

EXPERIMENTAL  
ARTICLES

## Selenium Tolerance of Yeasts

V. I. Golubev\* and N. V. Golubev\*\*

\*Skryabin Institute of Biochemistry and Physiology of Microorganisms, Russian Academy of Sciences, pr. Nauki 5,  
Pushchino, Moscow oblast, 142290 Russia

\*\*Mendeleev University of Chemical Technology, Moscow, 125820 Russia

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**Abstract**—Selenium tolerance of yeasts widely varies: the growth of some yeasts can be inhibited by a selenium concentration as low as  $10^{-4}$  M, whereas others can grow in the presence of  $10^{-1}$  M selenium. Homogeneous yeast taxa are characterized by a certain level of selenium tolerance, and heterogeneous taxa show a variable level of tolerance to selenium. In general, ascomycetous yeasts are more tolerant to selenium than basidiomycetous yeasts. Among the ascomycetous yeasts, the genera *Dekkera* and *Schizosaccharomyces* exhibited the lowest and the species *Candida maltosa*, *Hanseniaspora valbyensis*, *Kluyveromyces marxianus*, and *Yarrowia lipolytica* the highest tolerance to selenium. Among the basidiomycetous yeasts, the genera *Bullera*, *Cryptococcus* and *Holtermannia* showed the lowest and the species *Cryptococcus curvatus*, *Cr. humicola*, and *Trichosporon* spp. the highest tolerance to selenium. The selenium tolerance of yeasts depends on the composition of the growth medium, in particular, on the presence of sulfate, sulfur-containing amino acids, and glutamine in the medium.

**Key words:** selenium, yeasts, tolerance.

Selenium is an essential chemical element for living organisms, since it is a constituent of many enzymes. The presence of selenium in diets prevents the development of cancer tumors [1]. In some regions, the “selenium yeast” *Saccharomyces cerevisiae* is used to compensate for the deficiency of this element in human and animal diets [2]. At the same time, elevated levels of selenium may be toxic to organisms. The toxicity of selenium can be explained by the fact that this element is an analogue of sulfur. Accordingly, when selenium is present in great amounts, it incorporates into sulfur-containing amino acids and then in proteins, changing their conformation and functional activity [3].

In spite of the wide use of the selenium yeast, the effect of selenium on yeasts has not yet been studied in depth. The aim of the present work was to investigate selenium tolerance in a diversity of yeasts.

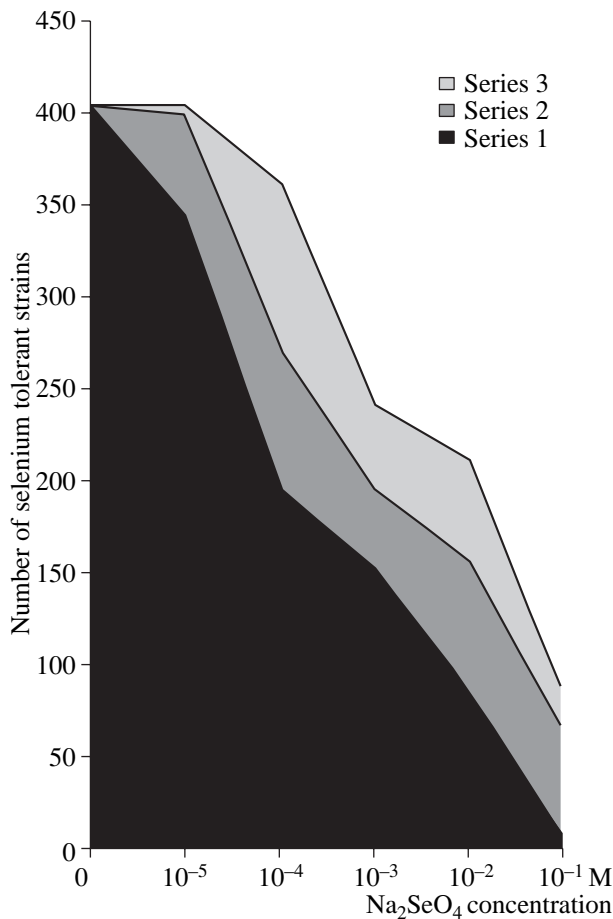
### MATERIALS AND METHODS

The yeast strains studied in this work (404 strains of 181 species belonging to 40 genera) were either obtained from the All-Russia Collection of Microorganisms (VKM) or were isolated by us during the microbiological survey of the Prioksko-Terrasnyi Zapovednik (Oka Terrace Nature Reserve) [4]. The tolerance of yeasts to selenium was tested by growing them at 25°C in a liquid medium containing (g/l) glucose, 10.0; peptone, 5.0; yeast extract, 1.0; agar, 20.0; and  $\text{Na}_2\text{SeO}_4$  at concentrations varying tenfold from  $10^{-5}$  to  $10^{-1}$  M. This medium was inoculated with yeast

cells grown on wort agar for 2–3 days. The effect of glutamine (5 g/l),  $(\text{NH}_4)_2\text{SO}_4$  (3 g/l), and the sulfur-containing amino acids methionine and cysteine (3 g/l) on the selenium tolerance of yeasts was studied by adding these compounds at indicated concentrations to an agar synthetic medium lacking  $\text{Na}_2\text{SeO}_4$  and amino acids. Cell suspensions used for inoculation contained  $10^6$  cells/ml. The control cultures were grown in the same media as the experimental cultures, but without  $\text{Na}_2\text{SeO}_4$ .

**Table 1.** Yeast species and the number of strains whose growth is partially suppressed by  $10^{-5}$  M  $\text{Na}_2\text{SeO}_4$  and completely inhibited by  $10^{-4}$  M  $\text{Na}_2\text{SeO}_4$

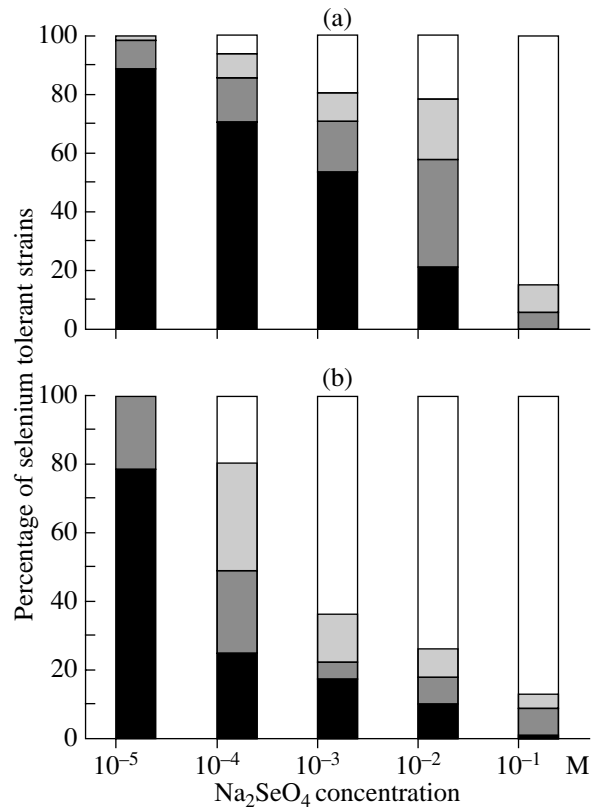
<i>Bullera alba</i> (5)	<i>Cr. phenolicus</i> (1)
<i>B. armeniaca</i> (1)	<i>Cr. terreus</i> (7)
<i>B. pseudoalba</i> (1)	<i>Cr. terricola</i> (1)
<i>Candida silvae</i> (1)	<i>Cr. vishniacii</i> (2)
<i>Cryptococcus chernovii</i> (1)	<i>Dekkera anomala</i> (2)
<i>Cr. dimennae</i> (1)	<i>Holtermannia corniformis</i> (2)
<i>Cr. flavus</i> (1)	<i>Pichia norvegensis</i> (1)
<i>Cr. fuscescens</i> (1)	<i>Rhodotorula yakutica</i> (1)
<i>Cr. hungaricus</i> (1)	<i>Schizosaccharomyces pombe</i> (2)
<i>Cr. marinus</i> (1)	



**Fig. 1.** The distribution of yeast strains according to the ability to grow in the glucose–peptone medium in the presence of different concentrations of  $\text{Na}_2\text{SeO}_4$ . Series 1, good growth; series 2, slow latent growth; and series 3, poor growth.

#### μMRESULTS

$\text{Na}_2\text{SeO}_4$  at a concentration of  $10^{-5}$  M did not influence the growth of almost all of the yeast strains under study on glucose–peptone agar, except for some species of the genus *Cryptococcus*, whose growth was somewhat suppressed at this selenium concentration and was partially or completely inhibited at a selenium concentration of  $10^{-4}$  M (Tables 1 and 3). Most of the yeast strains could grow at the last selenium concentration, but only half of them grew poorly at a selenium concentration of  $10^{-3}$  M.  $\text{Na}_2\text{SeO}_4$  at a concentration of  $10^{-2}$  M inhibited the growth of most of the yeast strains (Figs. 1 and 2), and only some strains could grow at a selenium concentration of  $10^{-1}$  M (Table 2). In the last case, the growth was poor, and the yeasts produced pink- to red-colored colonies, whereas normally (i.e., in the absence of  $\text{Na}_2\text{SeO}_4$ ) they are colorless. In the course of further incubation, colorless secondary colonies appeared on the surface of the red colonies.



**Fig. 2.** The distribution of (a) ascomycetous and (b) basidiomycetous yeasts according to the ability to grow in the glucose–peptone medium in the presence of different concentrations of  $\text{Na}_2\text{SeO}_4$ . The dark portions of the bars indicate the percentage of yeast species capable of growing at the particular concentration of  $\text{Na}_2\text{SeO}_4$ .

Among the ascomycetous yeasts studied, the genera *Citeromyces*, *Clavispora*, *Dekkera*, and *Schizosaccharomyces* showed the lowest tolerance to selenium (the members of these genera could grow at selenium concentrations not exceeding  $10^{-4}$  M). At the same time, almost all species of the genera *Aciculoconidium*, *Hanseniaspora*, *Kazachstania*, *Kluyveromyces*, *Mastigomyces*, *Nadsonia*, *Saccharomyces*, *Saccharomycoides*, *Schizoblastosporion*, *Wickerhamia*, and *Yarrowia* were able to grow at a selenium concentration of  $10^{-2}$  M. Among the basidiomycetous yeasts studied, only two cryptococci, *Cr. curvatus* and *Cr. humicola*, and some species of the genus *Trichosporon* exhibited high selenium tolerance (Table 2), whereas the other basidiomycetous yeasts studied could grow at selenium concentrations not exceeding  $10^{-4}$  M.

Some yeast genera were found to be heterogeneous with respect to selenium tolerance. For instance, the maximal concentrations of selenium appropriate for the growth of *Candida* spp. varied from  $10^{-5}$  to  $10^{-1}$  M. Similarly, some yeast species also exhibited heterogeneity with respect to selenium tolerance (Table 3).

**Table 2.** Yeast species and the number of strains which are able to grow in the presence of  $10^{-1}$  M  $\text{Na}_2\text{SeO}_4$ 

<i>Candida maltosa</i> (3)	<i>Tr. coremiiforme</i> (1)
<i>Cryptococcus curvatus</i> (3)	<i>Tr. cutaneum</i> (6)
<i>Cr. humicola</i> (4)	<i>Tr. dulcitum</i> (1)
<i>Hanseniaspora valbyensis</i> (2)	<i>Tr. gracile</i> (1)
<i>Kluyveromyces marxianus</i> (23)	<i>Tr. moniliiforme</i> (1)
<i>Trichosporon aquatile</i> (1)	<i>Yarrowia lipolytica</i> (19)
<i>Tr. brassicae</i> (1)	

Sulfates in the growth medium prevented the inhibitory effect of selenium on yeasts. For instance, in the presence of  $(\text{NH}_4)_2\text{SO}_4$  and  $10^{-3}$  M  $\text{Na}_2\text{SeO}_4$  in the medium, good growth was observed for members of the genera *Bullera*, *Cryptococcus*, *Cystofilobasidium*, *Dekkera*, *Erythrobasidium*, *Fibulobasidium*, *Filobasidium*, *Holtermannia*, *Pseudozyma*, *Rhodotorula*, *Schizosaccharomyces*, and *Tsuchiyaea*, which grew poorly, if at

all, in the presence of  $10^{-4}$  M selenium when ammonium sulfate was absent in the medium. Of interest is the fact that ammonium sulfate did not exert any beneficial effect on the highly selenium tolerant yeasts (Table 2), whereas the sulfur-containing amino acids methionine and cysteine enhanced the growth of these yeasts in the presence of  $10^{-1}$  M  $\text{Na}_2\text{SeO}_4$ . Glutamine also increased the selenium tolerance of yeasts about tenfold.

## DISCUSSION

As can be seen from the data presented, almost all of the yeasts studied grew well in the presence of  $10^{-5}$  M  $\text{Na}_2\text{SeO}_4$ , and only some of them were inhibited by  $10^{-4}$  M  $\text{Na}_2\text{SeO}_4$ . However, as the selenium concentration was raised further, the number of yeasts capable of growth at a particular selenium concentration steeply declined (Fig. 1).

**Table 3.** Yeast taxa which are heterogeneous with respect to selenium tolerance

Species (synonyms) and strains	$\text{Na}_2\text{SeO}_4$ concentration in the medium			
	$10^{-4}$	$10^{-3}$	$10^{-2}$	$10^{-1}$
<i>Candida rugosa</i> BKM Y-1511 ( <i>C. rugosa</i> var. <i>elegans</i> , T) 67T	s +	p +	– s	– p
<i>Cryptococcus albidus</i> BKM Y-714 ( <i>Torula albida</i> , T), 751 ( <i>Torulopsis nadaensis</i> , T), 1983, 1984, 2222 1531 ( <i>Cr. genitalis</i> , A), 1539 ( <i>Cr. albidus</i> var. <i>ovalis</i> , T), 1954, 1955, 2561 1646 ( <i>Cr. albidus</i> var. <i>kuetzingii</i> , T), 2223T	p s +	– p p	– – p	– – –
<i>Cr. laurentii</i> BKM Y-720 ( <i>Torulopsis carnescens</i> , T), 1291 ( <i>Rhodotorula peneaus</i> , T), 1594 328 ( <i>Rh. aurea</i> , T), 1032, 1595 ( <i>Cr. laurentii</i> var. <i>flavescens</i> , T), 1987, 2244 1627, 1628, 1665i	– p s	– – p	– – p	– – –
<i>Kluyveromyces thermotolerans</i> BKM Y-894T 533 ( <i>K. veronae</i> , T), 534, 2317	– +	– s	– p	– –
<i>Rhodotorula mucilaginoso</i> BKM Y-83 ( <i>Cr. ludwigii</i> ), 755 ( <i>Torulopsis nitritophila</i> , T), 1128, 1152, 1308, 1324, 1325, 2058, 2286, 2297 80 ( <i>Cr. corallinus</i> , T), 691 ( <i>Torula aelotiana</i> , T), 1117 ( <i>Cr. pararoseus</i> , T), 1119 ( <i>Cr. rubrorugosus</i> , T), 1123, 1320, 1323, 2283 ( <i>Rh. matritensis</i> , T) 7 ( <i>B. carbonei</i> , T), 17 ( <i>Blastodendron simplex</i> , T), 87 ( <i>Cr. radiatus</i> , T), 253 ( <i>Mycotorula cisnerosi</i> , T), 339T, 341 ( <i>Rh. rubra</i> , T), 343 ( <i>Rh. rubra</i> var. <i>longa</i> , T), 344 ( <i>Torulopsis sanniei</i> , T), 718 ( <i>Torulopsis biourgei</i> , T), 747 ( <i>Torulopsis mannitica</i> , T), 1127, 2650 ( <i>Rh. grinbergsii</i> , T)	p s +	– p s	– – p	– – –
<i>Trichosporon porosum</i> BKM Y-3T 2866, 2867	p +	– c	– p	– –

Note: T denotes "type strain." "+," "s," "p," and "–" stand for "good growth," "slow growth," "poor growth," and "no growth," respectively.

In general, ascomycetous yeasts are found to be more selenium tolerant than basidiomycetous yeasts (compare Figs. 2a and 2b). The only genus of ascomycetous yeasts that is tolerant to high selenium concentrations is *Trichosporon*. This observation does not contradict the fact that the list of highly selenium tolerant yeasts (Table 2) includes two cryptococci and no *Tr. pullulans*, since it is known that yeast taxa are extremely heterogeneous [5]. In particular, the results of rDNA sequencing suggest that the species *Cr. curvatus* and *Cr. humicola* belong to the order *Trichosporonales*, whereas the species *Tr. pullulans* belongs to a different order, *Cystofilobasidiales* [6, 7]. Undoubtedly, these three species need reclassification, as a result of which *Cr. curvatus* and *Cr. humicola* may appear in the genus *Trichosporon* Behrend, and *Tr. pullulans*, in the genus *Tausonia* Babjeva.

The high tolerance of the genus *Trichosporon* to selenium is a unique feature among basidiomycetous yeasts and, hence, selenium tolerance can be used for taxonomic purposes and for the selective isolation of this genus from natural and clinical sources. Similarly, media with increased concentrations of  $\text{Na}_2\text{SeO}_4$  can be used for the selective isolation of some ascomycetous yeasts, such as those of the genus *Yarrowia* (Table 2).

Noteworthy is the fact that heterogeneous yeast taxa also exhibit heterogeneity with respect to selenium tolerance. Taxonomic heterogeneity is especially pronounced in the genus *Candida* and in the yeast species *Cr. albidus* [8], *Cr. laurentii* [9], and *Rh. mucilaginosus* [10]. These species, as well as some others, exhibit high heterogeneity of their strains with respect to selenium tolerance (Table 3) and, hence, their synonymy should be approved by modern methods. Type strains, which are the oldest isolates of particular species, show decreased tolerance to selenium. Consequently, the possibility cannot be excluded that their low selenium tolerance resulted from their long-term maintenance under laboratory conditions.

The degree of selenium tolerance of a yeast is determined by its ability to distinguish selenium and sulfur in the metabolic processes in which these chemical elements are involved. The anions  $\text{SeO}_4$  and  $\text{SO}_4$  compete with one another already at the stage of their uptake by a cell, which is accomplished by the same transport system [11]. The competition between these anions for mutual permeases may explain the beneficial effect of  $(\text{NH}_4)_2\text{SO}_4$  on the yeasts with low or medium tolerance to selenium. At the same time, this mechanism of selenium detoxication by ammonium sulfate probably does not work in the yeasts with high tolerance to selenium. These yeasts may accomplish other detoxication mechanisms, such as the reduction of the Se ion to elemental Se [12, 13], as is evident from the formation of

pink- to red-colored colonies on agar media with a high content of  $\text{Na}_2\text{SeO}_4$ . In this case, the synthesis of sulfur-containing amino acids remains suppressed, as revealed by the growth-stimulating effect of the sulfur-containing amino acids added to the medium. Another possible mechanism of selenium detoxication is its incorporation into the amino acids  $\gamma$ -glutamylmethylselenocysteine and  $\gamma$ -glutamylmethylselenocystathionine, which are not used by cells for protein synthesis [3]. The fact that this mechanism does function in yeasts is evident from the detoxicating effect of high concentrations of glutamine in the growth medium.

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