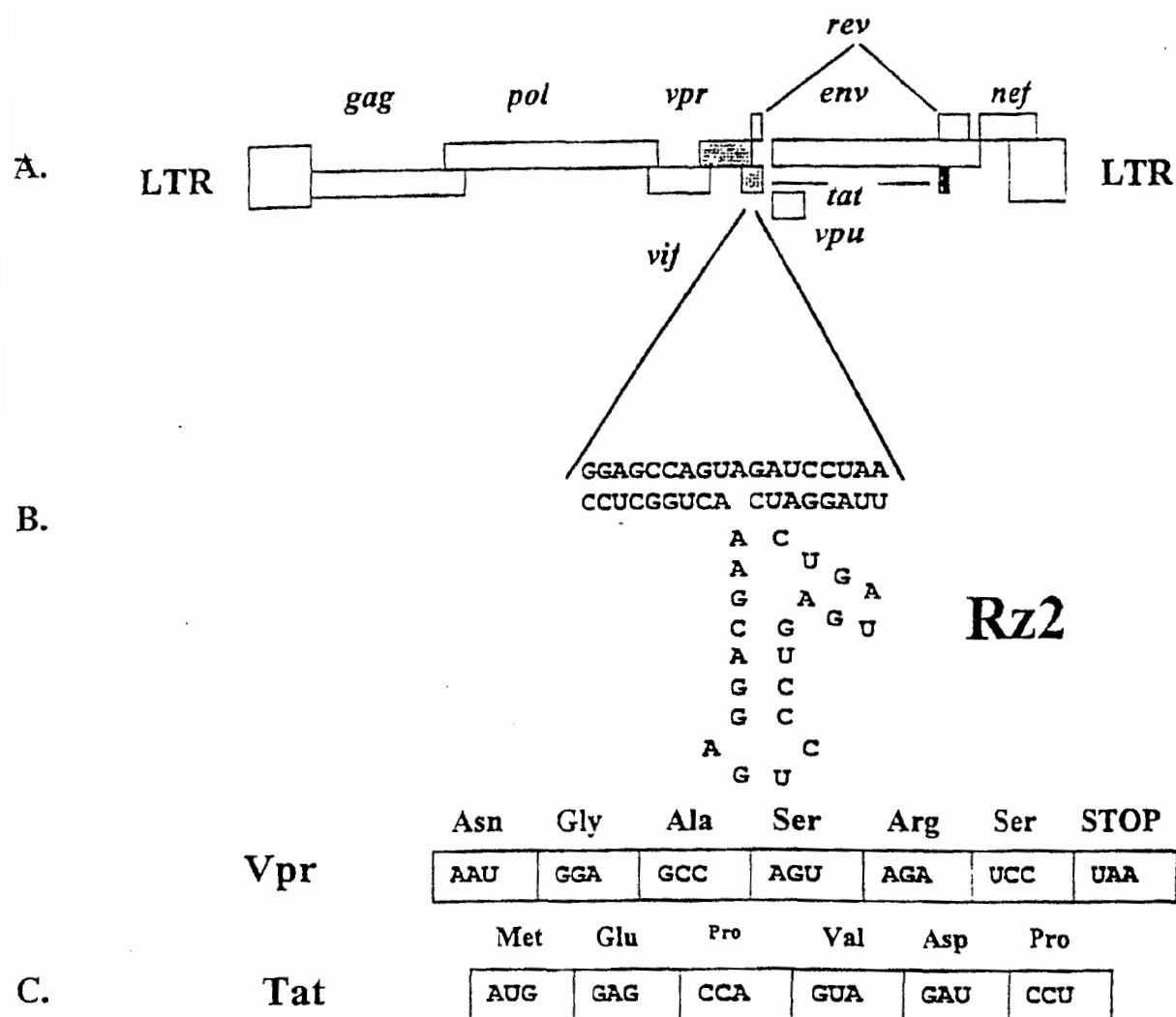




CD34+ Cell Product Processing

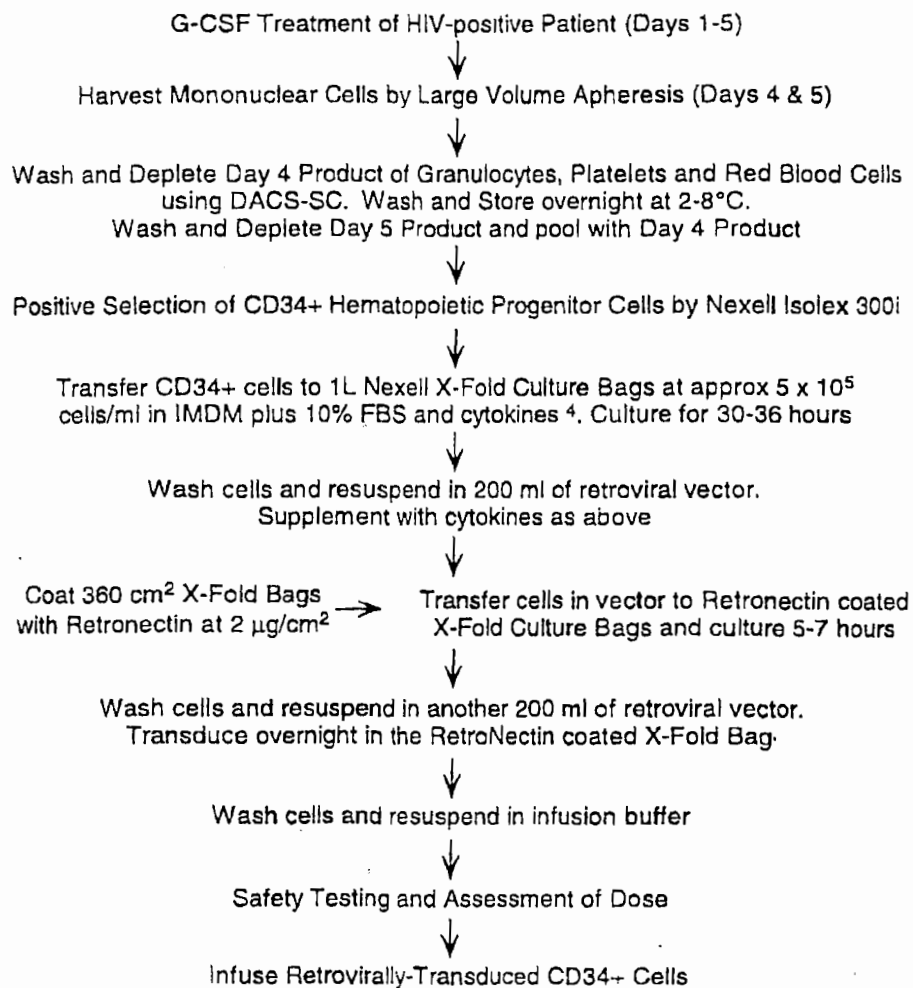
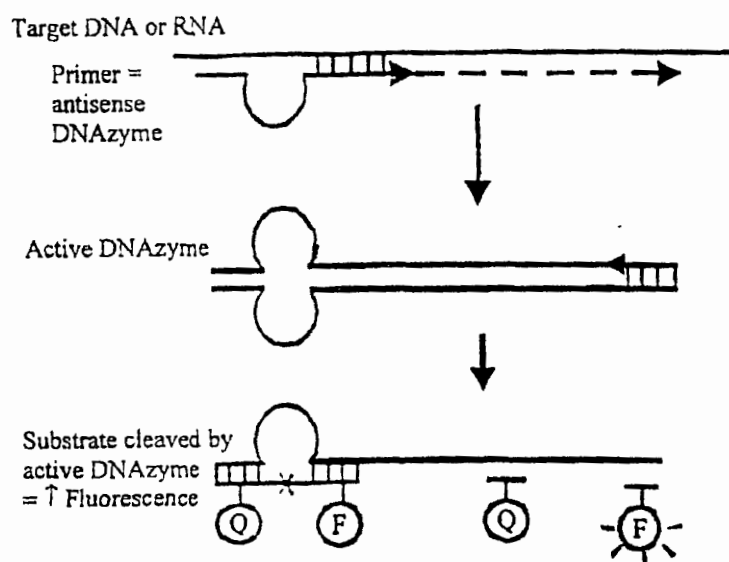


Figure 2

A



B

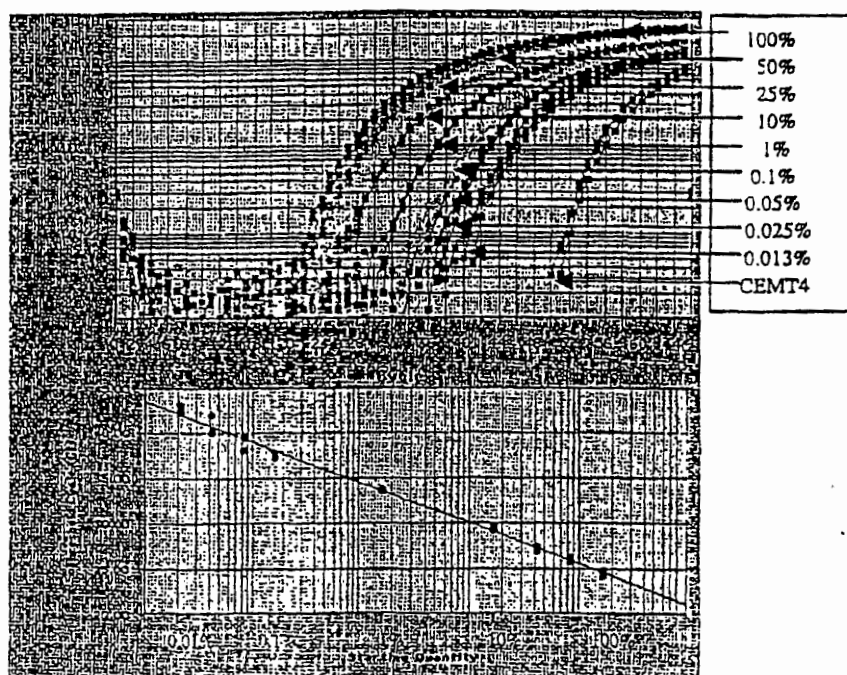


Figure 3.

(A) Schematic representation of the DzyNA PCR detection method
 (B) Quantitation of *neo^R* gene by DzyNA PCR generation of a standard curve which allows for estimate of percentage marking in an 'unknown' sample.

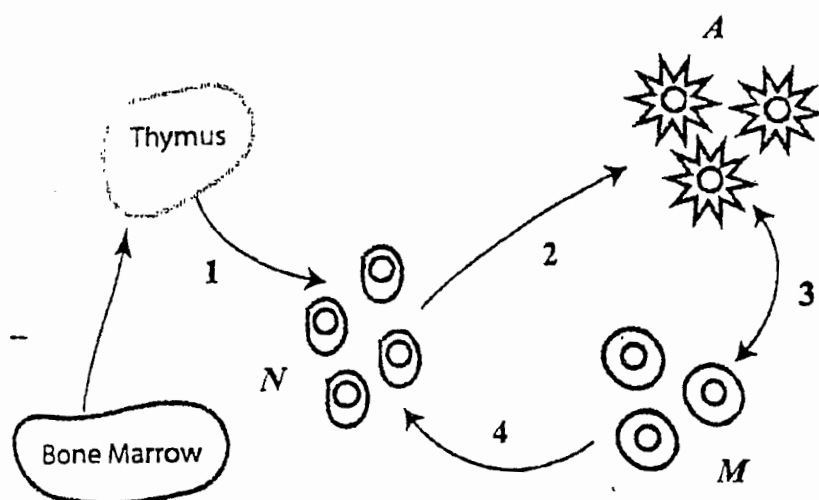


Figure 4: Diagram of the CD4+ T lymphocyte model: naive cells N, activated cells A, memory cells M.

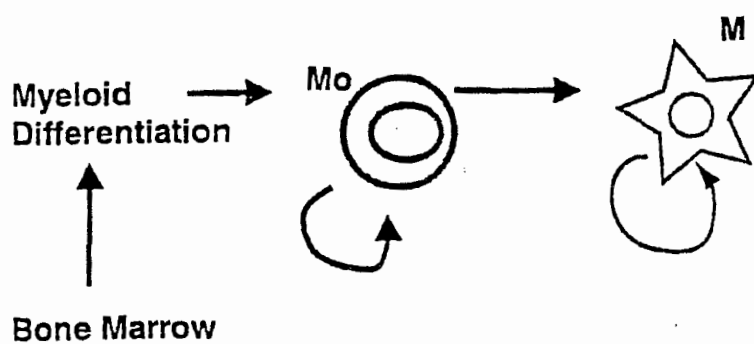


Figure 5: Diagram of the macrophage model: Monocyte Mo, Macrophage M.

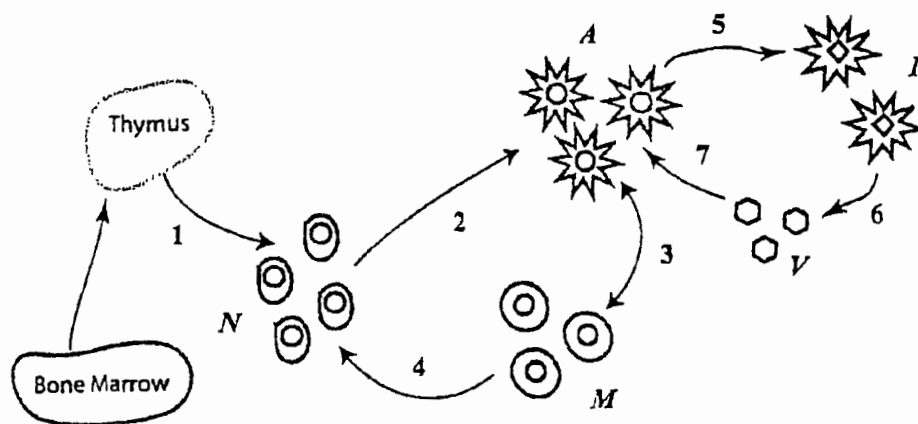


Figure 6: Diagram of the CD4+ T lymphocyte model: naive cells N, activated cells A, memory cells M, free HIV virus V.

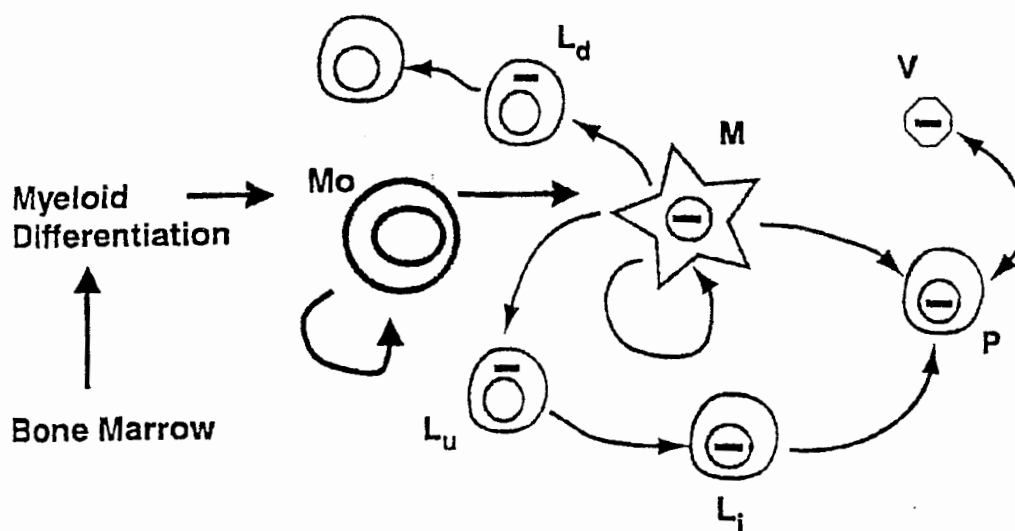


Figure 7: Diagram of the Macrophage Model: free virus V, productively infected cells P, infected macrophages M, resting CD4+ T cells with defective unintegrated HIV DNA L_d, resting CD4+ T cells with competent unintegrated HIV DNA L_u, resting CD4+ T cells with integrated HIV DNA.