REGULATION OF UNBALANCED REDOX HOMEOSTASIS INDUCED BY THE EXPRESSION OF WILD-TYPE HIV-1 VIRAL PROTEIN R (NL4-3VPR) IN FISSION YEAST

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The wild-type viral protein R (Vpr) of human immunodeficiency virus type 1 exerts multiple effects on cellular activities during infection, including the induction of cell cycle G_2 arrest and the death of human cells and cells of the fission yeast *Schizosaccharomyces pombe*. In this study, wild-type Vpr (NL4-3Vpr) integrated as a single copy gene in *S. pombe* chromosome was used to investigate the molecular impact of Vpr on cellular oxidative stress. NL4-3Vpr triggered an atypical response in early (14-h), and a well-regulated oxidative stress response in late (35-h) log-phase cultures. Specifically, NL4-3Vpr expression induced oxidative stress in the 14-h cultures leading, to decreased levels of superoxide anion (O_2^{-}), hydroxyl radical ('OH) and glutathione (GSH), and significantly decreased activities of catalase, glutathione peroxidase, glucathione reductase, glucose-6-phosphate dehydrogenase and glutathione S-transferase. In the 35-h cultures, elevated levels of O_2^{-} and peroxides were accompanied by increased activities of most antioxidant enzymes, suggesting that the Vpr-induced unbalanced redox state of the cells might contribute to the adverse effects in HIV-infected patients.

Keywords: HIV-1 - Vpr - oxidative stress - fission yeast - Schizosaccharomyces pombe

INTRODUCTION

The human immunodeficiency virus type-1 (HIV-1) viral protein R (Vpr) appears to be responsible for the progressive depletion of CD4⁺ lymphocytes in individuals with AIDS [18]. The wild-type Vpr (NL4-3Vpr) is localized in the mitochondria, nucleus

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Abbreviations: CAT, catalase; DHR 123, dihydrorhodamine 123; EPR, electron paramagnetic resonance; ET, ethidium bromide; GPx, glutathione peroxidase; GR, gluthathione reductase; GSH, glutathione; GSSG, glutathione disulfide; GST, gluthatione S-transferase; G6PD, glucose-6-phosphate dehydrogenase; HIV-1 Vpr, human immunodeficiency virus type-1 viral protein R; NL4-3Vpr, wild-type Vpr; PBN, N-*tert*-butyl-α-phenylnitrone; ROS, reactive oxygen species; *S. pombe, Schizosaccharomyces pombe;* SOD, superoxide dismutase

and cytoplasm [7], where it destroys the cellular regulation of the host: (i) it alters the cell morphology (elongated "cdc" phenotype cells) [24], (ii) it prevents cell proliferation by arresting cells in the G2 phase of the cell cycle [13, 24], (iii) it causes apoptosis through binding to the inner mitochondrial membrane and the release of cytochromes responsible for apoptosis [12], (iv) it activates caspase 8, the main effector of the receptor-mediated apoptotic pathway [27], (v) it causes oxidative stress in HIV-1-infected host cells [22, 28] and (vi) it stimulates the replication of HIV and the development of immunodeficiency [5, 14, 26]. We demonstrated earlier that the internal oxidative stress caused by wild-type Vpr in *Schizosaccharomyces pombe (S. pombe)* is partly suppressed by hydrogen peroxide (H₂O₂)-induced adaptive processes [3]. A mutation of wild-type Vpr may cause altered pathways and target molecules during HIV pathogenesis [16], indicating its structural importance. As an example, the C-terminal end of Vpr is crucial for G2 arrest, while the N-terminal α -helix is important for nuclear localization [7].

HIV-infected human deficiencies in the antioxidant enzyme system have been detected [8]. During HIV pathogenesis, reactive oxygen species (ROS) accumulate as biochemical by-products, causing an unbalanced redox state of the cells. The overproduction of ROS such as the superoxide anion (O_2^{-}) , hydroxyl radical (•OH) and peroxides (H₂O₂) may be related to an increased activation of polymorphonuclear leukocytes [2], DNA damage or impairment of the T-cell function during HIV infection [4]. A number of studies have reported semiquantitative measurements of the total ROS in HIV-infected cells [for a review, see 22]. Both enzymatic and nonenzymatic antioxidant molecules are responsible for maintaining the cellular oxidative balance, including superoxide dismutases (SODs), glutathione peroxidase (GPx), glutathione S-transferase (GST), glutathione reductase (GR), catalase (CAT), glucose-6-phosphate dehydrogenase (G6PD), glutathione (GSH), etc.

The aim of the present study was to determine ROS separately and to quantify the changes in the antioxidant system (e.g. GSH and the most important antioxidant enzymes) during the internal oxidative stress induced by wild-type Vpr in fission yeast. To acquire more detailed information, early-log (14-h) and stationary (35-h) phase cultures were analyzed. The data obtained with wild-type Vpr permits a comparison with our earlier published data on the mutant F34IVpr protein [28].

MATERIALS AND METHODS

Chemicals

All of the chemicals used in this study were of analytical grade and were bought from Sigma-Aldrich Ltd., Budapest, Hungary. Dihydroethidium was purchased from Fluka, Buchs, Switzerland.

Cell growth and gene expression in fission yeast cells

The NL4-3Vpr-carrying *S. pombe* strain RE007 (*leu1-32 ura4-D18 VprNL4-3::ura4+ade6-M210 h-*) and its parental strain SP223 (h-, leu1-32 ura4-294 ade6-216) were used. The processes of gene integration and induction under the control of the fission yeast *nmt1* promoter were described previously [20]. Cells were cultured at 30 °C in minimal growth medium (EMM) with 200 μ g ml⁻¹ and 75 μ g ml⁻¹ of the required amino acids and base [28], and were investigated in the early-log (14-h) and stationary (35-h) phases. Addition of 20 μ M thiamine to the growth medium caused 98% gene repression [20, 31, 32]. All experiments were carried out with the strain RE007 under gene-inducing and gene-repressing conditions in comparison with the parental strain SP223. Elimination of the effects of Vpr is not possible as the repression is not complete. The Vpr-repressed cultures are therefore not suitable as a control [20, 31, 32]. Taking this into account, we used the parental strain SP223 as a control. Other conditions and the microscopic examination of cells were as described earlier by Antal and Pesti [3].

Determination of living cell number, G2 arrest, septation index analysis, measurements of intracellular H₂O₂ and O₂ - concentrations, specific antioxidant enzyme activities and Cr(VI) reduction ability and formation of •OH

These methods have been described in details by Stromajer-Rácz et al. [28].



Fig. 1. Growth kinetics of *S. pombe* parental strain SP223 (●) and its NL4-3Vpr-carrying strain RE007 under gene-inducing (▼) and gene-repressing (■) conditions in EMM medium at 30 °C

RESULTS

Measurement of growth inhibition, G2 arrest and septation index

In the presence of thiamine (gene-repressing conditions), the NL4-3Vpr-carrying cells showed no significant inhibition of growth rate in comparison with the parental strain SP223 (Fig. 1). In the absence of thiamine (gene-expressing conditions), the NL4-3Vpr expression caused a 38.5% inhibition of growth in the 35-h EMM liquid medium in comparison with the parental strain SP223 (Fig. 1).

The cell cycle G2 arrest of RE007 was determined under *vpr*-expressing and *vpr*-repressing conditions in comparison with the parental SP223. Vpr induced 5% and 90% cell cycle G2 arrest (the elongated cells exhibited the "cdc" phenotype) of the



Fig. 2. Cell length as indicative of the cell cycle G2 arrest "cdc phenotype" in *S. pombe* parental strain SP223 and NL4-3Vpr derivative strain RE007 under gene-inducing conditions at 14 and 35 h. Bar represents 10 μm



Fig. 3. Septation indices of *S. pombe* parental strain SP223 (●) and its NL4-3Vpr derivative strain RE007 under gene-inducing (♥) and gene-repressing (■) conditions

Table 1
Summary of the effects of NL4-3Vpr on cellular enzymes involved in oxidative stress

	14 h		35	5 h
	SP223	RE007	SP223	RE007
ETa	1.04±0.05	0.46±0.09***	0.012±0.001	0.038±0.01***
GSH ^a	135.13±11.83	23.65±1.16***	41.04±6.99	51.24±4.44
GSSG ^a	0.68±0.06	0.10±0.05***	0.52±0.32	4.50±0.8***
GSH/GSSG	198.35	218.98	78.33	11.38*
SOD ^b	20.97±2.28	19.52±3.71	4.59±0.45	5.99±0.97*
SOD ^b Mn	1.47±0.38	2.93±0.8**	0.66±0.48	0.34±0.22
SOD ^b CuZn	19.50±5.46	16.59±5.51	3.93±2.92	5.64±3.71
CAT ^c	6.41±0.50	4.70±0.34***	6.82±0.34	8.11±2.15
GPxd	4.40±0.79	0.94±0.36***	3.01±0.48	6.43±0.7***
GRd	21.75±3.09	7.80±0.23**	30.07±1.68	43.66±4.81***
G6PD ^d	135.26±29.11	36.66±6.29***	206.26±22.44	253.98±7.94**
GST ^d	13.27±2.03	8.11±2.34***	18.67±1.92	30.59±2.39***

All cells were grown under gene-inducing conditions, i.e. without thiamine in the growth medium and enzymatic activities were measured at 14 and 35 h after the depletion of thiamine. Specific production values were expressed as means ± SD, calculated from the data of four independent experiments. ET: superoxide anion calibrated to ethidium bromide. *p < 5%; **p < 1%; **p < 0.1%. Values were calculated via the Student *t*-test. aSpecific concentrations are given in nmol (mg protein)⁻¹. bSpecific activities are given in unit (min mg protein)⁻¹. eSpecific activities are given in µmol (min mg protein)⁻¹. dSpecific activities are given in nmol (min mg protein)⁻¹.

NL4-3Vpr-repressing and NL4-3Vpr-expressing cells, respectively, in the 35-h cultures (Fig. 2), but not in the 14-h cultures (Fig. 2).

The septation index data showed that the parental strain SP223 displayed normal mitotic division, whereas the NL4-3Vpr-expressing RE007 cells (but not at the high level of the NL4-3Vpr-repressing cells) exhibited a G2 arrest with halted septation between 21 and 27 h of cell division (Fig. 3). Moreover, significantly decreased cell numbers, with a 77% and a normal 92% living cell rate were detected in the 35-h cultures of the *vpr*-expressing and control strains, respectively (data not presented). In this cell population, the majority of the *vpr*-expressing cells (90%) exhibited the "cdc" phenotype (Fig. 3).



Fig. 4. Time courses of dihydrorhodamine (DHR 123) oxidation of *S. pombe* NL4-3Vpr-expressing RE007 (\bigtriangledown) and its parental strain SP223 (\bullet) in 14-h and 35-h cultures. The conversion of DHR 123 by 5×10^6 cells ml⁻¹ at 21 °C was monitored in a Perkin-Elmer fluorimeter ($\lambda_{ex} = 488$ nm and $\lambda_{em} = 525$ nm). Data are from four independent measurements; the error was <7.5%

Measurements of intracellular H₂O₂ *and* O₂⁻⁻ *concentrations and specific antioxidant enzyme activities*

Increased or significantly increased intracellular H₂O₂ concentrations, as detected via the oxidation of dihydrorhodamine 123 (DHR 123) to rhodamine, were measured at 14 h and 35 h, respectively, in the NL4-3Vpr-expressing cells of strain RE007 (Fig. 4). The specific concentration of O₂⁻ calibrated to ethidium bromide (ET) for NL4-3Vpr-expressing cells was decreased in the 14-h cultures, but significantly increased in the 35-h cultures (Table 1). In comparison with the parental strain SP223, the GSH and GSSG concentrations were lower in the 14-h and higher in the 35-h cultures under NL4-3Vpr-expressing conditions (Table 1). These alterations were accompanied by a decreased and an elevated activity of GST in the 14-h and 35-h cultures, respectively (Table 1). NL4-3Vpr-expressing cells increased the activity of SOD_{Mn} in the 14-h cultures and those of the total SODs in the 35-h cultures (Table 1), and decreased activities of all the antioxidant enzymes (except the SODs) in the early-log-phase (14-h) samples and an up-regulating effect (except for CAT) in the late log-phase (35-h) samples (Table 1) in comparison with the parental SP223 cells. The 14-h cultures of NL4-3Vpr-expressing cells demonstrated decreased specific activities of CAT, GR, G6PD and GST. No changes were measured in the specific activities of SODs (Table 1).

Cr(VI) reduction ability and formation of 'OH

Samples of disrupted cells were treated for 10 min with $K_2Cr_2O_7$. An EPR spectroscopy signal arising from Cr(V) (Fig. 5) at g = 1.9554 indicated the rapid reduction of Cr(VI) to Cr(V). Significantly increased Cr(V) concentrations were detected under NL4-3Vpr-expressing conditions in both the 14-h and the 35-h cultures in compari-

Strain	Treatment	14 h		35 h	
		Cr(V)	PBN-OH	Cr(V)	PBN-OH
SP223	PBN + Cr(VI)	3.07	2.42	2.45	0.0045
RE007	PBN + Cr(VI)	3.68	0.48	3.34	0.0008
SP223	PBN + Cr(VI) + NADPH	56.09	0.64	6.45	0.0801
RE007	PBN + Cr(VI) + NADPH	18.80	0.16	5.88	0.1540

 Table 2

 Cr(V) and PBN-OH spin adduct production of disrupted parental strain SP223 and its NL4-3Vpr-carrying strain RE007

All cells were grown under the gene-inducing conditions, i.e. without thiamine in the growth medium and samples were taken 14 and 35 h after the depletion of thiamine. Cr(V) and PBN-OH spin adduct production was measured by EPR. Concentrations of Cr(V) and PBN-OH are given in μ M. The SD of the mean values was <10%. A representative EPR spectrum is to be seen in Fig. 5.



Fig. 5. Representative EPR spectra of disrupted cells of S. pombe parental strain SP223 (-) and its NL4-3Vpr-carrying strain RE007 (- -) under gene-inducing conditions at 14 h, recorded 10 min after mixing in 0.1 M PBN-containing buffer solution (pH 7.2), 2 mM K₂Cr₂O₇ and 2 mM NADPH. The field scan was 10 mT. Numerical data are to be seen in Table 2

son with the parental strain SP223 (Table 2). However, the addition of both 2 mM $K_2Cr_2O_7$ and 2 mM NADPH resulted in an approximately 3-fold decrease in the Cr(VI) concentration in the 14-h cultures (Table 2), suggesting that an unbalanced redox state of the cells had developed by this time point. The addition of Cr(VI) and NADPH resulted in decreased and increased 'OH production, respectively, in the cells of RE007.

DISCUSSION

NL4-3Vpr expression-induced growth inhibition, cell cycle G2 arrest and change in septation index

The expression of NL4-3Vpr caused an inhibition (38.5%) of growth in the 35-h cultures in comparison with the parental strain SP223, demonstrating the inhibitory effect of cell multiplication of this protein. These data are in agreement with earlier observations [3, 32]. For comparison, the multiplication of the mutant F341Vpr-expressing cells stopped at 20 h and started again at 26 h [28].

It has been demonstrated previously that the expression of wild-type NL4-3Vpr in S. pombe induces elongated cells that are delayed or arrested at the G2/M cell cycle transition [3, 29, 30]. The G2 arrest mediated by Vpr utilizes the cellular signaling pathway whose physiological function is to recognize replication stress and which is strongly associated with the oxido-reduction balance of cells [33]. In our work, the Vpr-induced G2 arrest was confirmed microscopically and by determination of the septation index based on the cell fraction in a population with septum, indicating mitotic cell division (Figs 2 and 3). A rapid decrease in septum formation is detected when a G2 arrest occurs in comparison with actively growing controls, around 10 to 20% of the cells containing septa [21]. From this aspect, our data are in line with the earlier immunoblot findings that the vpr expression starts at 15-17 h, leading to morphological changes in the 17-h cultures [19]. The G2/M cell cycle block may be detected in the 35-h cultures, leading to apoptosis preceded by morphological changes [1, 30]. Chang et al. [6] found that wild-type *vpr*-expressing S. pombe cells began to exhibit elongation at 19 h and were lengthened progressively at 24 h. Morphological changes were observed at 20 h in mutant F34Ivpr-expressing S. pombe cultures, in which 20% of the cells were in the mitotic phase, with more than half displaying elongated morphology [28].

Previously presented data suggested that the *vpr*-repressing cultures included a small percentage of cells in which expression of the Vpr protein was observed [28]. In our work, two ages of cultures were used, so as to allow a more accurate analysis of the NL4-3Vpr function. In the early-log-phase (14-h) of the cultures, visible morphological changes could not be detected, but the morphological changes were already manifested in the stationary-phase cells (35 h) because of the completion of G2/M blocking.

NL4-3Vpr-induced depletion of GSH and increased intracellular ROS production

It was proved that the GSH and GSSG concentrations increased with the age of the cultures, due to the increasing NL4-3Vpr expression. The elevated activity of GST in the older cultures may be a consequence of its function (GST exports GSSG from the cells if the ratio GSH/GSSG is unbalanced) [15].

An increase in ROS production was observed in the NL4-3Vpr-expressing strain RE007 as compared with its parental strain SP223. The specific concentration of O_2^{+} for the NL4-3Vpr-expressing cells was decreased in the 14-h cultures, but significantly increased in the 35-h cultures. Surprisingly, the wild-type Vpr-induced H₂O₂ concentration was only about half of that induced by the mutant F341Vpr [28]. Our results are in good agreement with the earlier findings that the wild-type Vpr stimulated elevated O_2^{+-} and total ROS production in *S. pombe* cells, as observed semi-quantitatively by a fluorescence method [25].

The 35-h cultures exhibited elevated antioxidant enzyme activities (except for the SODs and GPx), in agreement with the findings of Lee et al. [17], explaining the increased oxidative stress tolerance of the older cells. These alterations in the NL4-3Vpr-expressing cells might results from the decreased concentration of GSH, the changed concentration of H_2O_2 and the disturbed regulation of the oxidative stress process via the MAPK-regulated transcription factors *pap1* and *atf1*. Since, the specific activities of all the examined antioxidant enzymes in the NL4-3Vpr-expressing cells were up-regulated and the concentration of GSH, as the first indicator of oxidative stress [11], was elevated in comparison with that in the parental strain SP223 in the 35-h cultures, the results allow the presumption that, under the wild-type NL4-3Vpr-induced conditions, the cells were able to adapt to the oxidation stress conditions due to the elevated levels of O_2^{-} and H_2O_2 through the up-regulation of the activities of all the antioxidant enzymes.

NL4-3Vpr induced a decreased 'OH concentration in spite of the higher intracellular O₂- and H₂O₂ levels

The redox state of cells can be characterized by the simultaneous measurement of the reduction of Cr(VI) to Cr(V) and the 'OH concentration by EPR spectroscopy. Cr(VI)is reduced by antioxidants (GSH, etc.) and the GR/NADPH system, and hence the quantity of Cr(V) indicates the total reduction capacity of the cell [11, 23]. The 'OH concentration measured as a N-tert-butyl- α -phenylnitrone-OH (PBN-OH) adduct provides information about the Fenton $(Cr(V)+H_2O_2 \rightarrow Cr(VI)+OH+OH^-)$ and Haber-Weiss $(O_2^{\bullet} + H_2O_2 \rightarrow OH + OH)$ reactions [11]. These data support the idea that Vpr-expressing cells are in an oxidized redox state [23]. In agreement with the results obtained with mutant F341Vpr-expressing cells [28], significantly decreased •OH production was detected in the NL4-3Vpr-expressing cells after the addition of Cr(VI) in both the 14-h and the 35-h cultures, in spite of the high concentrations of O_2 (in the 35-h cultures) and H_2O_2 (in both cultures). These phenomena might be consequences of a disrupted metal homeostasis (decreased concentrations of redoxactive metals, e.g. Fe²⁺ and Cu²⁺) of strain RE007 [10]. As already suggested, this may be an atypical reaction because the addition of NADPH to the Cr(VI)-tolerant strain of S. pombe resulted in increased 'OH production [9].

Comparison of the data relating to the 14-h and 35-h cultures of both strains permits the conclusion in the case of the Vpr-expressing strain that higher oxidative stress was observed in the older cultures due to the already occurred Vpr expression in the cells. The decreased 'OH production and SOD activities and the increased concentrations of O_2 - and H_2O_2 and the CAT activity suggest a slower Fenton reaction, which may be caused by disrupted metal homeostasis. The high level of O_2 definitely causes DNA oxidative damage, which may also lie in the background of the apoptotic and G2 arrest phenomena. In the case of the parental strain, this correlation

was not observed, but a slight increase of the H_2O_2 was detected in consequence of the aging of the cultures.

To summarize, the expression of the wild-type Vpr caused an unbalanced redox state in strain RE007 in both the 14-h and the 35-h cultures in comparison with the parental strain SP223. In both cultures, the changes in the concentrations of the ROS were accompanied by changes in the GSH concentration and in the activities of specific antioxidant enzymes, which indicates a partial adaptation of the cells since the fenton reaction seems to be slowed down as a consequence of NL4-3Vpr cells adapted to the elevated concentration of H₂O₂ and O₂⁻⁻, intensifying the activity of the antioxidant enzyme system (except the SODs) in the recovery from the Vpr-induced unbalanced redox state. A typical regulation of detoxification could be observed. In the system of GSH/ROS/antioxidants, e.g. in the 35-h cultures, the increased concentration of H₂O₂ implies alterations in the activities of the detoxification enzymes CAT and GPx.

A comparison of the results with the mutant F341*vpr*-expressing strain [28] led to the conclusion that the mutant F341*vpr* gene displays a more pronounced response to oxidative stress and a higher adaptation than those for the wild-type *vpr*-expressing cells.

The results of this study may contribute to the knowledge relating to redox control, which can be an important therapeutic strategy for oxidative stress-associated disorders during HIV infection.

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