Lyophilized yeast powder for adjuvant free thermostable vaccine delivery

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Freeze-dried yeast powder for long-term storage of immunogen under the non-refrigerated condition

List of abbreviations: SEM, Scanning electron microscope; GVAP, Global Vaccine Action Plan; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2; COVID-19, coronavirus disease of 2019; APCs, antigen-presenting cells

Abstract

Thermolabile nature of commercially available vaccines necessitates their storage, transportation and dissemination under refrigerated condition. Maintenance of continuous cold chain at every step increases the final cost of vaccines. Any breach in the cold chain, even for a short duration results in the need to discard the vaccine. As a result, there is a pressing need for the development of thermostable vaccines. In this proof of concept study, we showed that *E. coli* curli-GFP fusion protein remains stable in freeze-dried yeast powder for more than a 13 and 6 months when stored at 30 °C and 37 °C respectively. Stability of the heterologous protein remains unaffected during the process of heat-inactivation and lyophilization. The mass of lyophilized yeast powder remains almost unchanged during the entire period of storage. Expressed protein remains intact even after two cycles of freeze and thaws. The protease deficient strain appears ideal for the development of whole recombinant yeast-based vaccines. The cellular abundance of expressed antigen in dry powder after a year was comparable to freshly lyophilized cells. SEM microscopy showed the intact nature of cells in powdered form even after a year of storage at 30 °C. Observation made in this study showed that freeze-dry yeast powder can play a vital role in the development of thermostable vaccines.

Introduction

In the last century, vaccines have proved to be one of the most important medical interventions in the fight against infectious diseases. Eradication of smallpox in 1980 (reviewed in **Fenner 1982)** and soon polio (reviewed in **Norrby *et al.* 2017)** are important success stories associated with the benefits of vaccines in public health. Widespread application of vaccines also leads to a sharp decline in newer cases of tuberculosis, hepatitis, measles, tetanus and other infectious diseases (**Versteeg *et al.* 2019**). The importance of vaccines can also be underscored, by the fact that there is a race for vaccine development against novel coronavirus (SARS-CoV-2) responsible for present COVID-19 pandemic **(Callaway 2020)**. As per WHO estimates, every year, around 2-3 million lives are saved by vaccine application **(Cruz-Resendiz *et al.* 2020).** Still, millions of individuals (out of which majority are children below 5, years of age) die from vaccine-preventable diseases. Despite the significant efforts at every step, millions of people are left unvaccinated, thus putting them at the risk of getting the infection at some point in life. According to GVPA, 2011-2020 review report approximately 19.4 million infants did not receive lifesaving vaccines in 2018 (https://www.who.int/immunization /globalvaccine_action_plan/en/). Among the various issues, associated with presently licensed vaccines (summarized by **Kumar and Kumar 2019),** inherent thermolabile nature of these vaccines makes their storage, transportation and dissemination a daunting task. In the field of vaccines, this is commonly known, as the "Cold Chain" problem and, at every step, maintenance of refrigerated condition (2-8 °C in general) is a must. In many instances, maintenance of continuous cold chain raises the cost of the vaccines by about 80 % (**Bandau *et al.* 2003; Das 2004**). Exposure of vaccine to sub-optimal temperature even for short duration leads to vaccine degradation and dramatic loss in efficacy or potency which forces almost 50 % of the vaccines to be discarded before their application (**Brandau *et al.* 2003; Hill *et al.* 2016**). This scenario becomes even more relevant in resource-poor settings in countries of Asia and Africa, which lack access to vaccines **(Chen and Kristensen 2009; Das 2004).** Therefore, improving the thermostability of available vaccines at ambient temperature is highly desirable.

To address, the problem of poor thermostability and short shelf life of vaccines at ambient temperature, different approaches have been taken in the past. In the case of human enterovirus type 71, biomineralization of virus particle improves thermal stability significantly (**Wang *et al.* 2013**). In another common approach, modification in a liquid formulation was found encouraging. For example, the addition of stabilizers like deuterium oxide, proteins, $MgCl_2$, and non-reducing sugars in vaccine formulation improves their thermal stability (**Milsstein *et al.*

1997; Alcock _et al._ 2010). Similarly, the addition of anionic nanogold particles and PEG

(polyethene glycol) improve the thermal stability of some of the vaccines significantly **(Pelliccia**

_et al_. 2016). Recently the coating of bacterial cells or viral particles in a thin film of sugar gave

promising results **(Leung _et al_. 2019; Bajrovic _et al._ 2020).** All the above-highlighted

approaches used the addition of one or more chemicals in vaccine preparation which,

necessitating additional safety tests and other regulatory procedures. Moreover, the application

of all the aforementioned approaches kept the vaccine stable only for a short duration depending

upon vaccine and storage temperature. Use of these approaches makes the entire process

lengthier and more cumbersome. Therefore, a procedure, which is simple, cost-effective, safe,

while improving the thermal stability and shelf life of a vaccine at ambient temperature will be

desirable.

It is well known and established, that the freeze-drying or lyophilization of biomolecules and

biopharmaceuticals (proteins, lipids, nucleic acids) improves their stability as well as shelf life

significantly **(Emami _et al._ 2018; Liao _et al._ 2004; Song _et al._ 2017).** Lyophilization has been

shown to improve the stability of whole cells (example bacterial cell) and viral particles that are

regularly used in conventional vaccines preparation **(Wang _et al._ 2012; Maa _et al._ 2004;**

Garmise _et al._ 2007). Although purified protein antigen can also be lyophilized and can be used

as vaccine but poor immunogenicity and fast body clearance of pure protein remains an

important concern**.** Apart from that application of purified antigen (subunit vaccines) will require

use of adjuvant and will involve protein purification which will further increase cost, time and

will affect the number of doses that can be available. On top of that lyophilized protein antigen

will again require cold chain (**reviewed by Vartak _et al._ 2016; Kumru _et al._ 2014**). The natural

adjuvant nature of yeast cell walls makes it possible to use lyophilized recombinant cells without

the addition of an external adjuvant **(Stubbs _et al._ 2001)**. Unlike subunit vaccine, the yeast-based

vaccines do not require protein purification. Yeast cells are efficiently taken up, by APCs **(Xiang**

_et al._ 2006). Handling of recombinant yeast is much easy and safer compared to infectious

biological entities. Most importantly, the application of inactivated yeast cells is found safe and

well-tolerated in human subjects **(Gaggar _et al._ 2014)**.

This forced us to investigate, whether whole recombinant yeast lyophilized powder can keep the

heterologous protein (acting as an immunogen) intact when stored, at ambient temperature. In

this proof of concept study, we showed that protein antigen remains stable in lyophilized yeast

powder for more than a year when stored at 30 °C. Stability of heterologous fusion protein

remains unaffected during the process of heat-inactivation and lyophilization. Apart from this,

the mass of lyophilized yeast powder remains essentially unchanged. The observations from this

study will help in developing a thermostable vaccine with long shelf life even under non-

refrigerator condition (2-8 °C) using a yeast-based platform.

Materials and methods

Yeast strains

Haploid auxotrophic PPY12h (*arg4 his4*) **(Gould *et al.* 1992)** and protease deficient SMD1163

(*pep4 prb1 his4*) **(Gleeson *et al.* 1998)** *Pichia pastoris* strains, were used in the entire study**.**

Media

YPAD media (1 % yeast extract, 2 % peptone, 0.05% adenine and 2 % dextrose), SD+CSM-His

(0.17 % YNB without amino acid and ammonium sulfate, 0.5 % ammonium sulfate, 0.08 %

CSM-His, 2 % agar, 2 % dextrose) **(Kumar 2019)**.

Cloning of *E. coli* curli protein

Synthetic construct coding for *E. coli* curli was synthesized by a commercial vendor (Genewiz,

New Jersey, USA) into a pUC57-Amp vector. Fragment coding of curli ORF was excised from

the vector and subcloned into *Pichia pastoris* integrating vector pIB2 **(Sears *et al.* 1998)** into

which cycle 3 GFP, was cloned previously **(Crameri *et al.* 1996)**. The sequence of ORF was the

same as that of the original construct for which sequence, was deposited in Genebank with the

following accession number MH264502 **(Kumar 2018)**. Expression of the CSGA-GFP fusion

protein was under the constitutive GAP promoter (Glyceraldehydes-3 phosphate dehydrogenase).

The combined mass of the fusion protein was 42.9 kDa. It is, important to note that no codon

optimization for the construct was used in this study.

Yeast transformation

Pichia pastoris (now officially *Komagataella phaffii* or *K. phaffii*) **(Kurtzman 2009; Kumar *et***

***al.* 2020)** transformation was performed as described previously **(Kumar 2019)** and briefly

mentioned here. The final plasmid (pRK10) was linearized by digestion with *EcoNI* which

cleaves within the *HIS4* marker gene. Linearized plasmid was, transformed into PPY12h, and

SMD1163 strain using the electroporation method (GenePulser Xcell from Biorad).

Transformants were selected, on *His⁻* plates and positive transformants were confirmed, both by

the fluorescent microscopy and western blot.

Protein extraction from regular cycling cells

Presence of *E. coli* CSGA-GFP protein in *P. pastoris* was confirmed by detecting CSGA-GFP

fusion protein using polyclonal rabbit anti-GFP antibodies (from Life Technologies USA; cat #

A-11122). Amount of protein was normalized, based on either the number of cells or by dry

mass of cells. Cells were treated, with 12.5 % TCA and samples were incubated at -80 °C for one
hour or overnight. On completion of incubation, samples were thawed, at room temperature and,
the samples were centrifuged, at 12000 g for 8 min. The supernatant was, discarded and, the
pellet was, washed twice with 80 % chilled acetone. Finally, protein pellet was air-dried and was
resuspended, in 150 µL 1 % SDS and 0.2 N NaOH. Then 150 µL 2X dye (100 mM Tris HCl
pH6.8, 200 mM DTT, 4 % SDS, 0.2 % bromophenol blue, 20 % glycerol) was added to the
sample and samples were heated for 5 min at 95 °C using dry heating block. Samples were
cooled down, to room temperature, vortexed, spanned and, an equal amount or volume of
samples was loaded, in each well of 10 % SDS-PAGE along with pre-stained protein marker
(Biorad, cat # 161-0376). Samples were run at constant 100V till the dye front reaches the
bottom of the gel **(Kumar 2019)**.

Western blot

On completion of SDS-PAGE run, proteins were transferred onto nitrocellulose membrane by
wet transfer (at constant 100 V for 1 hour at 4 °C) as described elsewhere **(Towbin *et al.* 1979)**.
The blotting membrane was incubated in blocking buffer (5 % nonfat skimmed milk powder in
TBST) (TBST, 137 mM NaCl, 2.7 mM KCl, 19 mM Tris base, 0.1 % Tween 20) for 1 hour and
then incubated with anti-GFP antibodies overnight at 4 °C under gentle shaking condition.
Primary antibodies were removed and, the membrane was washed thrice with TBST. The
membrane was again incubated with IRDye®800CW goat anti-rabbit secondary antibodies (from
LI-COR, USA; cat # 926-32211) for one hour at room temperature. The membrane was again
washed thrice with TBST and blot was scanned using LI-COR Odyssey software (from LI-COR,
Nebraska, USA) as per manufacturer instructions.

Heat inactivation of recombinant *P. pastoris*

A single colony of recombinant *P. pastoris,* expressing *E. coli* CSGA-GFP under GAP promoter
was inoculated into 5 mL of YPAD tube. The tube was incubated, at 30 °C, 230 rpm for
overnight growth (source of inoculum). The overnight grew culture was used for the inoculation
of 500 mL YPAD in 2.8 L flasks. The flasks were incubated, at 30 °C, 230 rpm for 48 hours.
Cells were pelleted by centrifugation at 3000 *g* for 5 minutes. Cells were washed twice with
sterile water. Finally, the cell pellet was, resuspended in 25 mL sterile water and tube was
incubated in a water bath maintained at 56 °C for 90 min for heat-inactivation of yeast cells
(Haller *et al.* 2007). After 90 min, tubes were taken out of a water bath, cooled to room
temperature. A small volume (10 µL) of cell suspension was taken out and plated on YPAD plate
to check for the presence of viable cells. The remaining cell suspension was, centrifuged at 3000

*g f*or 5 min, the supernatant was, discarded and, pellet in tubes was put for lyophilization as
described below.

Lyophilization (freeze-drying) of whole recombinant *P. pastoris*

Recombinant *Pichia pastoris* was lyophilized as described elsewhere **(Patterson *et al.* 2015)** and
briefly mentioned here. Cells were harvested in 50 mL falcon tube by centrifugation at 3000 g
for 5 minutes and removal of supernatant. Lyophilization was performed on the cell pellet using
the AdVantage 2.0 Bench Top Freeze Dryer/Lyophilizer (from SP Scientific). Samples in tubes
were regularly checked, for the formation of lyophilized powder. On completion of
lyophilization, tubes were, taken off and the combined mass of tube and lyophilized yeast
powder was noted.

Storage of freeze-dried yeast powder

212 On completion of the process of freeze-drying, tubes were removed from the lyophilizer and
213 caps were tightly closed. Tubes were kept in cardboard boxes and stored in a separate incubator
214 (in the dark away from direct sunlight, no humidity) operating at 30 °C and 37 °C. Then the
215 weight of tubes was noted, at a regular interval. All steps were performed under normal lab
216 environment (i.e. not in an inert and sterile atmosphere).

Protein extraction from lyophilized powder

218 A known amount of lyophilized yeast powder was, taken in a sterile Eppendorf tube. The powder
219 was resuspended in 200 uL of 12.5 % TCA and stored at -80 °C for 1 hour or overnight. The
220 subsequent procedure is the same as described above for protein extraction.

Freeze and thawing of lyophilized yeast powder

222 As above, a known amount of lyophilized yeast powder was, taken into fresh sterile Eppendorf
223 tubes. Tubes were, stored at -20 °C and -80 °C for, two hours followed by incubating the tubes at
224 30 °C for two hours. This process was repeated, depending on the number of freeze-thaw cycles.
225 A known amount of yeast powder was also taken and incubated at 30 °C as a control. Protein
226 extraction was performed as described above.

Field Emission Scanning Electron Microscopy

The shape and surface morphology of lyophilized yeast cells stored at 30 °C for a year was
analyzed, by Field Emission Scanning Electron Microscopy (ZEISS Gemini Sigma 500 VP, Carl
Zeiss Microscopy LLC, NY, USA). Samples were prepared, by mounting a small amount of
lyophilized yeast powder on a double-sided carbon tape on a metal SEM stub. Gently spread the
yeast powder across the surface of the tape. Compressed air was, used to remove loose
lyophilized yeast cells. Yeast samples were sputter-coated with a 3 nm thin layer of Gold-

Palladium for ensuring the conductivity for electrons beam. A beam strength of 5.0 kV and a
working distance in the range of 8–9 mm was, used to visualize yeast samples.

Fluorescence Microscopy

Expression of *E. coli CSGA-GFP* fusion protein in both SMD116 and PPY12h *Pichia pastoris*
was confirmed, by fluorescence microscopy. For each strain, 100 uL cycling cells, were taken
into sterile Eppendorf tube, cells were pellet down by centrifugation at 3000 g for 3 min. The
supernatant was discarded, and the cell pellet was resuspended, in 1 mL sterile Milli Q water. 5
uL of cell suspension was transferred, onto a glass slide. Cells were fixed by adding an
equivalent volume of 1 % agarose. Images were captured, using plan apochromat 100× 1.40-NA
oil immersion objective on a motorized fluorescence microscope (Axioskop 2 MOT plus; Carl
Zeiss) coupled to a monochrome digital camera (AxioCam MRm; Carl Zeiss). Images analysis
was performed, using Axio vision software **(Kumar 2019)**.

Results

The basic assumption behind the study

In our previous study, we showed that stationary phase yeast cells were able to keep the
heterologous protein intact when stored at room temperature (23-25°C) for a year
(Kumar,2018). Unfortunately, the gradual degradation of cells over time suggests that long-term
storage of protein antigen even within cells in a liquid environment is not feasible **(Kumar 2018)**
Therefore, we looked at other ways for long-term storage of cells at ambient temperature. It is a
common observation that active dry yeast powder sold commercially is stable (a significant
proportion of cells retain viability) for almost two years when stored under non-refrigerated
conditions (at 23-25 °C). Even under recommended conditions of storage and handling most of
the available vaccines remain potent for up to 1.5 to 3 years. Therefore it raised the possibility
that dry yeast powder can act as a simple yet an important, means for long-term storage of
immunogen (vaccine) under room temperature (ideally around two years) **(Clenet 2018; Chen *et***
***al.* 2010; Galazka *et al.* 1998; Chen and Zehrung 2013)**. Through this proof of concept study,
we tested whether, lyophilized yeast cells can be used, to store protein immunogen at ambient for
year or more. For this, we expressed *E. coli* curli-GFP fusion protein (*CSGA-GFP*) in *P.*
pastoris. Curli is a bacterial surface protein (fimbri) involved in cell adhesion and biofilm
formation **(Barnhart *et al.* 2006; Nguyen *et al.* 2014)**. Curli is capable of raising immune
response, thereby acting as a good immunogen **(Barnhart *et al.* 2006)** and a suitable, candidate
for the proof of concept study.

Expression of *E. coli CSGA-GFP* in *Pichia pastoris*

E. coli curli was subcloned into *P. pastoris* integrating vector pIB2 as described in the materials

and methods section. Cartoon presentation of the final plasmid used for the expression of *E. coli*

CSGA-GFP fusion protein is shown, in Figure 1A. For checking the expression of the CSGA-

GFP fusion protein, we randomly selected eight colonies from the transformant plate and patched

them on a fresh selection plate (CSM-HIS plate). Next day, the expression of the fusion protein

was first confirmed, by fluorescent microscopy (Figure 1B for PPY12h background). Later in a

separate experiment, the same plasmid was transformed, into SMD1163 strain. Integration of

plasmid and expression of CSGA-GFP was confirmed by fluorescent microscopy (Figure 1C).

Expression of the fusion protein, CSGA-GFP, was further confirmed by western blot. 2 OD_{600nm}

of cells were used for protein extraction for western blot. Blot image in Figure 1D shows that all

the selected colonies were positive. Ponceau-S stain image of the same blot is shown, as a

loading control (Figure 1E) which confirm the proper loading of protein in each well. Specific

anticipated bands are highlighted, by an arrow pointing towards them. We do not get any band in

empty vector control strain used as a negative control. The absence of any signal from negative

control shows the specificity of GFP antibodies used in the experiment. The expression of the

heterologous protein was endogenous. Overall we can say that we were able to express the fusion

protein in *P. pastoris.*

Heat inactivation and lyophilization do not affect the stability of expressed antigen

Yeast species especially *S. cerevisiae* and *P. pastoris* are non-pathogenic and are in the list of

GARS and routinely used for the production of diverse biomolecules for human consumption

(**Ramchuran *et al.* 2005; Basanta *et al.* 2010**). Even then. direct administration of live

recombinant yeast cells into human subjects is not advisable from the point of safety. However,

the level of immune response mounted on the application of whole recombinant yeast is

independent of live or dead nature of yeast (**Lu *et al.* 2004; Franzusoff *et al.* 2005**) raising the

possibility for application of inactivated recombinant yeast. Heat-inactivation of yeast is rapid,

simple and more convenient compared to chemical based-inactivation of bacterial or viral

particles for vaccine preparation. The combined effect of heat-inactivation and lyophilization on

the stability of heterologous protein was missing and through this study, we tried to fill that gap.

Heat inactivation of yeast cells was performed as described in the materials and methods. Heat

inactivation was confirmed by plating small volume of cell suspension on YPAD plates (Figure

2B) along with untreated control cycling cells (Figure 1A). Our present western blot data showed

that the stability of heterologous proteins remains essentially unaffected during heat-inactivation

(Figure 2C) and heat-inactivation followed by lyophilization (Figure 2E). Proper loading and transfer of protein are shown by Ponceau-S stained blots (Figure 2D and 2E respectively). Protein amount was normalized, by the weight of cells used for protein extraction. Results of Figure 2C are in accordance with our previous study **(Kumar 2018)**. Based on the present data, it can be said that together heat-inactivation and lyophilization have no affect protein stability.

The protease deficient yeast strain improves the stability of the expressed fusion protein

In our previous section, we showed that process of heat-inactivation and heat-inactivation followed by lyophilization does not affect the stability of CSGA-GFP in yeast cells. But we do see a very prominent band around 25 kDa region of the blot. Whether the observed bands were due to degradation of the fusion protein on heat-inactivation and lyophilization, or whether they were a result of protease activities in cytosol and vacuoles was not clear. To sort out this, we transform the same construct into protease deficient strain (SMD1163), which lacks Prb1 (cytosolic) and Pep4 (vacuolar). Our present western blot data pointed towards the idea that observed lower bands are the result of vacuolar protease action (Figure 2G). Like previous blots, we again observed lower bands in PPY12h strain used as a control (Figure 2G, well1), but failed to detect the same bands in SMD1163 (Figure 2G, well 2,3). The presence or absence of lower bands in PPY12h and SMD1163 respectively is not due to the difference in the amount of protein loaded into each well was confirmed by Ponceau S stained blot image (Figure 2H). Therefore, it can be concluded that lower observed bands are not due to heat-inactivation or lyophilization but may be due to the action of cellular proteases. Our observation is supported by the fact that, during stress, autophagic pathways get activated, which forces the degradation of cellular components in vacuoles or lysosomes **(Klionsky *et al.* 2007; Takeshige *et al.* 1992; Kumar *et al.* 2020)**. Therefore lower bands detected in PPY12h and their absence in SMD1163 may be due to autophagic degradation of fusion proteins in vacuoles and GFP (26.7 kD), which is quite stable in vacuoles (whose free release in the vacuole is used in autophagic assays) was detected by antibodies **(Klionsky *et al.* 2007; Takeshige *et al.* 1992)**.

Mass of freeze-dried yeast powder remains unchanged during storage

After confirming that heat-inactivation, as well as freeze-drying, do not affect the stability of expressed protein, we increase the volume of culture to 1 L. PPY12h strain expressing CSGA-GFP was grown in bulk volume (1 L). Cells were, harvested heat-inactivated and freeze-dried as described in Materials and Methods. After lyophilization, tubes were taken off from lyophilizer, screwed the cap and weighed. Initial mass was taken and stored at 30 °C for one year. The weight of the tube was, then checked, regularly and sometimes the tubes were opened, for a short

time. Table 1 is showing the data for the mass of powder stored at 30 °C. Similarly, we checked,

the weight of freeze-dried yeast powder stored at 37 °C for six months (Table 2). Present data

showed that mass of freeze-dried powder remains essentially unchanged under both conditions of

storage. The slight variation observed in weight may be due to moisture that might be entered

into tubes when tubes were open for some time. Note in Table 1 data is shown for two

conditions. In one condition cells were lyophilized after-heat inactivation (Table 1 second and

third column) and in another, cells were lyophilized without heat-inactivation (table 1 fourth and

fifth column). Based on the data shown in tables, it can be said, that the mass of stored

lyophilized cells does not change significantly.

Cells in lyophilized powder retain their intactness

Next we asked whether cells in lyophilized powder also remain intact by checking the

morphology and surface appearance of cells using SEM microscopy. A small amount of

lyophilized powder (figure 3A cells were lyophilized without heat-inactivation and, figure 3B

cells were lyophilized after heat-inactivation) was taken and SEM microscopy was performed.

SEM images showed that the cells remain intact in lyophilized powdered form when stored at 30

°C for more than a year (figure 3C and 3D). Although cells were intact under both the condition,

we observed a difference in the texture of the powder. Freeze dry powder of cells without heat-

inactivation appears less compact and less dense (figure 3A) whereas heat-inactivated powder

appear more compact and denser (figure 3B). Thus, it can be said, that cells in lyophilized

powdered form remain intact for a year even when stored under ambient or room temperature.

Expressed antigen remains stable in lyophilized yeast powder stored at 30 °C temperature

Although SEM data in the previous section confirmed intact nature of lyophilized yeast cells

after a year of storage at 30 °C, whether expressed protein also remained intact was unknown. To

check the stability of the expressed protein, we took equal amounts of lyophilized powder from

both the condition and extracted the protein. An equal amount of whole cell lysate was loaded on

10 % SDS-PAGE and, the fusion protein was detected using anti-GFP antibodies. Our present

western blot data showed that expressed protein remains stable in powdered yeast stored at 30 °C

for a year (figure 4). The expressed protein remains stable under both the conditions in which

cells were lyophilized without heat-inactivation (figure 4A) and in which cells were lyophilized

after heat-inactivation (figure 4B). Before incubation in blocking buffer, blots were stained with

Ponceau S for showing the loading control (figure 5C and 5D for figure 4A and 4B respectively).

Encouraged from the stability of the protein in lyophilized powder stored at 30 °C for a year, we

checked whether expressed protein also remains stable in lyophilized yeast cells stored, at 37 °C

for six months. Our present western blot data showed that expressed protein remains stable in powdered yeast even when stored at 37 °C at least for six months (figure 4E). Loading control is shown in figure 4F. Therefore, it can be, concluded that expressed protein remains stable in lyophilized yeast cells when stored at a temperature ranging from 30 °C -37 °C for quite a long time.

Freeze and thaw do not affect protein stability in powdered yeast

Most of the commonly used vaccines are stored at 2-8 °C, while some of them are also stored at -15 °C to -50 °C (e.g. measles, mumps and rubella or MMR$_I$ is stored at +8 °C to -50 °C, https://www.merckvaccines.com/mmr/storage-handling/). However for most vaccines, exposure to sub-optimal temperature (i.e. more than 2-8 C or below freezing temperature) even for a short duration is known to reduce vaccine potency dramatically) (**Brandau *et al.* 2003; Hill *et al.* 2016**). An ideal vaccine should remain stable and retain potency even when stored under the non-refrigerated condition as well as under accidental freeze condition. For investigating the effect of freeze and thaw on the stability of the heterologous protein in lyophilized yeast cells, known amount of yeast powder was taken into separate tubes. Tubes were stored at -20 °C and -80 °C for two hours. Freeze and thaw were performed in one cycle and two-cycle. For each condition, samples were taken, in duplex. After a freeze, thawing was performed by incubating samples at 30 °C for two hours. Our present western blot data showed that expressed protein in lyophilized yeast powder remain stable following one as well as two cycles of freeze and thaw at both -20 °C (figure 5A) and -80 °C (figure 5B). Ponceau S stained image of the membrane (as a loading control), is shown next to the blot. A known amount of yeast powder, stored at 30 °C separately was taken, as a control. Conditions of freeze and thaw are mentioned, in the materials and methods section. Based on present data, it can be said that expressed protein remains stable both under the non-refrigerated condition for a year as well as under freeze and thaw condition.

The abundance of expressed protein after a year was similar to fresh cells

In the previous section, we showed that the expressed CSGA-GFP fusion protein was stable in lyophilized yeast powder when stored at 30 °C for a year and also survived two cycles of freeze and thaw. But whether the expressed protein deteriorated over time and to what extent still needed to be determined. To answer this question, we compared the level of expressed protein in yeast powder stored for a year to the freshly lyophilized yeast cells. Before comparing, the level of fusion protein we checked the overall protein content in the two samples (Figure 6A). Although an equal amount of whole-cell lysate, was loaded for each sample, we still see a slightly low level of proteins in a sample from year-old powder (well 6,7) compared to freshly

lyophilized cells (well 4,5). Apart from this we also observed relatively more background in a

year old sample. Whether the observed difference in the level of proteins and background is due

to the difference in protein extraction efficiencies or protein degradation is unclear. But we do

find a problem in re-suspending the powder entirely even on vigorous vortex. We also loaded an

equal amount of cell lysate from normal cycling cells as control (well 2,3).

After looking at the overall proteome of lyophilized cells stored for a year and freshly

lyophilized cells, we proceed to compare the level of the expressed fusion protein in two

samples. Our present data (Figure 6B) shows that level of the CSGA-GFP fusion protein is

almost similar in freshly lyophilized cells (figure 6B, well 4) to that of one-year-old powder

(figure 6B, well 2,3). Note we used the same sample for western blot which, were used in

running SDS-PAGE shown in figure 6A. And the slight difference which appears in two samples

may be due to the issues mentioned above. Loading control is shown, by Ponceau-S stained

image of the same blot (figure 6C). To capture any possible difference in samples blot images

were captured at low intensity of exposure (data for high exposure, is not shown).

Discussions

Storage, transportation and distribution of vaccines before their final application are still presents

difficulties. On one hand, exposure of vaccine to temperatures more than the recommended

temperatures (generally 2-8 °C) leads to vaccine degradation, denaturation and finally loss of

vaccine potency or efficacy **(Brandau *et al.* 2003; Hill *et al.* 2016)**. On the other hand, exposure

of vaccines to below freezing conditions also affect vaccine potency **(Lloyd *et al.* 2015; Kumru**

***et al.* 2014)**. Therefore, maintenance of optimum conditions from point of manufacturing till the

final application is required which is more difficult in less well-developed countries. A solution

to this is the development of thermostable vaccines. Availability of thermostable vaccines will be

an important, step in global immunization step **(Lee *et al.* 2017)**. The thermostable nature of

future vaccines will not only make vaccine transport, storage and distribution more convenient

and economical but, will also help in saving a huge, amount of life-saving vaccines. Currently,

the development of thermostable vaccines are yet unmet, with little success in which thermal

stability was observed (only for a few months at best and that too in few cases) (**Leung *et al.***

2019; Chu *et al.* 2016; Mistilis *et al.* 2017; Hassett *et al.* 2013, 2015; Chen *et al.* 2010;

Ohtake *et al.* 2010, 2011; Lovalenti *et al.* 2016). By using thin-film coating, one study was able

to keep the adenovirus stable at room ambient temperature for three years (**Bajrovic *et al.* 2020).**

It will be interesting to see whether this approach can be applied to other vaccines. But this study

showed the possibility that a vaccine may be stored and transported at ambient temperature. So

far, most of the approaches used for enhancing the thermal stability and shelf life of vaccines at
ambient temperature require the addition of different chemicals. Assessing the safety of each
chemical added in formulation requires more regulatory clearance, time and money. Moreover,
whether the developed formulation applies to a wide range of vaccines also remain a matter of
future research. Unlike conventional vaccine preparation which requires growth of bacterial
culture or viral particles followed by formalin-based inactivation which itself is quite lengthy,
growth and heat-inactivation of yeast are rapid and more economical.

The degradation of expressed immunogen by vacuolar proteases may pose a serious issue in the
development of a whole recombinant yeast-based vaccine. The degradation of immunogen will
not only reduce the level of the immunogen in cells but may also affect the level of the immune
response. It is, therefore, important to use protease deficient strain (like SMD1163) **(Gleeson *et
al.* 1998)**. Compared to budding yeast, the post-translation modification *P. pastoris* is closer to
those of higher eukaryotes **(Cregg *et al.* 1993)**. The fact that P. pastoris is better fermenter is
other added advantages **(Cereghino and Cregg, 2000)** when a large amount of vaccine is needed
such as during a pandemic.

Surprisingly, we do not observe any bacterial contamination in yeast powdered stored for the
year even though we opened the tubes several times in a year. This can be the result of heat-
inactivation followed by lyophilization. This observation showed, that lyophilized yeast powder
can be, stored for a long time even in the absence of antibiotics. Several conventional vaccine
preparations involve the use of antibiotics such as MMR which not only increase the cost of
vaccines production but also cause an allergic response in some people **(Chung, 2014)**.
Although we have shown the data for the stability of heterologous protein for a year (at 30 °C)
and six months (at 37 °C), we expect that expressed protein may remain stable for much longer
(as a result we are still keeping our samples at the aforementioned condition). Further, it will be
interesting to see whether recombinant yeast lyophilized powder stored at ambient for a year or
more can mount a specific immune response *in vivo* models like a mouse. Previous observation
that APCs can process freeze-dried yeast cells and can mount an immune response relates to our
present and future study in this direction **(Patterson *et al.* 2015).** It is almost impossible to
mimic the real scenario, but our present data showed that lyophilized yeast powder, can prevent
immunogen degradation due accidental freeze and thaw that was lacking in other approaches
used for the development of thermostable vaccines. Yeast-based vaccines, which are in different
phases of clinical trials may present opportunities where the present method for developing

thermostable vaccines can be adopted directly **(Kumar and Kumar, 2019; Ardiani** *et al.* **2010).**
Moreover, there is no obvious barrier why our approach cannot be directly adopted for subunit
vaccines where protein immunogen have been already characterized **(Agwale** *et al.* **2002; Zhang**
et al. **2015; Schiller and Lowy, 2015)**.
Overall, we have shown that a dry yeast powder is a simple yet effective way for long-term
storage of vaccines (immunogen) under non-refrigerated condition. Data shown in this report
showed that yeast-based approach for thermostable vaccine development prevents vaccine
deterioration both at high temperature (above 2-8 °C) as well as below freezing condition (below
0 °C) which is common in countries of Europe, North America and parts of Asia. We believe
that this cost effective freeze-dried yeast powder approach may be one solution to the cold chain
problem and will go towards attaining the universal immunization program.

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Authors Contribution

RK conceived, designed, performed the experiments, analyzed the data and wrote the
manuscript. BNK perform SEM and fluorescence microscopy.
489
490 **Conflict of interest**
491 The authors declare that no conflict of any kind exists.

492 **Declaration**
493 Authors declare that the present study does not involve any human subjects or animal handling
494 by any author mentioned in this study.
495

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Legends

Tables

Table 1. Mass of lyophilized yeast powder stored at 30 °C for one year

Table 2. Mass of lyophilized yeast powder stored at 37 °C for six months

Figures

For Graphical abstract

Schematic showing the basic workflow of the entire study.

Figure 1. Expression of *E. coli* CSGA-GFP fusion protein in *P. pastoris*. (A) Map of the final

plasmid used for expression of the CSGA-GFP fusion protein. Expression of a CSGA-GFP

fusion protein in (B) PPY12h and (C) SMD1163 strain. Scale bar represents 5 uM. (D) Image of

western blot showing expression of the CSGA-GFP fusion protein in PPY12h. Bands of interest are pointed by arrows towards them, while free GFP is pointed by an Asterisk mark. (E) Proper loading and protein transfer are shown, by Ponceau S stained blot image. Well, 1 (pre-stained protein marker), well 2 (empty vector as a negative), well 3-8 (transformants).

Figure 2. Heat-inactivation and freeze-drying do not affect protein stability. (A) Control untreated viable cells. (B) Complete loss of viability on heat-inactivation. (C) Western blot image showing the effect of heat-inactivation on protein stability. Well, 1, 2 (untreated cycling cells); well 3 4 (heat-inactivated cells). (D) Proper loading and protein transfer is shown by Ponceau S stain blot image. Western blot image showing the effect of freeze-drying on protein stability (E) well 1, 2 (control cycling cells); well 3,4 (lyophilized yeast cells). (F) Proper loading and protein transfer is shown by Ponceau S stain blot image. Bottom free GFP bands are due to vacuolar protease (G) well 1 (PPY12 strain); well 2-3 (SMD1163, protease deficient strain, lack Prb1 cytosolic and Pep4 vacuolar protease). (H) Proper loading and protein transfer is shown by Ponceau S stain blot image. Bands of interest are pointed by arrows towards them, while free GFP is pointed by an Asterisk mark.

Figure 3. Cells in lyophilized powder remain intact for more than a year. Image of lyophilized yeast powder stored at 30 °C for more than a year (A) cells was lyophilized without heat-inactivation and (B) cells were heat-inactivated before lyophilization. SEM images showing external morphology of lyophilized cells after one year (C) lyophilized cells without heat-inactivation and (D) lyophilized cells after heat-inactivated.

Figure 4. Expressed protein remains stable in powdered yeast. (A) Stability of CSGA-GFP in freeze-dried yeast powder stored at 30 °C after one year. (B) Stability of CSGA-GFP in heat-inactivated lyophilized yeast powder stored at 30 °C after one year. (C) Stability of heat-inactivated lyophilized yeast powder stored at 37 °C after six months. In each case, proper loading is shown, by Ponceau S stained blot image. Note, the protein amount was normalized, based on yeast powder used for protein extraction. Bands of interest are pointed by arrows towards them, while free GFP is pointed by an Asterisk mark. Note in figure Y stand for yes and N for no for condition mentioned in the figure.

Figure 5. Freeze and thaws do not affect the stability of the expressed protein. (A) Effect of freeze and thaw at -20 °C. Control (lyophilized powder stored at 30 °C), well 1,2 (once the cycle of freeze and thaw), well 3,4 (two-cycle of freeze and thaw). (B) Effect of freeze and thaw at -80 °C. Same common control as above, well 1,2 (one cycle of freeze and thaw), well 3,4 (two-cycle of freeze and thaw). Note, in all case the protein amount was normalized based on the weight of powder and, each sample was taken in duplex. Loading and proper transfer are shown through Ponceau S stained of same blots. Also, a control sample is common in panel A and B.

Figure 6. The abundance of heterologous protein in a year old powdered and freshly lyophilized cells are similar. (A) The efficiency of protein extraction from different samples. Cycling cells without lyophilized (well 2,3), an equivalent number of cells after lyophilized powder (well 4,5), an equivalent amount of lyophilized powder stored at 30 °C for one year (well 6,7), well 1 for protein marker. (B) The abundance of expressed protein is similar in freshly

706 lyophilized cells and year-old powder lyophilized powder after one year at 30 C (well 2,3),
707 freshly prepared powder (well 4) and protein marker (well 1). The same blot was Ponceau S
708 stained before blocking in skimmed milk powder in TBST shown as a loading control (C).

709
710 **Tables**
711 **Table 1. Mass of lyophilized yeast powder stored at 30 °C for one year**
712

Month	Heat inactivated		Without heat inactivated	
	Tube 1	Tube 2	Tube 1	Tube 2
0	16.88	16.97	17.06	17.26
2	16.97	16.96	17.14	17.29
4	16.95	16.91	17.14	17.27
6	16.98	16.93	17.13	17.26
8	16.95	16.91	17.15	17.21
10	16.96	16.93	17.16	17.21
12	16.97	16.95	17.15	17.22

713
714 **Note: Net mass is in grams. Reported mass is the combined mass of tube and powder.**
715
716 **Table 2. Mass of lyophilized yeast powder stored at 37 °C for six months**
717

Month	Tube 1	Tube 2
0	1.02	0.98
1	1.02	1.01
2	1.05	1.01
3	1.05	1.00
4	1.04	1.00
5	1.05	1.00
6	1.05	1.00

718
719 **Note: Net mass is in grams (reported are the values after deducting mass of empty tube)**

Figure 1. Expression of E. coli CSGA-GFP fusion protein in *P. pastoris.* (A) Map of the final plasmid used for expression of the CSGA-GFP fusion protein. Expression of a CSGA-GFP fusion protein in (B) PPY12h and (C) SMD1163 strain. Scale bar represents 5 uM. (D) Image of western blot showing expression of the CSGA-GFP fusion protein in PPY12h. Bands of interest are pointed by arrows towards them, while free GFP is pointed by an Asterisk mark. (E) Proper loading and protein transfer are shown, by Ponceau S stained blot image. Well, 1 (pre-stained protein marker), well 2 (empty vector as a negative), well 3-8 (transformants).

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