

can promote this right to a *chance of cure*. This statement was approved in principle by the following representatives of these study groups in San Diego in December 2003:

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Chronic myeloid leukemia-specific T-cell responses

In chronic myeloid leukemia (CML), the chimeric p210 fusion protein resulting from the bcr-abl fusion gene produced by the t(9;22)(q34; q11) translocation, in virtue of the unique sequence of amino acids contained in the junctional regions, which is CML-specific,

furnished the rationale for a peptide vaccine strategy in this disease.¹ Although treatment of CML has been revolutionized by imatinib mesylate (see previous articles in this journal)²⁻¹² a portion of patients are or become resistant to this drug,¹³ so that alternative treatments have been proposed.¹⁴⁻¹⁷ Theoretically, CML-specific T cells might be useful for therapeutic purposes. As discussed by Posthuma and co-workers in this issue,¹⁸ a prerequisite for a CML-specific T-cell response is proteasomal degradation of intracellular BCR/ABL protein resulting in presentation of BCR/ABL-specific oligopeptides by HLA class I or HLA class II molecules on the membrane of leukemic cells and the presence of T-cells with a T-cell receptor (TCR) that can recognize these peptide-HLA complexes. Until now there is no definite proof of the existence of such CML-specific responses *in vivo*. In their work, Posthuma and co-workers¹⁸ show that proteasomal degradation of BCR/ABL protein can generate a CML-specific HLA-A*0301 restricted peptide, but high-avidity T-cells recognizing this BCR/ABL-specific antigen could not be demonstrated.

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Molecular basis of von Willebrand disease and its clinical implications

Von Willebrand's disease (VWD) is an autosomally inherited bleeding disorder caused by a deficiency or abnormality of von Willebrand factor (VWF). Castaman and co-workers¹ have recently reviewed the molecular basis of VWD in this journal. VWD has a prevalence of about 1% in the general population, but the figure for clinically relevant cases is lower (about 100/million inhabitants). Most cases appear to have a partial quantitative deficiency of VWF (type 1 VWD) with variable bleeding tendency, whereas qualitative variants (type 2 VWD), due to a dysfunctional VWF,

are clinically more homogeneous. Type 3 VWD is rare and the patients have a moderate to severe bleeding diathesis because of the virtual absence of VWF, and a recessive pattern of inheritance.

In this issue, Hilbert and co-worker² report studies on a new mutation in the VWF gene that causes type 2A VWD. These patients may be poorly responsive to desmopressin and should receive FVIII/VWF concentrates in cases of prolonged mucosal bleeding and major surgery. This study provides an example of the relevance of molecular studies for clinical practice.

Other studies on VWF or VWD recently published in this journal are listed below.³⁻⁸

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