FEBS Lett.* 553 (2003) 387–390.

[81] S.C. Hand, M.A. Menze, M. Toner, L. Boswell, D. Moore, LEA proteins during water stress: not just for plants anymore, *Annu. Rev. Physiol.* 73 (2011) 115–134.

[82] M.J. Oliver, L. Guo, D.C. Alexander, J.A. Ryals, B.W.M. Wone, J.C. Cushman, A sister group contrast using untargeted global metabolomic analysis delineates the biochemical regulation underlying desiccation tolerance in *Sporobolus stapfianus*, *Plant Cell* 23 (2011) 1231–1248.

[83] K. Nguyen, S. Kc, T. Gonzalez, H. Tapia, T.C. Boothby, Trehalose and tardigrade CAHS proteins work synergistically to promote desiccation tolerance, *Commun. Biol.* 5 (2022) 1046.

[84] J.D. Hibshman, B. Goldstein, LEA motifs promote desiccation tolerance *in vivo*, *BMC biol.* 19 (2021) 1–22.

[85] C. Walters, Temperature dependency of molecular mobility in preserved seeds, *Biophys. J.* 86 (2004) 1253–1258.

[86] D. Ballesteros, C. Walters, Detailed characterization of mechanical properties and molecular mobility within dry seed glasses: relevance to the physiology of dry biological systems, *Plant J.* 68 (2011) 607–619.

[87] C. Walters, Understanding the mechanisms and kinetics of seed aging, *Seed Sci. Res.* 8 (1998) 223–244.

[88] O. Leprince, C. Walters-Vertucci, A calorimetric study of the glass transition behaviors in axes of bean seeds with relevance to storage stability, *Plant Physiol.* 109 (1995) 1471–1481.