SLX4-SLX1 Protein-independent Down-regulation of MUS81-EME1 Protein by HIV-1 Viral Protein R (Vpr)**

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Evolutionarily conserved structure-selective endonuclease MUS81 forms a complex with EME1 and further associates with another endonuclease SLX4-SLX1 to form a four-subunit complex of MUS81-EME1-SLX4-SLX1, coordinating distinctive biochemical activities of both endonucleases in DNA repair. Viral protein R (Vpr), a highly conserved accessory protein in primate lentiviruses, was previously reported to bind SLX4 to mediate down-regulation of MUS81. However, the detailed mechanism underlying MUS81 down-regulation is unclear. Here, we report that HIV-1 Vpr down-regulates both MUS81 and its cofactor EME1 by hijacking the host CRL4-DCAF1 E3 ubiquitin ligase. Multiple Vpr variants, from HIV-1 and SIV, down-regulate both MUS81 and EME1. Furthermore, a C-terminally truncated Vpr mutant and point mutants R80A and Q65R, all of which lack G₂ arrest activity, are able to downregulate MUS81-EME1, suggesting that Vpr-induced G₂ arrest is not correlated with MUS81-EME1 down-regulation. We also show that neither the interaction of MUS81-EME1 with Vpr nor their down-regulation is dependent on SLX4-SLX1. Together, these data provide new insight on a conserved function of Vpr in a host endonuclease down-regulation.

Primate lentiviruses encode several common accessory proteins, including viral protein R (Vpr),² viral protein X (Vpx), viral protein U, negative regulatory factor, and viral infectivity factor. These proteins were once thought to be non-essential but were later found to be critical for *in vivo* viral replication (1, 2). Among these proteins, Vpr and Vpx are closely related and similar in size, at about 100 amino acids (3). Vpr is present in all types of primate lentiviruses, including HIV-1, HIV-2, and SIV, although Vpx is only found in HIV-2 and some SIV strains (1, 4, 5).

Vpr, as a virion-associated protein, is implicated in many processes that promote HIV-1 infectivity. It is packaged into viral particles and has been suggested to be involved in nuclear import of the viral pre-integration complex, a key step for successful infection in non-dividing cells such as macrophages (6-9). Vpr has been implicated as a transcriptional activator and viral replication activator (10-12). Vpr was also reported to induce T-cell apoptosis and to induce differentiation in a rhabdomyosarcoma cell line (13-17).

Vpr expression in cycling cells results in G₂/M cell cycle arrest by inducing the ataxia telangiectasia and Rad3-related (ATR)-mediated DNA-damage checkpoint, which may favor viral replication (18-21). However, the detailed mechanism of Vpr-induced G₂/M arrest is not yet clear. Several lines of evidence suggest that G₂ arrest requires interaction of Vpr with the CRL4 E3 ubiquitin ligase complex, comprising DDB1, CUL4, and RBX1, via its cellular partner DCAF1 (22-30). DCAF1 is a substrate receptor component of the CRL4 (31–33). One plausible hypothesis is that Vpr usurps the CRL4-DCAF1 complex to promote down-regulation of unknown cell cycle regulatory factors via the proteasome-dependent pathway (34-36). In this case, down-regulation of the unknown factor(s) would activate ATR-mediated checkpoint signaling and induce G₂ arrest. In line with this hypothesis, mutagenesis studies have shown that a Vpr mutant with reduced DCAF1 binding, Q65R, lacks the ability to induce G_2 arrest (22, 25, 28, 37). Furthermore, single mutation at Arg-80 (R80A) results in abrogation of Vpr-mediated G₂ cell cycle arrest without affecting DCAF1 interaction (10, 22, 25, 38-40). These reports suggest that the unidentified cell cycle regulatory factor(s) is recruited to the CRL4-DCAF1 complex via the C terminus of Vpr.

The MUS81-EME1 and MUS81-EME2 complexes are conserved DNA structure-selective endonucleases and play essential roles in homologous recombination and replication fork processing and repair (41, 42). The C-terminal regions of EME1 (total of 570 residues) and EME2 (total of 379 residues) are highly homologous with more than 60% sequence similarity in human (43, 44). Multiple studies have suggested that these complexes are involved in maintenance of genomic stability and damaged DNA processing (43-47). Purified recombinant MUS81-EME1 recognizes and preferentially cleaves several DNA structures, including 3'-flap, aberrant replication forks, and nicked Holliday junctions (45, 48). Loss of MUS81 or EME1, or their disruption, results in genomic instability and hypersensitivity to DNA cross-linking agents such as mitomycin C and platinum compounds, due to DNA repair defects (49, 50).

Together with another structure-selective endonuclease, SLX4-SLX1, MUS81-EME1 defines one of two major Holliday junction resolution pathways in human cells (51, 52). SLX4 provides a scaffold for multiple factors, and its C terminus binds



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² The abbreviations used are: Vpr, viral protein R; SIV, simian immunodeficiency virus; Vpr^{HIV-1} NL⁴⁻³, Vpr isolated from HIV-1 NL4-3 clone; Vpx^{SIVmac}, Vpx isolated from SIV infecting macaque monkey; Vpx, viral protein X.

the catalytic SLX1 to form an SLX4-SLX1 complex. SLX4-SLX1 further associates with MUS81-EME1, forming a stable holoenzyme termed SLX4com (51). The role of MUS81-EME1 in Holliday junction resolution and DNA interstrand cross-link repair requires its interaction with SLX4 in human cells (51–53).

Recently, human SLX4com, comprising SLX4, SLX1, MUS81, and EME1, was reported to be a target of HIV-1 Vpr (54). In particular, Vpr was shown to directly bind SLX4 and was postulated to untimely activate the SLX4com via the CRL4-DCAF1 complex. These interactions were proposed to lead to aberrant DNA cleavage and cell cycle arrest (54). Additionally, MUS81 was reported to be down-regulated by Vpr, which was further confirmed by another two groups (55, 56). However, the detailed mechanisms of MUS81 down-regulation and its relation to Vpr-induced G2 arrest are not clear. In the work described here, we report Vpr-mediated down-regulation of both MUS81 and its partners EME1 and EME2. Vpr executes this task by hijacking host CUL4-DCAF1 E3 ubiquitin ligase and targeting MUS81-EME1 and MUS81-EME2 for degradation in a proteasome-dependent manner, which could be inhibited by MG132. Multiple Vpr proteins, from either HIV-1 or SIV isolated from chimpanzee, down-regulate MUS81 and EME1, suggesting that it is a conserved function of Vpr. Strikingly, we further show that MUS81-EME1 down-regulation does not require SLX4-SLX1. We also find that Vpr mutants defective for G₂ arrest are still capable of down-regulating MUS81-EME1, suggesting that G_2 arrest is not correlated with MUS81-EME1 down-regulation.

Results

HIV-1 Vpr but Not SIV Vpx Down-regulates MUS81-To confirm previous reports of Vpr-induced down-regulation of MUS81, we co-transfected MUS81 and HIV-1 NL4-3 Vpr plasmids into HEK293T cells. As shown in Fig. 1A, the level of MUS81 was dramatically decreased in a Vpr dose-dependent manner. Because MUS81 forms a complex with its cofactor EME1 (or EME2), we wondered whether EME1 influences the stability of MUS81. We therefore co-transfected EME1 with MUS81 (Fig. 1B). The level of MUS81 increased in accordance with the amount of EME1-expressing plasmid (Fig. 1B, lanes 1-4). When Vpr was present, the level of MUS81 also increased with EME1 (Fig. 1*B*, *lanes* 5-8); however, the level of MUS81 was always lower in the presence of Vpr. These results suggest that MUS81 is down-regulated by Vpr irrespective of its quaternary state. We also interrogated whether endogenous MUS81 could be down-regulated by Vpr. When Vpr alone was transfected into HEK293T cells, endogenous MUS81 indeed decreased with increasing amounts of Vpr (Fig. 1C).

Because Vpr and Vpx are closely related and have evolved from a common ancestor, we investigated whether Vpx could also down-regulate MUS81 (Fig. 1*D*). The level of MUS81 was not affected by co-transfection of Vpx (Vpx^{SIVmac}) from SIV that infects the macaque monkey. The level of human SAMHD1, a known target of Vpx^{SIVmac}, however, was significantly lower with increasing amounts of Vpx^{SIVmac}. Together, these results suggest that MUS81 is down-regulated by Vpr^{HIV-1 NL4-3} but not by Vpx^{SIVmac}. To further investigate Vpr-dependent down-regulation of MUS81, chase assays were carried out with an inhibitor of *de novo* protein synthesis, cycloheximide. In the presence of cycloheximide, the intensity of the band corresponding to MUS81 decreased significantly over an 8-h period (Fig. 1*E*). Importantly, the same band disappeared more rapidly in the presence of Vpr. Vpr also decreased after cycloheximide treatment. Together, these results suggest that Vpr enhances down-regulation of MUS81.

Both MUS81 and Its Co-factor EME1 or EME2 Are Downregulated by HIV-1 Vpr-MUS81 associates with EME1 or EME2 and functions as a DNA structure-selective endonuclease in mammalian cells. Therefore, we asked whether EME1 and EME2 are also down-regulated in a Vpr-dependent manner, along with MUS81. HEK293T cells were transiently cotransfected with MUS81, EME1 (or EME2), and VprHIV-1 NL4-3expressing plasmids, and protein levels were interrogated 48 h after transfection. In agreement with previous results, MUS81 decreased in a Vpr dose-dependent manner. As shown in Fig. 2, A and B, both EME1 and EME2 were down-regulated by Vpr^{HIV-1 NL4-3}. Endogenous MUS81, EME1, and EME2 were also down-regulated by the Vpr (Fig. 2C). Because EME1 and EME2 behaved similarly in response to $Vpr^{\rm HIV-1\;NL4-3}$ co-expression, the remainder of our studies focused on MUS81-EME1.

To explore whether Vpr proteins from other strains also function in both MUS81 and EME1 down-regulation, we examined the cellular levels of MUS81 and EME1 after co-transfection with Vpr from viruses isolated from diverse species, including chimpanzee SIV PTT and PTS (Fig. 2*D*), and HIV-1 group M, N, and O (Fig. 2*E*). Vpr from all these strains efficiently downregulated both MUS81 and EME1. Sequence analysis of these Vpr proteins suggests a conserved N-terminal and central region (Fig. 2*F*). The C-terminal region of these Vpr proteins, however, is variable. This may imply that the C-terminal part is not important for MUS81-EME1 down-regulation.

MUS81-EME1 Down-regulation Is Mediated via a Ubiquitin/Proteasome-related Pathway-To determine whether MUS81-EME1 down-regulation is mediated via a ubiquitin/ proteasome-related pathway, we treated MUS81-EME1/Vpr co-transfected cell cultures with MG132, an inhibitor of the proteasome. We used VprHIV-1 NL4-3 for this experiment and the remainder of our studies. MG132 effectively restored MUS81 and EME1 levels (Fig. 3A). This suggests Vpr-dependent down-regulation of MUS81-EME1 requires the proteasome. The level of Vpr increased with MG132 treatment. This is consistent with the previous observation that Vpr is also degraded by the proteasome (28). To further examine the role of the proteasome-dependent pathway, we co-expressed ubiquitin with MUS81, EME1, and Vpr. As shown in Fig. 3B, ubiquitin expression resulted in a strong poly-ubiquitination signal in whole cell extracts. Immunoprecipitation with an anti-FLAG antibody (Fig. 3C), to pull down FLAG-tagged MUS81, revealed stronger poly-ubiquitination signals (lane 2) in the presence of Vpr. Taken together, these results suggest that Vpr induces down-regulation of MUS81-EME1 via ubiquitination and a proteasome-dependent pathway.





FIGURE 1. **Vpr but not Vpx down-regulates MUS81.** *A*, MUS81 was transiently co-transfected with increasing amounts of Vpr^{HIV-1} NL4-3</sup>. At 48 h post-transfection, cells were harvested and analyzed by Western blotting. *B*, MUS81 was co-transfected with increasing amounts of EME1 and empty vector or Vpr^{HIV-1} NL4-3</sup> plasmids. The same amounts of MUS81 and EME1 plasmids were used for *lanes 1* and *5*, *lanes 2* and *6*, *lanes 3* and *7*, and *lanes 4* and *8*. *C*, HEK293T cells were transfected with increasing amounts of Vpr^{HIV-1} NL4-3</sup> plasmid. The endogenous level of MUS81 was analyzed by Western blotting. *D*, MUS81 or SAMHD1 was co-transfected with an increasing amount of Vpx^{SIVmac}. *E*, HEK293T cells were incubated with 20 μ g/ml cycloheximide, 40 h after transfection with the indicated plasmids. The levels of MUS81 and Vpr were analyzed by Western blotting after the indicated times. The intensity of bands corresponding to MUS81 and Vpr were quantified from four independent experiments.

Vpr usurps the CRL4-DCAF1 E3 ubiquitin ligase to downregulate its host cellular targets via the ubiquitination/proteasome pathway. Previous studies by Laguette *et al.* (54) suggested that Vpr uses the CRL4-DCAF1 E3 ubiquitin ligase to down-regulate MUS81. Thus, we investigated the role of CRL4-DCAF1 in Vpr-induced MUS81 down-regulation. Transient co-expression of DCAF1 resulted in significant enhancement of Vpr-induced MUS81 down-regulation (Fig. 4*A*). To further confirm the role of DCAF1 in MUS81-EME1 down-regulation, we performed titration of DCAF1 plasmid with a stable amount of control plasmid or Vpr plasmid (Fig. 4*B*). In the absence of Vpr, the levels of MUS81 and EME1 were similar, despite the increasing level of DCAF1; however, when Vpr was present, DCAF1 indeed dramatically promoted down-regulation of both MUS81 and EME1. Therefore, DCAF1 and Vpr cooperate in MUS81-EME1 down-regulation. Of note, consistent with a previous report (28), overexpression of DCAF1 resulted in stabilization of Vpr. Furthermore, knockdown of endogenous DCAF1 by siRNA reduced Vpr-dependent down-regulation of MUS81 (Fig. 4*C*).





FIGURE 2. **Both MUS81 and its cofactors EME1 and EME2 are down-regulated by Vpr from multiple species.** *A* and *B*, MUS81 and EME1 plasmids (*A*) or MUS81 and EME2 plasmids (*B*) were transiently co-transfected with increasing amounts of Vpr^{HIV-1 NL4-3} plasmid. At 48 h post-transfection, cells were harvested and analyzed by Western blotting. *C*, endogenous levels of MUS81, EME1, and EME2 were analyzed with increasing amounts of Vpr^{HIV-1 NL4-3} plasmids. At 48 h post-transfection, cells were harvested and EME1 plasmids were co-transfected with increasing amounts of plasmid encoding Vpr isolated from SIV that infects chimpanzee, Vpr^{cpz-PTT} and Vpr^{cpz-PTT} (*D*), or encoding Vpr from HIV-1 group M, N, or O, Vpr^{HIV-1 M}, Vpr^{HIV-1 N}, and Vpr^{HIV-1 O} (*E*). All experiments were repeated three times with equivalent results. *F*, sequence alignment of the Vpr proteins used in *A*–*D* was performed with Clustal Omega (68) and colored by ESPript 3.0 (69).

To further confirm the role of DCAF1 in MUS81-EME1 down-regulation, we performed co-immunoprecipitation experiments with FLAG-tagged MUS81 (Fig. 4*D*). Interestingly, DCAF1 associated with the MUS81-EME1 complex even in the absence of Vpr (Fig. 4*D*, *right-panel*, *lane* 9). Vpr, however, required DCAF1 to associate with MUS81-EME1 (Fig. 4*D*, *right panel*, compare *lanes* 8 and 10). These results suggest that DCAF1 is a natural partner of MUS81-EME1 and bridges the interaction between MUS81-EME1 and Vpr. To further examine the association of components of the CRL4 E3 ubiquitin ligase in this complex, we co-expressed DDB1, DCAF1, Vpr, and MUS81-EME1. As shown in Fig. 4*E*, DDB1, the substrate adaptor of the CRL4, could also be detected in the MUS81-EME1 co-immunoprecipitation fractions, regardless of whether Vpr was present. Taken together, the MUS81-EME1

complex interacts with the CRL4-DCAF1 E3 ubiquitin ligase and is down-regulated by the same ligase subverted by Vpr.

 G_2 Arrest and MUS81-EME1 Down-regulation Are Separate Functions of HIV-1 Vpr—Vpr proteins from different viral strains, with diverse C termini, all retained their ability to down-regulate MUS81-EME1 (Fig. 2). We thus wondered whether the Vpr C-terminal domain is required for MUS81-EME1 down-regulation. We first tested MUS81-EME1 down-regulation with a C-terminally truncated mutant Vpr (residues 1–79). As shown in Fig. 5A, the MUS81-EME1 complex was efficiently down-regulated by Vpr(1–79). To further explore the interaction between Vpr(1–79) and MUS81-EME1, we performed a MUS81 co-immunoprecipitation experiment (Fig. 5B). Vpr(1–79) was observed to coimmunoprecipitate with MUS81 (Fig. 5B, lane 5). Thus, res-





FIGURE 3. Vpr mediates MUS81-EME1 down-regulation via the ubiquitin/proteasome pathway. *A*, MUS81 and EME1 plasmids were co-transfected with empty vector plasmid or Vpr plasmid. At 42 h post-transfection, DMSO or MG132 (10 μ m final concentration) in DMSO were added to the cell culture. Six hours later, cells were harvested and analyzed by Western blotting. *B*, MUS81 and EME1 plasmids were co-transfected with empty vector or Vpr and/or ubiquitin plasmid as indicated. Cells were treated with MG132 (10 μ m final concentration) at 42 h post-transfection for 6 h. *C*, cell lysates from *B* were subjected to co-immunoprecipitation (*IP*) with anti-FLAG antibody, and elution fractions from co-immunoprecipitation with FLAG peptides were analyzed by Western blotting. All experiments were repeated three times with similar results.

idues 80–96 of Vpr are not required for MUS81-EME1 interaction and down-regulation.

Because Vpr residues in the C-terminal region are important for its G_2 arrest activity, we asked whether Vpr-induced G_2 arrest activity is independent of MUS81-EME1 down-regulation. We thus tested a Vpr mutant, R80A, that is deficient in inducing cell cycle arrest (10, 22, 38, 39). The R80A mutant still down-regulates MUS81-EME1 to levels comparable with the wild type protein (Fig. 5C). Furthermore, Vpr R80A associates with MUS81-EME1 as well as DDB1/DCAF1 (Fig. 5D). This suggests the function of MUS81-EME1 down-regulation by Vpr is independent of its activity in G₂ arrest. An analysis of Vpr Q65R, another mutant defective in G_2 arrest (22, 25, 28, 37), further supports this conclusion (Fig. 5, *E* and *F*). Vpr Q65R was previously shown to be impaired in DCAF1 binding (22, 25, 26). This was confirmed in our co-immunoprecipitation assay (Fig. 5G). Thus, interaction of Vpr with MUS81-EME1 and DCAF1 may overcome the negative effect of Q65R mutation in DCAF1 binding (Fig. 5*F*).

To ensure Vpr mutants are defective in cell cycle arrest activity, HEK293T cells were transiently transfected with empty vector plasmids, Vpr WT, residues 1–79, R80A, and Q65R (Fig. 5, H-L). Cells expressing Vpr WT showed significant accumulation at G₂/M phase. In contrast, Vpr mutants had minimal to moderate effects on cell cycle arrest.

SLX4 Is Not Required for DDB1-DCAF1-Vpr-MUS81-EME1 Complex Formation—Previous studies by Laguette *et al.* (54) suggested that Vpr directly binds the C-terminal region of SLX4. Because MUS81-EME1 directly associates with SLX4-SLX1, SLX4 may mediate the interaction between MUS81-EME1 and CRL4-DCAF1-Vpr. In this case, Vpr may promote MUS81-EME1 and DDB1-DCAF1 complex formation via

SLX4, and co-expression of SLX4 would result in facilitated Vpr-dependent down-regulation of MUS81-EME1. To address this possibility, a C-terminal domain of SLX4 (residues 1430-1834), competent for Vpr, SLX1, and MUS81-EME1 binding (54, 57), was co-transfected with MUS81-EME1 with or without Vpr (Fig. 6A). Surprisingly, co-expression of SLX4 did not promote MUS81-EME1 down-regulation. In fact, with increasing amounts of SLX4, the levels of MUS81 and EME1 were restored even when Vpr was present. This result does not support the model that Vpr down-regulates MUS81-EME1 through SLX4. On the contrary, SLX4 competes with CRL4-DCAF1-Vpr for MUS81-EME1. In other words, MUS81-EME1 down-regulation by Vpr in our system is likely independent of the SLX4/Vpr interaction. However, the truncated SLX4 may sequester Vpr or CRL4-DCAF1-Vpr and interfere with MUS81-EME1 down-regulation. Nevertheless, siRNA-mediated knockdown of SLX4 did not affect down-regulation of MUS81 and EME1 in the presence of Vpr (Fig. 6B).

To further test this hypothesis and to minimize the influence of unknown factors present in cells, we performed an *in vitro* pulldown assay with purified recombinant proteins (Fig. 6*C*). GST fusion DDB1 in complex with DCAF1 was assayed for MUS81-EME1 binding with or without Vpr (Fig. 6*D*). We detected an interaction between DDB1/DCAF1 and MUS81-EME1 only in the presence of Vpr. Taken together, these results suggest Vpr-mediated down-regulation of MUS81-EME1 does not require SLX4-SLX1.

Discussion

Despite being classified as an accessory factor, Vpr has multiple essential functions in the HIV-1 life cycle (4). Vpr primarily executes its biological activity by engaging DCAF1, a sub-



FIGURE 4. **CRL4-DCAF1E3 ubiquitin ligase is involved in Vpr-mediated MUS81-EME1 down-regulation.** *A*, MUS81 and EME1 plasmids were co-transfected with increasing amounts of Vpr or balanced empty vector plasmid and empty vector or DCAF1 plasmids as indicated. 48 h later, cells were harvested, and some lysates were analyzed by Western blotting with appropriate antibodies. *B*, MUS81 and EME1 plasmids were co-transfected with increasing amounts of DCAF1 or balanced empty vector or Vpr plasmids. *C*, DCAF1 siRNA or control siRNA was transfected 24 h prior to co-transfection of Vpr, MUS81, EME1, or balanced empty vector plasmids as indicated. *D*, MUS81 and EME1 plasmids were co-transfected with Vpr and/or DCAF1 plasmids, which were balanced by empty vector plasmids. 42 h later, MG132 (10 μ M final concentration) in DMSO or DMSO only was added to the cell culture. After 6 h, cells were harvested, and some lysates were analyzed by Western blotting. *Co-immunoprecipitation (IP)* with anti-FLAG antibody was performed and analyzed by immunoblotting. *E*, plasmids expressing MUS81, Vpr, DCAF1, and DDB1 were co-transfected into cells as indicated. Protein levels in cell lysates and co-immunoprecipitation were analyzed as described above. All experiments were repeated three times with similar results.

strate receptor component of the CRL4 E3 ubiquitin ligase (58). Current and prevailing data support the hypothesis that Vpr recruits cellular targets to the CRL4-DCAF1 E3 ubiquitin ligase for proteasome-dependent degradation. Previous reports by Laguette et al. (54) suggested that SLX4com, comprising SLX4, SLX1, MUS81, and EME1, is a target of Vpr and is loaded onto the CRL4-DCAF1 E3 ubiquitin ligase. In particular, the recruitment of SLX4com to DCAF1 by Vpr was shown to enhance the structure-specific endonuclease activity of MUS81-EME1. Knockdown of any component of SLX4com abolished Vpr-induced G₂ arrest. Furthermore, Vpr mutants, previously shown to be defective in G₂ arrest, failed to form a stable complex with SLX4com and DCAF1 (54). These data implied that hyperactivity of MUS81-EME1, induced by Vpr, would produce an abnormal number of DNA breaks, turning on the DNA damage repair pathway, and ultimately leading to G₂ arrest. A subsequent independent study further supported an interaction between SLX4com and CRL4-DCAF1 subverted by Vpr (55). Berger *et al.* (55) investigated the G_2 arrest phenotypes of a panel of SIV Vpr proteins in human and simian grivet cells and established a correlation between G2 arrest and Vpr/SLX4com interaction. Interestingly, association of SLX4com with CRL4-DCAF1-Vpr was found to down-regulate MUS81. A possible

model arising from these observations argues that Vpr interacts with SLX4com via the C-terminal region of SLX4 (Fig. 7A). Spatial arrangement of SLX4com allows poly-ubiquitination of MUS81, mediated by CRL4-DCAF1-Vpr, resulting in its proteasome-dependent degradation (54, 55). In this report, we provide evidence that MUS81-EME1 interacts with the CRL4-DCAF1 complex irrespective of Vpr and that the level of the endonuclease is regulated by Vpr binding to CRL4-DCAF1 and MUS81-EME1 (Fig. 7B). Co-immunoprecipitation experiments showed Vpr-independent interaction between DCAF1 and MUS81-EME1 (Fig. 4, D and E), whereas an in vitro pulldown experiment with purified recombinant proteins suggested a role for Vpr in mediating interaction between DCAF1 and MUS81-EME1 (Fig. 6, C and D). This suggests that an unknown cellular factor(s) may facilitate association of these proteins in cells. Surprisingly, Vpr-mediated down-regulation of MUS81/1EME1 does not require SLX4-SLX1.

Down-regulation of MUS81-EME1 appears to be an evolutionarily conversed function of Vpr. We show that Vpr proteins isolated from multiple HIV-1 groups are capable of downregulating MUS81-EME1. Furthermore, Vpr proteins from SIVcpz virus, isolated from two chimpanzee species, also induced MUS81-EME1 down-regulation. These SIV Vpr pro-







teins are ancestors to those from the HIV-1 virus, generated by zoonosis (59, 60). These observations suggest that removal of the structure-selective endonuclease activity of MUS81-EME1 may be essential for the HIV-1 life cycle. However, our study suggests that this activity is not related to induction of cell cycle arrest at G₂, one of the most well documented Vpr-dependent phenotypes. It is well established that the C terminus of Vpr is required for G_2 arrest, arguing that the C terminus may bind and load cell cycle-related factors onto the CRL4-DCAF1 E3 ubiquitin ligase for proteasome-dependent degradation (10, 22, 25, 38-40). Several lines of evidence in this report argue that MUS81-EME1 is recruited to CRL4-DCAF1 by Vpr, but not via its C terminus. First, the Vpr variants that we tested for MUS81 down-regulation show significant diversity at the C terminus (Fig. 2F). Second, Vpr mutants, defective in G₂ arrest still downregulate MUS81-EME1 (Fig. 5). Consistent with our observation, another report showed that the same Vpr mutants, Q65R and R80A, caused MUS81 degradation (56). Further exploration is required to understand how Vpr mediates G₂ arrest by interacting with SLX4com, and the independent biological pathway of MUS81 down-regulation, and the roles of these processes in the HIV-1 life cycle.

DCAF1 is an essential component in cell cycle progression and DNA replication, and its functions are mainly ascribed to its association with the CRL4 E3 ubiquitin ligase (58). Significantly, DCAF1 associates with chromatin during early S and G₂ phases, and silencing of DCAF1 limits the cells progressing through the S phase (61). Therefore, it is not surprising to find that DCAF1 associates with several chromatin-bound factors including histone H2A, p53, HDAC1, and TET2/3, modulating transcriptional activation of specific genes (62-65). Therefore, it is tempting to speculate that DCAF1 regulates the cellular activity of MUS81-EME1 as well as SLX4com. It will be interesting to investigate how DCAF1 affects the biochemical activities of these structure-specific endonucleases, including resolution of Holliday junctions and repair of DNA interstrand cross-links. Understanding these processes may provide clues about the pathological advantage of MUS81 down-regulation by Vpr.

Experimental Procedures

Cloning and Plasmid Constructions—HIV-1 NL4-3 Vpr (Vpr^{HIV-1 NL4-3}) was cloned into the pcDNA3.1 vector (Life Technologies, Inc.) with an HA tag. Vpr^{HIV-1 NL4-3} with R80A and Q65R mutants were prepared using the QuikChange mutagenesis kit (Agilent Technologies). Various Vpr clones from HIV-1 group (O, M, and N), SIVcpz PTT and PTS in pCG

vector, were gifts from Dr. Jacek Skowronski at Case Western Reserve University. Vpr^{HIV-1 NL4-3} was cloned into pET41 vector (EMD Bioscience). FLAG-tagged SLX4 (residues 1430– 1834), FLAG-tagged MUS81, and Myc-tagged EME1 or EME2 were cloned into pcDNA3.1 vector. HA-tagged DCAF1, FLAGtagged DCAF1, and V5-tagged DDB1 were cloned into pcDNA3.1 vector. For *Escherichia coli* expression, MUS81 and EME1 were cloned into pET28 and pET21 vectors, respectively. All other clones were described previously (66).

Protein Purification and GST Pulldown—DCAF1 and GST fusion DDB1 were expressed in insect cell Sf21 (Life Technologies, Inc.). GST, GST-His₆-Vpr, Vpr, and MUS81-EME1 were expressed in *E. coli* Rosetta 2 (DE3) or BL21, and proteins were purified as described previously (48, 66). Purified GST, GST-His₆-Vpr, GST-DDB1/DCAF1, or GST-DDB1/DCAF1/Vpr were incubated with GST beads for 3 h with shaking in PBS buffer with 0.5% Nonidet P-40 (binding buffer). Excess unbound protein was washed by binding buffer. Purified MUS81-EME1 was incubated with these GST beads for 3 h with shaking in binding buffer. After washing three times, binding proteins were eluted by 20 mM glutathione in binding buffer.

Mammalian Cell Lines, Transfection, and Western Blotting-Human embryonic kidney cell lines (HEK293T from ATCC) were maintained in advanced DMEM, supplemented with 1% (v/v) 100× glutamine, and 10% (v/v) fetal bovine serum at 37 °C, 5% CO $_2$. Typically, 3 \times 10 6 cells were plated on a 100-mm dish a day before transfection. Transfection was performed with a mixture of pCDNA plasmids encoding specific cDNAs using Lipofectamine 3000 (Life Technologies, Inc.), according to the manufacturer's protocol. Empty pcDNA vector plasmid was used as a balance. Cells were harvested 48 h post-transfection. After washing with PBS, cells were lysed with sonication in lysis buffer (50 mM Tris, 150 mM NaCl, 0.5% Nonidet P-40, pH 7.5) with Complete protease inhibitor mixture (Sigma). For protein analysis by Western blotting, some cell lysate was mixed with 1:5 (v/v) of $5 \times$ loading buffer. After centrifugation at 10,000 \times g for 10 min, 20 μ l of anti-FLAG M2 magnetic beads (Sigma) were added to the cell lysate supernatant for co-immunoprecipitation assay. The beads/cell lysate mixture were incubated with shaking at 4 °C for 5 h followed by washing with lysis buffer three times. Bound proteins were eluted with FLAG peptides at a concentration of 0.1 mg/ml and subjected to SDS-PAGE and Western blotting analysis with appropriate antibodies. The antibodies used in this study included anti-HA (Covance), anti-Myc (Covance), anti-FLAG (Abnova), anti-V5 (Sigma), anti-actin (Sigma), anti-DCAF1 (Santa Cruz Biotech-



FIGURE 5. **Vpr mutants that lack G₂ arrest activity down-regulate MUS81-EME1.** *A*, MUS81 and EME1 plasmids were co-transfected with WT Vpr or a C-terminally truncated Vpr mutant 1–79 plasmid, and empty vector plasmid was added to balance total plasmid in HEK293T cells. After 48 h of transfection, cells were harvested, and lysates were analyzed by immunoblotting. Intensity of bands corresponding to MUS81 and EME1 was quantified from four independent experiments. *B*, DDB1, DCAF1, MUS81, EME1, and Vpr(1–79)-encoding plasmids, along with empty vector plasmids, were transfected as indicated. FLAG-tagged MUS81 proteins were immunoprecipitated (*IP*) with anti-FLAG antibody from cell lysates and analyzed with immunoblotting. *C*, MUS81 and EME1 plasmids were co-transfected with Vpr R80A or Vpr WT plasmids. *D*, combinations of MUS81, EME1, DDB1, DCAF1, and Vpr R80A were transfected as indicated. Proteins associated with MUS81 were analyzed as described above. *E*, MUS81 and EME1 plasmids were co-transfected with Vpr Q65R plasmid. *F*, Vpr Q65R plasmids were transiently expressed in cells along with combinations of MUS81, EME1, DDB1, and DCAF1 plasmids. BUS81-associated proteins were analyzed as describe above. *G*, Vpr WT or Q65R mutant plasmids were co-transfected with DDB1 and DCAF1 plasmids. FLAG-tagged DCAF1 proteins were analyzed as describe above. *G*, Vpr WT or Q65R mutant plasmids were co-transfected with DDB1 and DCAF1 plasmids. FLAG-tagged DCAF1 proteins were analyzed as describe above. *G*, Vpr WT or Q65R mutant plasmids were co-transfected with DDB1 analyzed by Western blotting. *H* and *J*, cell cycle analysis of HEX293T cells transiently transfected with empty vector plasmid (*H*), Vpr WT (*I*), Vpr(1–79) (*J*), Vpr R80A (*K*), or Vpr Q65R (*L*) plasmid. The ratio of cells in G₂/M relative to those in G₁ is indicated.





FIGURE 7. **Two possible modes of Vpr-induced MUS81-EME1 down-regulation**. *A*, SLX4-SLX1 mediates CRL4-DCAF1-Vpr and MUS81-EME1 interaction and promotes MUS81-EME1 down-regulation. *B*, MUS81-EME1 is a natural partner of CRL4-DCAF1, and Vpr promotes MUS81-EME1 poly-ubiquitination and down-regulation. An additional cellular factor(s) may bridge the interaction between MUS81-EME1 and CRL4-DCAF1 when Vpr is not present.

nology), anti-MUS81 (Santa Cruz Biotechnology), anti-EME1 (Santa Cruz Biotechnology), anti-EME2 (Santa Cruz Biotechnology), anti-SLX4 (LSBio), anti-GST (Santa Cruz Biotechnology), anti-Vpr (NIH AIDS Reagent), anti-goat IgG (Santa Cruz Biotechnology), anti-rabbit IgG (Sigma), and anti-mouse IgG (Sigma).

Cycloheximide Chase Assay—Cells were transiently transfected with a mixture of pCDNA plasmids encoding MUS81 and empty vector or Vpr using Lipofectamine 3000. After 40 h post-transfection, cells were incubated with 20 μ g/ml cycloheximide (Sigma) to block *de novo* protein synthesis. Cells were harvested after 1, 2, 4, 6, and 8 h of cycloheximide treatment (0 h for no cycloheximide). Cell lysates were separated by SDS-PAGE and analyzed by Western blotting with appropriate antibodies. Intensity of bands was quantified by ImageJ (67).

siRNA-mediated Protein Knockdown—siRNA targeting DCAF1 (5'-CCACAGAAUUUGUUGCGCA-3', catalogue no. J-021119-10) and its non-targeting control siRNA (5'-UGGUUUACAUGUCGACUAA-3', catalogue no. D-001810-01), On-TARGETplus human SLX4 siRNA SMARTpool (5'-UCAAA-CGGCACUCAGAUAA-3', 5'-GCGGAGACUUUGUU-

GAAAU-3', 5'-CAAGUGAGCCCGAGGAACA-3', and 5'-UCAGAGCCGUCCCAAAUAA-3', catalogue no. L-014895-00), and its Control On-TARGETplus non-targeting siRNA (5'-UAAGGCUAUGAAGAGAGAUAC-3', 5'-AUGUAUUGGCCU-GUAUUAG-3', 5'-AUGAACGUGAAUUGCUCAA-3', and 5'-UGGUUUACAUGUCGACUAA-3', catalogue no. D-001206-14) were obtained from Dharmacon (GE Healthcare). HEK293T cells were transfected with siRNAs using Lipofectamine RNAiMAX according to the manufacturer's instructions. 24 h later, cells were transfected with pcDNA plasmids encoding specific cDNAs using Lipofectamine 3000 (Life Technologies, Inc.), according to the manufacturer's protocol. Total cell lysates were subjected to SDS-PAGE, and protein levels were analyzed by Western blotting with antibodies after 48 h.

Cell Cycle Assays—HEK293T cells seeded on 12-well plates were transfected with Vpr expressing pCDNA plasmids or empty vector. 48 h after transfection, the cells were treated with trypsin and washed twice with PBS. After fixation with 70% ethanol for 30 min, cells were washed twice with PBS and suspended in 1 ml of propidium iodide staining solution (0.1%



FIGURE 6. Vpr-dependent down-regulation of MUS81-EME1 is independent of SLX4. *A*, HEK293T cells were transfected with MUS81 and EME1 plasmids, with increasing amounts of SLX4 plasmid along with Vpr or empty vector plasmid as indicated. Cells were harvested, and proteins levels were analyzed by immunoblotting with appropriate antibodies. *B*, SLX4 siRNA or control siRNA was transfected 24 h prior co-transfection with the indicated combinations of Vpr, MUS81, EME1, or balanced empty plasmid. *C*, purified MUS81·EME1, GST, GST-His₆-Vpr, GST-DDB1·DCAF1, and GST-DDB1·DCAF1·Vpr complexes were separated by SDS-PAGE and stained with Coomassie Blue. *D*, GST, GST-His₆-Vpr, GST-DDB1·DCAF1, or GST-DDB1·DCAF1·Vpr complex was incubated with recombinant MUS81-EME1. Bound fractions were analyzed by Western blotting. All experiments were repeated three times with similar results.

Triton X-100, 10 μ g/ml propidium iodide, and 100 μ g/ml DNase-free RNase A in PBS). After 60 min of staining at room temperature, cells were sorted by flow cytometry (BD LSRII SORP). Data were analyzed using FlowJo software (Tree Star Inc., Ashland, OR).

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