Murine Endogenous Retrovirus MuERV-L Is the Progenitor of the “Orphan” Epsilon Viruslike Particles of the Early Mouse Embryo

David Ribet, Sophie Louvet-Valée, Francis Harper, Nathalie de Parseval, Marie Dewannieux, Odile Heidmann, Gérard Pierron, Bernard Maro, and Thierry Heidmann

Unité des Rétrovirus Endogènes et Éléments Rétroïdes des Eucaryotes Supérieurs, CNRS UMR 8122, Institut Gustave Roussy, Villejuif F-94805, and Université Paris-Sud, Orsay F-91405, France; Laboratoire de Biologie Cellulaire du Développement, CNRS UMR 7622, Université Pierre et Marie Curie, Paris F-75252, France; and Laboratoire de Réplication de l’ADN et Ultrastructure du Noyau, UPR 1983 Institut André Lwoff, Villejuif F-94801, France

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Viruslike particles which displayed a peculiar wheellike appearance that distinguished them from A-, B- or C-type particles had previously been described in the early mouse embryo. The maximum expression of these so-called epsilon particles was observed in two-cell-stage embryos, followed by their rapid decline at later stages of development and no particles detected at the zygote one-cell stage. Here, we show that these particles are in fact produced by a newly discovered murine endogenous retrovirus (ERV) belonging to the widespread family of mammalian ERV-L elements and named MuERV-L. Using antibodies that we raised against the Gag protein of these elements, Western blot analysis and in toto immunofluorescence studies of the embryos at various stages disclosed the same developmental expression profile as that observed for epsilon particles. Using expression vectors for cloned, full-length, entirely coding MuERV-L copies and cell transfection, direct identification of the epsilon particles was finally achieved by high-resolution electron microscopy.

Multiple viruslike particles are generated by mouse tissues under normal and/or pathological conditions. These have been initially characterized according to morphological criteria by electron microscopic examination and classified as type A, B, C, and epsilon particles depending on their site of assembly (i.e., within the cell cytoplasm or at the cell surface) and the structure of the mature particles (3, 24; reviewed in reference 16). These particles were suspected to be generated by so-called endogenous retroviruses (ERVs), as they can be observed in normal animals in the absence of any exogenous infection. Some of these particles were even observed in the developing embryo and, for some of them, at very early stages with, for instance, a peak of occurrence of particles of the epsilon type at the two-cell stage (23, 24). In most cases, the ERVs responsible for the formation of these particles have been identified, thanks to the complete sequencing of the mouse genome (15a), the in silico identification of elements with full coding capacity, and, finally, the cloning of functional copies that could be demonstrated to be responsible for the synthesis of the corresponding particles in appropriate ex vivo assays by electron microscopic analysis. Examples include the highly reiterated intracisternal A-type particles (IAPs) (7; reviewed in reference 12), the Mus musculus type D (MusD) intracytoplasmic elements (18, 19), and the endogenous murine leukemia virus and mouse mammary tumor virus (reviewed in reference 4). Yet, the epsilon particles, first identified more than 25 years ago in mouse embryos (23, 24) and later in some iododeoxyuridine-induced mouse cell lines (13, 14), have up to now remained “orphan.” Actually, these particles have a rather unusual morphology, with a wheel-shaped core disclosing an electron-lucent space surrounded by two concentric electron-dense rings that are separated by a radial array of internal spikes, not found in any present-day infectious retrovirus and at variance with the other types of particles for which similar morphologies can be found among infectious elements of animals. Along this line, we had previously identified a new family of ERVs (ERV-L) in humans (5) that we further demonstrated to be present in all placental mammals and, therefore, to have settled within the genome of a mammalian ancestor more than 70 million years ago (1, 2; see also reference 8). A phylogenetic analysis of their pol genes disclosed a rather unusual position among retroviral elements, with, again, no clear-cut relationship to the major groups of retroviruses (Fig. 1B). Despite its age, this family has most probably maintained some of its elements in an active state, since we could demonstrate relatively recent amplifications of members of this family of elements in the mouse, not observed in the rat, having occurred less than 10 million years ago (1, 2; see also references 6 and 10). Interestingly, in the mouse, most of these MuERV-L elements are full length and have complete open reading frames (ORFs) with identifiable gag and pol genes (but no env) (Fig. 1A and C), and transcripts from these elements have recently been detected in embryos at early developmental stages (9, 11, 20, 21). Altogether, it therefore seems plausible that the epsilon particles could be encoded by members of the MuERV-L family of elements, and experiments were therefore devised to test this hypothesis.
As a first step, we analyzed the developing embryos for MuERV-L protein expression at the very early stages where epsilon particle formation had been observed. To do so, we produced a recombinant protein, corresponding to the complete Gag polyprotein of a cloned MuERV-L element (GenBank accession no. Y12713), that we used to immunize rabbits and raise specific antibodies. Western blot analyses of cell extracts from Swiss (OF1) mouse embryos using this antisera (Fig. 2A, left panel) revealed a 66-kDa band most probably corresponding to the complete MuERV-L Gag polyprotein (theoretical molecular mass, 64 kDa), observed as well in the Western blot shown in Fig. 2A, right panel, with cells transfected with an expression vector for MuERV-L (see below). As illustrated in Fig. 2A, this MuERV-L Gag product shows a peak of expression in embryos at the two-cell stage (lane 2c, collected 36 h postfertilization), a much-lower expression level at the eight-cell stage (lane 8c, collected 54 h postfertilization), and no expression in immature oocytes (i.e., in the germinal vesicle) (lane GV) or blastocysts (lane Bl), as previously reported for the epsilon particles. Moreover, immunofluorescence analysis of the embryos with the anti-Gag antisera (Fig. 2B) revealed an intense intracytoplasmic staining at the two-cell stage (Fig. 2B, panel 2c), possibly at the level of...

FIG. 1. Structure and phylogeny of MuERV-L elements. (A) Genomic organization of an MuERV-L provirus, with the LTRs (in dark gray) flanking two ORFs homologous to the retroviral gag and pol genes (in light gray). The pol gene further discloses domains homologous to retroviral pro and dUTPase genes. A primer binding site (PBS) complementary to tRNALeu and a polypurine track (PPT) can be identified. (B) Phylogeny of retroviruses, based on their reverse transcriptase domain. The tree was constructed by the neighbor-joining method using the seven blocks of conserved residues found in the reverse transcriptase domain of all retroelements and was rooted with non-LTR retrotransposons. All sequences are readily accessible from GenBank and previous reports (e.g., reference 15). Percent bootstrap values obtained from 1,000 replicates are indicated. The retroviruses “endogenized” in the mouse genome are marked (*). The seven retroviral genera are indicated on the right. PFV, primate foamy virus; MuRRS, murine retrovirus-related DNA sequence; MuRV-Y, murine retrovirus Y associated; MmERV, Mus musculus ERV; GLN, murine leukemia virus; WDSV, walleye dermal sarcoma virus; ALV, avian leukemia virus; MMTV, mouse mammary tumor virus; JSRV, jaagsiekte sheep retrovirus; MusD, Mus musculus type D retrovirus; MPMV, Mason-Pfizer monkey virus; IAP, IAP with an envelope gene; HIV-1, human immunodeficiency virus type 1; HTLV-1, human T-cell leukemia virus type 1. (C) Search for full-length, entirely coding MuERV-L copies in the C57BL/6J genome. Blast search was carried out with the NCBI m36 mouse assembly (April 2006 release) and yielded 489 full-length MuERV-L copies among which 350 contain the two complete gag and pol ORFs.
the endoplasmic reticulum (ER), consistent again with the electron microscopic observations made in the past. This staining markedly decreases at the eight-cell stage (panel 8c), is not observed in cells transfected with a control plasmid (Fig. 3A). MuERV-L-transfected cells were also analyzed by electron microscopy (Fig. 3B, panels 1 and 2). Forty-eight hours posttransfection, the cells were fixed in 1.6% glutaraldehyde, postfixed in 1% osmium tetroxide, and embedded in Epon 812. Ultrathin sections were stained with uranyl acetate and lead citrate. The cells disclosed intracisternal particles (not observed in cells transfected with a control plasmid) with the canonical epsilon morphology: a central electron-lucent space surrounded by two concentric electron-dense rings separated by a radial array of internal spikes. Budding particles could also be observed at the level of the ER membrane (Fig. 3B, panel 2). Clearly, the structure and site of expression of the observed particles are similar to those of the naturally occurring epsilon particles found in the embryo (Fig. 3B, panel 3) (23). Of note, expression vectors for two other full-length, coding-competent MuERV-L copies (possibly corresponding to distinct MuERV-L burst events within the Mus species; see references 2 and 6) produced particles with an identical morphology.

In conclusion, the present study unambiguously demonstrates that the epsilon particles are indeed generated by the MuERV-L endogenous elements. This family of ERVs is therefore the most ancient “parasite” of mammals identified to date still able to generate viruslike particles. Their morphology—not recovered in any present-day infectious retrovirus—is rather unique, and a remaining question to be answered now concerns the possible role of these particles. In particular, the intracisternal localization of the generated epsilon particles is similar to that of the still-functional IAPs, which are also actively transcribed at stages of mouse embryonic development, albeit with a peak of expression occurring later, at the 8- to 16-cell stages (24), and which transcribe at a high rate. Experiments should now be devised to determine whether some present-day MuERV-L elements are still competent for autonomous retrotransposition or whether, as suggested by a recent report, they are mainly used by their host as “early” promoters dispersed throughout the genome (17).

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