

Modulation of Cellular Protein Trafficking by Human Immunodeficiency Virus Type 1 Nef: Role of the Acidic Residue in the ExxxLL Motif

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The *nef* gene contributes to the replication of primate lentiviruses by altering the trafficking of cellular proteins involved in adaptive immunity (class I and II major histocompatibility complex [MHC]) and viral transmission (CD4 and DC-SIGN). A conserved acidic leucine-based sequence (E₁₆₀xxxLL) within human immunodeficiency virus type 1 (HIV-1) Nef binds to the cellular adaptor protein (AP) complexes, which mediate protein sorting into endosomal vesicles. The leucine residues in this motif are required for the down-regulation of CD4 and for the up-regulation of DC-SIGN and the invariant chain of MHC class II, but the role of the acidic residue is unclear. Here, substitution of E160 with uncharged residues impaired the ability of Nef to up-regulate the expression of the invariant chain and DC-SIGN at the cell surface, whereas substitution with a basic residue was required for a similar effect on the down-regulation of CD4. All substitutions of E160 relieved the Nef-mediated block to transferrin uptake. E160 was required for the efficient interaction of Nef with AP-1 and AP-3 and for the stabilization of these complexes on endosomal membranes in living cells. Systematic mutation of the ExxxLL sequence together with correlation of binding and functional data leads to the hypotheses that AP-1 and AP-3 are major cofactors for the effect of Nef on the trafficking of transferrin, are less important but contribute to the modulation of the invariant chain and DC-SIGN, and are least critical for the modulation of CD4. The data suggest that the E160 residue plays a differential role in the modulation of leucine-dependent Nef-targets and support a model in which distinct AP complexes are used by Nef to modulate different cellular proteins.

The *nef* gene of primate lentiviruses is required for high-level viremia and the efficient pathogenesis of AIDS (12, 25, 44). These effects are at least partly due to the effect of Nef on the cellular protein trafficking environment. Nef alters the subcellular localization of a number of proteins, including CD4, DC-SIGN, transferrin receptor, tumor necrosis factor, LIGHT, CD28, class I major histocompatibility complex (MHC), and both mature and immature class II MHC (1, 27, 39, 40, 42, 43). These effects likely influence the efficiency of viral replication. For example, the down-regulation of the cell surface level of CD4 by Nef prevents the binding of the viral envelope glycoprotein (gp120) to CD4 on the surface of the virus-producing cell, preserving the infectivity of newly formed virions and potentially enhancing their release (26, 37). In contrast to CD4, Nef up-regulates the surface level of DC-SIGN, a C-type lectin expressed on dendritic cells that both allows the uptake of mannoseylated antigens and serves as an adhesion molecule, facilitating the interaction of dendritic cells with T cells during antigen presentation (17, 40). Although DC-SIGN binds gp120, the human immunodeficiency virus virions internalized into dendritic cells remain infectious and are subsequently transmitted to T cells, a process that Nef may facilitate. Finally, Nef disrupts the presentation of viral antigens by down-regulating class I MHC and mature class II MHC from the cell surface, while up-regulating the surface expression of the invariant chain, which normally chaperones the immature class II

complex to an endosomal compartment in which antigens derived from the extracellular space are processed (42).

The down-regulation of CD4, CD28, and transferrin receptor as well as the up-regulation of tumor necrosis factor, LIGHT, invariant chain, and DC-SIGN require two leucine residues within a C-terminal, solvent-exposed loop of the Nef protein (4, 9, 18, 27, 40, 42, 43). The leucine codons are conserved among human immunodeficiency virus type 1 (HIV-1) *nef* alleles, and their mutation leads to the complete loss of Nef's effect on these membrane proteins. The two leucines, together with the adjacent upstream sequence, conform to a class of intracellular sorting signals whose consensus sequence is E/DxxxLL. These sequences bind to the heterotetrameric adaptor protein (AP) complexes, which form part of the coat of a subset of vesicles that mediate transport within the endosomal system (21, 31). The family of AP complexes has four members: AP-1 mediates vesicular transport between the *trans*-Golgi network and endosomal compartments; AP-2 mediates transport between the plasma membrane and early/sorting endosomes; AP-3 mediates transport from the *trans*-Golgi network and early endosomes to late endosomes and lysosomes; and AP-4 may mediate transport from the *trans*-Golgi network to the basolateral plasma membrane of polarized cells.

The roles of these individual AP complexes in the effects of Nef on protein trafficking are unclear. The rate of internalization of CD4 from the cell surface is increased by Nef, suggesting involvement of AP-2 (1). In further support of a role for AP-2, Nef has been detected in clathrin-coated pits at the plasma membrane by electron microscopy (16), and a domi-

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