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Selenium status in the body and proliferative activity of malignant cells

Sir,

Avanzini and co-workers recently reported the results of their study on serum selenium concentrations in patients with newly diagnosed lymphoid malignancies.¹ Interestingly, as compared to controls, selenium (Se) levels were significantly lower in patients with non-Hodgkin's lymphoma (in a representative sample) in IV stage and/or in those with high grade disease. However, literature reports of serum Se levels in patients with lymphoid malignancy or solid tumors are discordant.²⁻⁴ These discrepancies may be due to case series that are not directly comparable among themselves or with healthy control cohorts.

It must be borne in mind that the elevated variability of serum Se may be due to factors other than cancer such as age, sex, body mass, dietary habits, life style (alcohol, smoking), intercurrent disease and medications.

When considering the various biological roles of Se (proliferation and expression of oncogenes by both normal and malignant cells, carcinogen metabolism, cellular immune response, prevention of oxidative stress, apoptosis) and when addressing the topic of whether Se is a risk factor or a protection against cancer,⁵ one must evaluate selenium levels both in serum and in biological material that integrates selenium intake and reflects its status over the medium and long term (4 months for red blood cells, 12 months or longer nails and hair). These studies are however methodologically complex, and at present various types of investigations (prospective, environmental, epidemiological, ecological) have failed to provide conclusive results.

Several factors may influence Se exchanges between labile pools, deposits and cancer tissue, and we wish to point out one important factor which has so far received little attention but which might influence selenium profiles, namely the proliferative activity of cancer cells. Of particular interest along this line of thought is the finding by Avanzini and co-workers of an inverse relationship between serum selenium and β_2 -microglobulin, an important index of the turnover of neoplastic cells.

In an ongoing study on Se levels in the serum and hair of women with breast cancer (unpublished data), we observed that patients recruited at an early clinical stage had lower serum Se and higher Se hair content with respect to patients at a more advanced stage or to healthy controls:

Stage 0-I (n=42): serum Se mean value 76.2±21.7 µg/L, hair Se content geometric mean 416.5 µg/g

Stage II-IV (n=44): serum Se mean value 81.5±22.4 µg/L, hair Se content geometric mean 335.2 µg/g

Controls (n=86): serum Se mean value 88.6±26.4 µg/L, hair Se content geometric mean 370.5 µg/g

Though our data fell short of statistical significance, due in part to the wide spread of values in the series, the findings are suggestive in light of the kinetic properties of tumor cells.⁶ Indeed the relationship between Se and cellular growth emerged from our *in vitro* studies that showed different Se accumulation and effect according to cell density in the lymphocyte cultures used in our experiments.⁷ Breast cancer shows a Gompertz-type growth curve with an exponential increase in the early proliferative phases. These events, whose underlying mechanisms also require further investigation in terms of host-tumor interactions, might influence different aspects of Se distribution in various body districts.

In conclusion, further experimental and epidemiological studies are warranted to better elucidate the relationships between Se status and carcinogenesis; however, determination of Se levels in subjects with malignancies will only contribute data useful for the clinician when it is performed across several body districts (blood, depots, healthy and diseased tissue) and interpreted in the light of the proliferative characteristics of the tumor and of the intricate tumor host relationships.

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Selenium and lymphoid malignancies (Reply)

Sir,

Piccinini *et al.*'s hypothesis that the proliferative activity of malignant lymphoid cells and of cancer cells in general might influence selenium status is interesting and is partially supported by prior data, including our own.¹ However, due to the limited evidence available on this topic, further data are needed to confirm that enhanced selenium uptake by neoplastic tissue may vary according to the mitotic activity of the cancer cells.

Piccinini and colleagues also addressed a fundamental issue in epidemiologic and clinical research on the health effects of selenium: the methodology for exposure assessment and, in particular, the use of biomarkers as surrogate measures of selenium exposure (represented in most individuals by dietary intake). Selenium content of serum, plasma, erythrocytes, whole blood, hair, toenails, and urine are among the biomarkers most frequently used in epidemiologic and clinical studies. Serum, plasma and urine selenium are short-term markers of exposure, whereas the remaining indicators tend to reflect long-term selenium intake. The limitations of these indicators as surrogate measures of intake have been reviewed.² Selenium-dependent glutathione peroxidase activity has also been evaluated as a possible biomarker of exposure, but it does not appear to be a reliable indicator of selenium intake since the correlation between the two parameters is not linear² and, what is more, glutathione peroxidase activity may be induced by oxidizing agents³ (including selenium itself).^{4,5}

In our clinical studies^{1,6} we evaluated selenium exposure through determination of serum selenium content, a sensitive short-term selenium marker,² because we were interested in a possible relationship between the clinical characteristics of lymphoid malignancies and recent changes in selenium status. Obviously the characteristic that makes serum selenium content of interest in clinical research, i.e. its ability to reflect short-term selenium intake, also represents a limitation in an epidemiologic setting, particularly in retrospective studies where selenium status is likely to be affected by the disease, at least in some body tissues. This is why we did not consider our results to be contradictory to the prior hypothesis of a direct association between selenium exposure and the risk of lymphoid malignancies,⁷ though they did not add any evidence to support this hypothesis.

Biomarkers, however, may not adequately reflect selenium intake due to factors such as gender, body mass, medical

treatment.⁸⁻¹⁰ Moreover, concurrent exposure to other factors such as cadmium or methionine may modify the distribution (and therefore the levels) of selenium in peripheral tissues.¹¹ An influence of cadmium on selenium metabolism might explain at least in part why lower selenium levels have generally been observed in smokers as opposed to non-smokers,^{8,9,12} in spite of the fact that tobacco smoke represents a source of selenium exposure.¹³ The limitations of biomarkers in exposure assessment have suggested the usefulness of evaluating average selenium exposure through estimation of usual dietary intake, but unfortunately this methodology is also subject to limitations due to the high variability in the selenium content of foodstuffs in several geographical areas.^{12,14} The degree of correlation between selenium intake, as estimated through food diaries or other techniques, and biomarkers as surrogate measures of exposure has been found to be satisfactory in some studies but not in all.^{8,10}

An important advantage to estimating dietary intake would be the possibility of determining the specific chemical forms of selenium to which the human body has been exposed before any *in vivo* metabolic conversion occurs. Whichever approach is adopted in the assessment of selenium exposure – dietary intake or biomarkers – other factors such as exposure to heavy metals, which can markedly influence the biological activity of selenium, should also be considered.⁵

Despite the complexity of epidemiologic studies on the relationship between selenium and cancer, we agree with Piccinini *et al.* on the need to investigate this issue further, especially in the light of the results of two recent studies that have analyzed the effect of selective long-term selenium exposure (self-administered or accidental) on cancer risk.^{7,12}

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When to perform peripheral blood progenitor cell collection in hematological patients?

Sir,

we read with interest the paper by Torretta *et al.*¹ on the experience of Pavia University regarding circulating progenitor cell collection in cancer patients. The authors stated that, using daily flow cytometric monitorization of CD34⁺ cells in the peripheral blood (PB), collections were started when these cells reached a value of 20 μ L. However, the possibility of harvesting even though circulating CD34⁺ cells were below 20 μ L (between 10 and 20 μ L) was always considered in relation to the particular clinical history, state of disease and therapeutic strategy adopted for each patient.

In agreement with this latter statement, we would like to describe our experience on this topic. As Torretta *et al.* reported, we usually start leukaphereses when white cells in the PB, evaluated in the morning just before collection, are greater than 1000 μ L and CD34⁺ cells greater than 20 μ L. However, when making clinical decisions, we have to consider that some patients, due to their clinical situation such as secondary myelodysplasia, exhausted marrow, heavy previous chemo- or radiotherapy may have serious problems in mobilizing CD34⁺ progenitor cells in the blood.^{2,3} This leads to the need for a larger number of procedures to obtain the optimum yield of progenitor cells for a safe engraftment. In this setting, some authors suggest a threshold dose of 2×10^6 /kg CD34⁺ cells, while others refer that less than 5×10^6 /kg, although able to restore hemopoiesis in most cases, can be responsible for delayed engraftment or defective platelet reconstitution and recommend collecting not less than 8×10^6 /kg CD34⁺ cells to ensure a rapid, complete and sustained hematopoietic recovery.^{4,5} We retrospectively analyzed our data on 54 patients suffering from hematological malignancies (21 multiple myelomas, 14 acute myeloid leukemias, 10 non-Hodgkin's lymphomas, 9 Hodgkin's disease) who received high-dose mobilizing chemotherapy plus growth factor administration (G-CSF and in some cases GM-CSF) at our Institution between April 1993 and July 1996. The total number of leukaphereses performed was 159 (mean number per patient 2.9, range 1-4). About 9 liters of blood were processed for each patient.

The amount of CD34⁺ cells collected at each leukapheresis was analyzed in relation to the number of CD34⁺ cells in the PB, in order to evaluate the predictive yield capacity of the latter. A

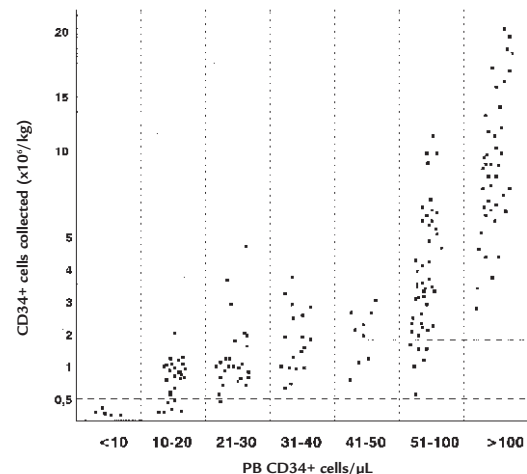


Figure 1. CD34⁺ progenitor cell content ($\times 10^6$ /kg) in leukaphereses in relation to the PB CD34⁺ cell concentration (μ L).

strong correlation was found between the two variables ($r = 0.86$; $p < 0.0001$), as suggested by linear regression analysis. The best results were obtained when the number of CD34⁺ cells in PB was greater than 50 μL , while the worst results were related to a level of less than 10 μL (Figure 1). Only in 5 cases, because of the particular clinical history of the patients, did we perform leukapheresis when circulating CD34⁺ cells were $< 10 \mu\text{L}$. None of these reached $0.5 \times 10^6/\text{kg}$ CD34⁺ cells (mean value collected was 0.23, range $0.05\text{--}0.4 \times 10^6/\text{kg}$). With more than 10/mL CD34⁺ cells in PB we collected no less than $0.5 \times 10^6/\text{kg}$, with rare exceptions. When CD34⁺ cells exceeded 20/mL we usually reached more than $1 \times 10^6/\text{kg}$. No statistical difference was found in patients with CD34⁺ cell values ranging from 21 to 50 μL . Finally, in our hands, more than 50 μL circulating CD34⁺ cells ensured a collection greater than $2 \times 10^6/\text{kg}$ in most patients.

Though the quality of a leukapheresis does not depend only upon the absolute number of CD34⁺ progenitors present,^{6,7} our data confirm that daily estimation of the circulating CD34⁺ cell number by flow cytometry may guide our clinical decisions and may offer a useful tool for predicting the number of procedures to perform.

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All-trans retinoic acid might also induce apoptosis in freshly isolated chronic myeloid leukemia cells

Sir,

we read with interest the recent letters by Martinelli and coworkers¹ and Zinzani and coworkers² on the induction of apoptosis by the nucleoside analogs fludarabine (FAMP), 2-chlorodeoxyadenosine (2-CdA) and 2-deoxycoformycin (DCF), whether used alone or in combination with α -interferon (α -IFN), in freshly isolated leukemic cells from chronic myeloid leukemia (CML). Apoptotic cell death, as demonstrated by electrophoretic gel DNA fragmentation pattern, was induced by both FAMP¹ and 2-CdA,² either alone or in combination with α -IFN, whereas DCF,² with or without α -IFN, failed to do so. The authors focused on the opportuneness of promoting further *in vitro* and *in vivo* studies with these two promising adenosine analogs, possibly employing assays able to measure pro-

grammed cell death. We agree with the authors about the timeliness of exploring new effective drugs capable of driving the CML clone into apoptosis, and on this point we would like to provide further evidence in support of their *in vitro* findings.

We recently tested the *in vitro* capability of FAMP and all-trans retinoic acid (ATRA) to drive peripheral myeloid cells from untreated Ph⁺ CML patients in chronic phase into apoptosis. Apoptosis was measured by using simple and reliable flow cytometric methods based on decreased forward light, increased right angle scatter and reduced propidium iodide fluorescence stainability.³ These methods, as compared to electrophoretic gel DNA fragmentation assays, which allow only bulk apoptosis measurement, are able to detect programmed cell death on a single cell basis. In our model, CML cells cultured alone in standard complete medium (RPMI 1640 plus 10% FCS) showed a low apoptotic cell rate (6.8% at 96 hours of culture) at all the different time points tested (24, 48, 72, 96 hours). By contrast, when cultures were performed in the presence of FAMP (5 μM) apoptosis reached 26.3% and 31.7% at 72 and 96 hours, respectively. Similar results were obtained when FAMP was substituted with ATRA (3 μM). This agent, which was previously shown to induce apoptosis in both acute promyelocytic leukemia⁴ and chronic lymphoproliferative disorders,⁵ also drove CML cells into apoptotic cell death (28.6% at 72 and 40.5% at 96 hours of culture). Apoptosis occurred mainly via terminal myeloid differentiation of the leukemic clone, as demonstrated by cytospin morphological and cytochemical examinations.

Taken together, these findings further support the importance of focusing on inducers of programmed cell death as promising new agents in the management of chronic phase CML, and provide a rationale for the employment of either purine analogs or ATRA in pilot clinical trials.

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Cryosupernatant in thrombotic thrombocytopenic purpura (TTP): is it really useful?

Sir,

Perotti *et al.* in their paper entitled *Cryoprecipitate-poor plasma fraction (cryosupernatant-CPP) in the treatment of thrombotic thrombocytopenic purpura at onset. A report of four cases*¹ demonstrated that CPP can induce a more rapid improvement in the clinical manifestations of TTP than fresh frozen plasma, while laboratory parameters show slow normalization. We treated six patients with plasma exchange (PE) and CPP (two relapses at four and six years, respectively; four cases at first diagnosis).

The clinical manifestations of these patients at onset are reported in Table 1.

Table 1. Laboratory parameters and clinical manifestations at onset in six patients with TTP.

	Hb g/dL	Plt x10 ⁹ /L	LDH U/dL	Neurological findings	Kidney findings
RR	6.5	9.0	2750	no	no
MD	6.8	5.0	1498	no	no
DA	11.0	2.0	5786	yes	yes
CM	7.5	13.0	2450	no	no
PLM	6.7	14.0	1743	yes	yes
BS	7.2	11	1570	yes	no

All patients received the first plasma exchange employing fresh frozen plasma, and CPP was used in the following procedures. PE was performed daily until normalization of the platelet count and serum LDH levels.

Two patients (B.S. and P.M.L.) died 8 and 5 days, respectively, after beginning plasma exchange, and the addition of high-dose iv immunoglobulins in one case did not produce any effect. Clinical remission was reached in four patients. In one case, neurological disorders at onset such as amnesia and paresthesia improved after three PE procedures.

Normalization of platelet count was obtained after 7, 11, 13 and 17 plasma exchange procedures, respectively. Our experience confirms the data reported by Perotti *et al.* regarding hematological recovery, while the neurological disorders that appeared during the treatment in two patients did not seem to be influenced by the addition of CPP.

These observations seem to limit the efficacy of CPP as compared to fresh frozen plasma in the treatment of TPP.

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Lymph node myeloid metaplasia associated with chronic neutrophilic leukemia

Sir,
chronic neutrophilic leukemia (CNL) is a rare disorder characterized by neutrophilia due to mature elements and organ infiltration, including both hepatosplenomegaly and lymph node infiltration.^{1,2} Around 78 cases have been reported up to now and no one else has described the presence of myeloid metaplasia in lymph nodes as we have in this study.

A 68-year-old man presented weight loss, enlarged lymph nodes, hepatosplenomegaly and ascites. WBC count was $156 \times 10^9/L$ (84% neutrophils, 1% monocytes, 2% basophils, 3% bands, 2% atypical lymphocytes, 3% metamyelocytes, 5% myelocytes). Hemoglobin was 8.7 g/dL, platelets $68 \times 10^9/L$, uric acid was 15.8 mg/dL, alkaline phosphatase 2457 U/L, LDH 915 U/L, granulocytic alkaline phosphatase 300 U. Renal and liver function was normal. Paracentesis revealed ascitic liquid with $3.6 \times 10^6/L$ cells (85% mature neutrophils). Bone marrow biopsy displayed increased cellularity and granulocytic hyperplasia compatible with CNL. Cytogenetic study was normal (46,XY). Molecular biology did not demonstrate a bcr-abl translocation. A lymph node biopsy showed massive substitution of the normal lymph node architecture by hematopoietic cells, including elements of all three series; within the granulo-

cytic line numerous polynuclear cells were conspicuously intercalated among immature cells. Megakaryocytes were also frequent, and some of them presented phagocytic phenomena. The patient was treated with hydroxyurea, which provided good WBC count control; however, two months later he developed hepatorenal failure and died on day +71 after diagnosis.

Myeloid metaplasia has not been previously described in CNL. Chronic neutrophilic leukemia was recently characterized as a distinct myeloproliferative disease with a specific molecular marker (bcr/abl with C3/A2 junction).^{3,4} Our finding of myeloid metaplasia is consistent with the myeloproliferative nature of CNL.

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Prolonged low doses of oral etoposide may be effective in individual patients with advanced lymphoproliferative disorders refractory to aggressive chemotherapy

Sir,
etoposide, a semisynthetic podophyllin derivate, is currently employed in the treatment of several malignancies. The antitumor efficacy of etoposide is highly schedule dependent and it has been demonstrated that five-day administration is superior to single day administration.¹ Oral etoposide has greatly facilitated the use of multiple-day schedules and the drug displays good activity in many extrahematological malignancies as well as in lymphomas.^{2,3} From December '93 to May '94, we treated with prolonged low doses of oral etoposide 24 patients with advanced hematological malignancies not eligible for further intensive approaches due to age >60 years and/or severe previous infective complications. Of these, ten patients had ALL, 8 AML, 4 NHL and 2 had CML.

Etoposide (50 mg/m² per day) was administered orally for 21 consecutive days; patients who showed a response or stable disease on day 28 received a second and a third cycle of the same treatment. Responder patients received maintenance etoposide at the same dosage 10 days/month until relapse. Three patients (12.5%) died during the first cycle of intracerebral hemorrhage (1 patient) or infective complications (2 patients); no other toxic deaths were observed. All other patients received at least 2 cycles. Two ALL and 1 NHL patient (12.5%) obtained a complete remission of short duration (2, 3, 7 months); 1 ALL and 2 NHL patients (12.5%) achieved a partial response. No patient with myeloproliferative disease attained a response. Data from responder patients are shown in Table 1.

As for toxicity, 15/24 patients (62.5%) suffered febrile episodes during treatment (4 sepsis, 6 bronchopneumonia, 1 fungal sinusitis and 4 fever of unknown origin). One patient developed an intracerebral hemorrhage, while 8/24 patients (33.3%) displayed cutaneous hemorrhagic manifestations. Ten patients (41.6%) had mild nausea (WHO < 2) and vomiting, and 6/24 (25%) mucositis (WHO 2).

Table 1. Data from responder patients.

Sex	Age	Disease	Disease status	Previous treatments	Response	N.cycles to response	Response duration (months)
F	22	ALL	2nd resistant relapse	VCR+DNR+CTX IDAC+IDA ARA-C+MTX+PDN	CR	2	2
M	34	ALL	resistant	AIL 7603 IDAC+IDA	CR	2	3
F	64	NHL	1st relapse	F-MACHOP	CR	3	7
F	41	ALL	1st relapse	MAGRATH	PR	3	4
F	77	NHL	1st resistant relapse	CHOP PROVECIPI	PR	2	4
M	22	NHL	resistant	MACOP-B +ABMT	PR	2	3

Despite the heterogeneity and low number of patients treated, some preliminary observations can be made. The 25% overall response rate achieved is encouraging and compares favorably with other single-agent approaches, as well as with intravenous etoposide.⁴

Previous studies have also reported encouraging results using oral etoposide in untreated elderly AML patients.⁵ By contrast, the results reported in ALL patients are disappointing.⁴ No AML patient in our study responded. However, ALL patients showed a 30% response rate (2/10 CR and 1/10 PR). Overall, oral etoposide achieved better results in advanced lymphoproliferative disease (6/14 responder patients) than in advanced myeloproliferative disease (0/10 responder patients). The reason for this behavior is unclear.

The toxicity of the schedule was acceptable and no patient discontinued the treatment because of nausea. As expected from the advanced disease status and heavy pretreatment of these patients, median response and survival duration were short. The use of oral etoposide in less advanced disease and/or its association with other drugs are possible ways of improving these results.

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Circulating antiplatelet antibody specificity in children with immune thrombocytopenic purpura at onset

Sir,
immune thrombocytopenic purpura (ITP) is caused by the interaction of platelet reactive antibodies with platelet surface antigens, which determines accelerated platelet destruction of antibody-coated platelets. Two forms of childhood ITP may

occur: a syndrome similar to adult chronic ITP, and an acute self-limiting form of the disease. Only few reports have been published about antiplatelet antibody specificity in paediatric ITP; some of them^{1,2} studied the specificity of circulating antiplatelet antibodies by testing patient sera by immunoblotting. However, certain conformational antigens on platelet membrane are destroyed by this technique. Moreover, these reports concerned small paediatric ITP populations. It was suggested that the presence of circulating anti-GPIIb/IIIa antibodies may be useful in differentiating acute from chronic ITP in children;³ however, in a recent paediatric survey⁴ no difference between the two ITP forms was found. We investigated the specificity of circulating antiplatelet antibodies of ITP children at onset, in order to assess whether it may represent a marker of evolution of the disease. Sera were collected from 74 ITP children (4 months to 13 years, mean age of 5.5 years) at onset before beginning therapy. Forty-nine patients recovered within 6 months from the initial diagnosis (acute ITP), whereas 25 patients developed chronic disease (mean duration 2.2 years, range 1 to 5 years). Antibody specificity was assessed by indirect MAIPA assay refined according to Kiefel *et al.*;⁵ we looked for anti-GPIIb/IIIa and anti-GPIb/IX IgG antibodies.

Anti-GPIIb/IIIa antibodies were found in 19/49 (38.8%) and in 8/25 (32.0%) acute and chronic ITP, respectively. Antibodies to GPIb/IX were detected in 15/49 (30.6%) acute ITP and in 7/25 (28.0%) chronic ITP. Thus, in our experience we did not find any significant difference between acute and chronic ITP, evaluating both anti-GPIIb/IIIa and anti-GPIb/IX antibodies. This study reports on the investigation of circulating antiplatelet antibodies specificity in the largest sample of acute and chronic ITP children at onset ever analyzed at our knowledge. We conclude that circulating antiplatelet IgG specificity in childhood ITP at onset does not represent a marker to the early recognition of those patients devoted to chronicize. We cannot rule out that autoantibodies against platelet antigens other than GPIIb/IIIa and GPIb/IX were responsible for thrombocytopenia in some of our patients; moreover, antiplatelet IgM should be investigated especially in acute ITP patients. Finally, more useful information could be obtained performing directly MAIPA on patient's platelets.

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Hodgkin's disease in Brazil: a clinicopathologic study

Sir,
the behavior of HD in Brazil, namely its epidemiology, histological distribution, clinical and pathological stages, and the

potent for cure with conventional treatment is not well-known but is of both practical and scientific importance. Spector *et al.*¹ described good results in treating 59 HD patients from rather low socioeconomic background using C-MOPP/ABV chemotherapy.² The authors carried out a study in 134 cases of adult patients (age above 15 years; 76 males and 58 females) from 1985 to 1994. HD appeared predominantly in males ($R=1.31$); the incidence peaked between 21 and 30 years and presented a descending curve after 30 years of age.

The most frequent histologic type was nodular sclerosis (NS)(50.4%) followed by mixed cellularity (MC) (34.6%), lymphocyte depletion (LD) (9.0%) and lymphocyte predominance LP) (5.2%) (Table 1). NS-I type prevailed over NS-II (30.8% and 19.6%, respectively). The cervical region, followed by the spleen and retroperitoneum, was the most frequent primary sites of disease. A bulky mass was found in 20% of the cases, mostly in the mediastinum. Bone marrow (BM) was involved in 16.5% of the patients: none LP type, 3.0% in NS, 6.8% in MC and over 40% in LD type. Anemia was directly related to BM infiltration ($p=0.003$). B symptoms as well as higher levels of mucoproteins were more frequent in males ($p=0.02$ and 0.03 , respectively). In contrast, early clinical stage occurred mainly in females ($p=0.004$). Probability of survival after 5 years was 83%. Disease free survival was 52%. No statistically significant differences were found between sexes, NS and MC types or NS-I and NS-II subtypes. Complementary factors that negatively influenced the survival were BM infiltration ($p=0.02$), lymphopenia ($p=0.01$) and serum mucoproteins above 7 mg/dL ($p=0.01$).

Table 1. Distribution of histologic types according to sex.

histologic type	Males		Females		Total	
	n.	%*	n.	%*	n	%
LP	4	5.2	3	5.2	7	5.2
MC	33	43.4	13	22.4	46	34.6
NSI°	17	22.3	25	43.1	41	30.8
NSII°	13	17.1	13	22.4	26	19.6
LD	8	13.1	4	6.9	12	9.0
unclassified	1	1.3	0	0	1	0.8
total	76	100	58	100	134	100

These findings demonstrate that the NS type is possibly the most frequent histologic type, at least in Southeast Brazil, a pattern similar to that found in developed countries.³

One interesting clinical feature concerns the period between the first sign of clinical symptoms and medical diagnosis. We observed that this period was significantly shorter in women and in NS-II type, when compared to other types, particularly NS-I. Among men, a history of disease up to 24 months could

be detected. On the other hand, the shorter disease history interval among NS-II patients may be due to more rapid progress of lymph node enlargement. This fact points to different biological behavior for NS-I and NS-II, although this difference did not influence response to therapy or prognosis.

One of the most important aspects in our study is related to the distribution of patients in clinical stages. The validity of the Ann Arbor staging system has been established.⁴ In our study, a large number of cases were in advanced stages (68.8%) and presented B symptoms (67.2%). This is different from developed countries, where such cases rarely exceed 50%. The larger extent of disease at diagnosis in our patients could also be documented by the number of sites of involvement, the prognostic significance of which has already been stressed.⁵ All these features point to a delay in diagnosis of HD in our population. This is also confirmed by the high incidence of BM involvement, anemia, high ESR, and mucoprotein levels, as well as a high frequency of lymphopenia. Nevertheless, response to treatment was adequate according to the results described for the same stages and with the same therapy schedules.

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MEETING ANNOUNCEMENTS

24th National Congress of the Italian Association of Pediatric Hematology and Oncology

Bologna, Italy, Palazzo dei Congressi
June 1-3, 1997

Any further information available through the Scientific
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