

Similarity between p15E of murine and feline leukaemia viruses and p21 of HTLV

SIR — Retroviral infections of a number of animal species are frequently associated with immunosuppression. The hydrophobic retroviral transmembrane protein p15E is highly conserved among type C and type D retroviruses¹. This protein may contribute to the immunosuppression associated with retroviral infections since it inhibits *in vitro* lymphocyte transformation in man², cats³ and mice⁴ and increases tumour incidence in cats given feline sarcoma virus⁵. The mechanism for the inhibition of human lymphocyte transformation by p15E was shown to be due to suppression of interleukin-2 (IL-2) production⁶. We have demonstrated *in vitro* suppression of human monocyte function⁶ and *in vivo* inhibition of murine macrophage accumulation to inflammatory foci by p15E from murine leukaemia virus (MuLV)⁷ or by tumour-associated proteins which are antigenically related to p15E^{6,8}.

Human T-cell leukaemia-lymphoma virus (HTLV) is the designation for a family of T-lymphotropic, exogenous type C retroviruses which have been isolated in a number of countries⁹⁻¹³. These viruses have been linked as aetiological agents to certain types of leukaemias and lymphomas and to the acquired immunodeficiency syndrome (AIDS)¹⁴⁻¹⁶. It has recently been reported that the polymerase gene products of HTLV, MuLV and avian Rous sarcoma virus are partially homologous in their amino acid sequence¹⁷. Since immunosuppression often accompanies the HTLV-associated T-cell malignancies and is characteristic of AIDS, we hypothesized that HTLV-viruses might have, in addition to their lymphocytotoxic effects, other features in common with immunosuppressive MuLV and feline leukaemia viruses (FeLV). We therefore compared the amino acid sequences for p15E of Friend¹⁸, Moloney¹⁹, AKV²⁰, and Gross²¹ murine leukaemia viruses (FLV, MoLV, AKV, GLV), FeLV²², and mink-cell focus-forming viruses of Moloney²³ and AKR²⁴ origin (MMCF, AMCF) with the sequences of the envelope proteins of HTLV-I²⁵ and HTLV-II²⁶. The sequences were analysed for homology using the PROTHOM computer program developed by Fristensky *et al.*²⁷.

As shown in the figure above there is a significant degree of homology (73%) between the various p15E proteins and HTLV-1 and HTLV-II which occurs in a 26-amino acid sequence located in the p21 region of HTLV. It is particularly noteworthy that the first 10 amino acids in this region are identical and that this region of homology occurs in almost the same region of the p15E and p21 molecules. The homology begins at residue 70 in p15E and at residue 377 of the HTLV-I envelope protein and at residue 373 of the HTLV-II envelope protein which both correspond to

HTLV _I ENV	Q N R R G L D L L F	W E Q	G G L C K	A L Q E	Q C R F	402
HTLV _{II} ENV	Q N R R G L D L L F	W E Q	G G L C K	A I Q E	Q C C F	398
FLV p15E	Q N R R G L D L L F	L K E	G G L C A	A L K E	E C C F	95
MLV p15E	Q N R R G L D L L F	L K E	G G L C A	A L K E	E C C F	
AKV p15E	Q N R R G L D L L F	L K E	G G L C A	A L K E	E C C F	
GLV p15E	Q N R R G L D L L F	L K E	G G L C A	A L K E	E C C F	
MMCF p15E	Q N R R G L D L L F	L K E	G G L C A	A L K E	E C C F	
AMCF p15E	Q N R R G L D L L F	L K E	G G L C A	A L K E	E C C F	
FeLV p15E	Q N R R G L D L L F	L K E	G G L C A	A L K E	E C C F	

Amino acid sequence homology between p21 of HTLV-I and HTLV-II and p15E of murine and feline leukaemia viruses. Residue 377 of the RTLV-I envelope and 373 of the HTLV-II envelope both correspond to residue 65 of the p21 protein.

residue 65 of the p21 transmembrane protein²⁶.

The significance of such a well conserved region occurring in murine, feline, and human retroviruses is not yet clear. However, since dramatic immunosuppression is often associated with these viruses and purified p15E is immunosuppressive in a variety of systems, it will be particularly important to determine whether p21 of HTLV can also inhibit immune functions.

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Islet activating protein analogous to p21 ras?

SIR — In a recent article on mutant *ras* protein and cell transformation, Rob Newbold considered the significance of new findings about p21 *ras* function and activation. In particular, similarities between p21 *ras* and the guanine nucleotide binding proteins which are responsible for stimulation and inhibition of adenylate cyclase were emphasized and an analogy drawn with the action of cholera toxin on G_s protein GTPase, implying that p21 *ras* might act in an analogous manner to cholera toxin, which exerts a strong proliferative stimulus on certain mammalian cells in culture.

I suggest that the analogy with cholera toxin is dubious in view of the fact that rat cells transformed by Kirsten sarcoma virus have been shown to contain less cyclic AMP and lowered adenylate cyclase activity. Therefore a closer analogy may be with the action of islet activating protein which reacts with Gi proteins to inhibit adenylate cyclase and reduce cyclic AMP. This would imply that decreased cyclic AMP assists proliferation.

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Heritable IQ — a reason to bother

SIR — The recent article from Teasdale and Owen¹ prompts one to wonder about IQ genes. If intelligence is substantially heritable, is it determined by genes in a positive sense, such that better allelic combinations of IQ genes produce more intelligent people? This may seem to be the most parsimonious explanation, but parsimony is in the eye of the beholder.

That simple genetic mechanisms can account for conspicuously low IQ stirs no