

Conclusions: Our results indicate that BAFF-induced inflammation-related signal transduction can lead to BCL10 nuclear translocation and NF- κ B activation. The autocrine BAFF signal transduction pathway may contribute to the HP-independence of gastric MALT lymphoma without t(11;18)(q21;q21).

9233

POSTER

Pharmacokinetic Intra-individual Variability of Imatinib – Consequences for Therapeutic Drug Monitoring in Chronic Myeloid Leukaemia

C. Bardin¹, X. Declèves¹, A. Vekhoff², L. Labat¹, F. Chast¹. ¹Hôtel-Dieu Hospital, Pharmacy-Pharmacology-Toxicology, Paris Cedex 04, ²St-Antoine Hospital, Clinical Hematology, Paris, France

Background: Imatinib is a competitive inhibitor of protein tyrosine kinase Bcr-Abl and is currently used for the treatment of chronic myeloid leukaemia (CML) and other digestive malignant pathologies. Trough imatinib plasma levels are associated with major molecular response in CML. Due to important interpatient variability, monitoring of imatinib plasma levels can be very useful especially in the case of treatment failure. The aim of this work was to determine intra-individual variability in imatinib plasma concentrations to a better knowledge of therapeutic drug monitoring (TDM) of imatinib in the real life.

Methods: Imatinib plasma concentrations were determined by high-performance liquid chromatography with UV detection at 262nm after liquid-solid extraction. Limit of quantification was set at 200 ng/mL. Blood samples were collected at steady-state (trough values before drug administration). Samples were collected and analysed over a period of 3 years with an interval of at least 3 months between each sample. Dosage regimens ranged from 200 mg/day to 600 mg/day. Results were expressed as mean \pm standard deviation and variability in plasma concentrations were presented as a CV expressed in percentage. A minimum of three measurements was needed to calculate CV%.

Results: 21 patients were evaluated with a mean number of 5 measurements per patients (3–10) and a total of 108 samples. Sex-ratio (M/F) was 0.48, mean age and weight were respectively 55 \pm 16 years and 76 \pm 19 kg. Concerning the 400 mg/day group of patients (61 samples), mean imatinib plasma concentration was 1130 \pm 640 ng/mL, ranging from 250 to 2800 ng/mL (total CV of 57%) with 18 subtherapeutic concentrations (a plasma threshold of 1000 ng/mL is associated with major molecular response). For all dosage regimens (21 patients), mean intra-individual variability of imatinib was 31%, ranging from 9.7% to 70%. Possible causes for pharmacokinetic variability are multiple: drug/drug interactions (due to metabolism of imatinib with CYP3A4 and transport by Pgp), poor compliance, genetic polymorphisms. A value of 31% for intra-individual variability can be considered low.

Conclusion: The high interindividual and relatively low values of intrapatient variability in plasma support therapeutic drug monitoring. Nevertheless mean plasma concentrations were often closed to therapeutic threshold and our data suggest the need of regular TDM measurement.

9234

POSTER

Cytogenetic Abnormalities in the Spleen Detected by FISH in Patients With Chronic Lymphocytic Leukemia/Small Lymphocytic Lymphoma

S. Norin¹, A. Wallblom¹, B. Sander², E. Kimby¹. ¹Karolinska University Hospital, Hematology Center, Stockholm, ²Karolinska University Hospital, Department of Laboratory Medicine, Stockholm, Sweden

Background: In chronic lymphocytic leukemia/small lymphocytic lymphoma (CLL/SLL), the prognostic importance of cytogenetic abnormalities in blood and bone marrow is well established. However, only limited data has been published regarding frequency and prognostic impact of aberrations in other tissues.

Materials and Methods: We identified 62 patients with CLL (n=57) and SLL (n=5) who underwent splenectomy between 1989–2010. Fluorescence in situ hybridization (FISH) could be successfully performed on spleen tissue in 60 cases for detection of the cytogenetic abnormalities 11q-, 13q-, 17p- and trisomy 12. The results were compared with available FISH analyses on blood and/or bone marrow. To avoid false-positive results due to incomplete nuclei present in the tissue sections, the cut-off rate for the deletions was set to 40%. For trisomy 12 the cut-off was set to 5%.

Results: Cytogenetic aberrations were detected in 75% of the patients; in 31 cases a single abnormality and in 16 multiple aberrations. The most common aberration was 13q-, detected in 69% of cases. There was a significant correlation between the frequency of 13q and 11q deletions in spleen and blood/ bone marrow.

In 4 of 27 cases, new abnormalities were detected in blood/bone marrow samples after splenectomy. Two cases with a heterozygous 13q deletion in the spleen developed homozygosity for 13q- in blood and a new 11q- clone

was found in two patients. One patient had both homo- and heterozygous 13q- clones in the blood before splenectomy as well as in the spleen, but in repeated blood samples taken nine to ten years later only a heterozygous clone remained.

After a median follow-up of 43 months after splenectomy, 23 of 62 patients are alive. Median time to next therapy was 9 (range 0–255+) months. Cytogenetic aberrations had a significant impact on overall survival dividing patients into three categories, (1) 13q- as a sole abnormality, (2) normal karyotype or trisomy 12 and (3) abnormalities involving 11q- or 17p- (p<0.05). Patients with 11q- and/or 17p- did also have shorter therapy-free survival (p<0.01).

Conclusions: In our study on splenectomized patients, cytogenetic aberrations could be reliably detected by FISH on paraffin-embedded sections and did influence overall and therapy-free survival. Clonal evolution could be found in a few patients.

9235

POSTER

BCL2 Expression Correlates With Surface Immunoglobulin Levels and Prognosis in Follicular Lymphoma Patients

A. Barreca¹, C. Martinengo², L. Righi³, A. Chiappella⁴, M. Ladetto⁵, L. Chiusa⁴, A. Stacchini⁴, R. Chiarle². ¹University of Turin, Department of Biomedical Sciences and Human Oncology, Turin, ²University of Turin, Department of Biomedical Sciences and Human Oncology Center for Experimental Medicine and Clinical Studies (Cerms), Turin, ³University of Turin at S. Luigi Hospital, Department of Clinical and Biological Sciences Division of Pathology, Orbassano, ⁴ASO San Giovanni Battista Hospital, Department of Biomedical Sciences and Human Oncology, Turin, ⁵University of Turin, Department of Experimental Medicine and Oncology, Turin, Italy

Background: Almost 90% of Follicular Lymphoma (FL) carry a t(14;18)(q32;q21), which juxtaposes the immunoglobulin heavy chain (IgH) enhancers to the Bcl2 gene and results in an overexpression of the BCL2 protein. Most cases of FL show heterogeneous intensity of IgH or BCL2 protein. The aim of this study was to correlate IgH and BCL2 expression and to ask whether different levels of BCL2 were associated to resistance to therapy and prognosis in FL patients.

Material and Methods: We analyzed 103 freshly isolated FL cases by flow cytometry and immunohistochemistry to assess BCL2 levels, and correlate BCL2 expression to survival. Isolated cells from FL patients and cell lines were studied for BCL2 regulation and resistance to apoptosis.

Results: We found a strong positive correlation between IgH and BCL2 expression in FL cases (P<0.0001) as well as in subpopulations of FL cells within individual patients. A concordant regulation of both IgH and BCL2 in lymphoma cells carrying a t(14;18) translocation was found. Remarkably, primary FL cells expressing high BCL2 were more resistant to Rituximab or doxorubicin than FL cells with low BCL2 isolated from the same patient. Finally, patients with low BCL2 levels had significantly higher probability of survival as compared to patients expressing high levels of BCL2 (P<0.05).

Conclusion: We show a strong correlation between IgH and BCL2 expression in FL cells, suggesting mechanisms that concordantly regulate both IgH and BCL2 transcription in translocated cases. Heterogeneous expression of BCL2 affects sensitivity to therapy and overall survival.

9236

POSTER

Phosphatidylinositol-3 Kinase I Inhibitor BKM120 Induces Cell Death in B-chronic Lymphocytic Leukemia Cells In-vitro

L. Amrein¹, M. Shawi¹, L. Panasci¹. ¹Segal Cancer Center, Oncology, Montreal QC, Canada

Background: B-chronic lymphocytic leukemia (CLL) is characterized by the accumulation of CD5+ B-lymphocytes that are long-lived *in-vivo* but die quickly by apoptosis when cultured *in-vitro*. The phosphatidylinositol-3 kinase (PI3K) cascade is a critical component of survival signalling including PI3K-activated Akt (phosphorylated Akt) which inhibits cell death pathways by inactivating pro-apoptotic proteins. However, PI3K has increased activity in CLL lymphocytes as compared to normal B lymphocytes. There are three classes of PI3Ks of which class I is the most clearly implicated in human cancer. The PI3K- δ inhibitor, CAL-101, promotes caspase-dependent apoptosis and abrogated protection from spontaneous apoptosis induced by CD40 in primary CLL lymphocytes *in-vitro*. Thus PI3K pathway appears to play a critical role in B-CLL cell survival. BKM120 is a pan class I PI3K inhibitor developed by Novartis. Phase I trials demonstrated that plasma concentrations of 3–5 μ M can be obtained. In view of the critical role of PI3K in CLL homeostasis, the activity of BKM120 was examined in CLL lymphocytes.

Material and Methods: BKM120 cytotoxicity was assessed by the MTT assay in primary B-CLL lymphocytes and the MEC-2 B-CLL cell line.