when compared with epithelial tumours, mesenchymal tumours of the gastrointestinal tract are rare, with gastrointestinal stromal tumours (GISTs) being the most frequent. Possibly derived from the interstitial cells of Cajal of the autonomic nervous system, the tumour cells in GIST characteristically express c-kit (CD-117; a transmembrane tyrosine kinase receptor) and can be detected with an antibody that binds to the C-terminal domain of this molecule, making the diagnosis of this tumour easier than in the past.7–18 In contrast, estimating the possible clinical outcome and predicting the biological behaviour of these tumours remain uncertain. Since 1984, when a subtype of stromal tumours, the so called gastrointestinal autonomic nerve tumour was described, numerous macroscopic, histological, cell kinetic, and molecular parameters, including the size and localisation of the tumours, the Ki-67 and mitotic indices, DNA ploidy, the S phase fraction, and the presence of specific mutations, have been suspected and partly proved to have an impact on the prognosis of the disease.19–23 In spite of extensive research, to date, there is no reliable single predictive factor useful for routine histopathology. Various grading systems have been suggested, which have included histological features such as the phenotype of the tumour cells (epithelioid versus spindle) together with pleomorphism and the mitotic count; or the grade of pleomorphism and the extent of necrosis together with the mitotic count. This has resulted in several different grading systems, but there is no widely accepted, consistently applied system yet in use.19–23

“In spite of extensive research, to date, there is no reliable single predictive factor useful for routine histopathology”

Recently, a new clinicopathological approach to assess the risk of aggressive behaviour was proposed by Fletcher et al.19

This scheme is based upon the assessment of the two parameters thought to be the most predictive and “simplest to determine”; namely: the mitotic count and the size of the tumour. This resulted in a four grade scale: very low, low, intermediate, and high risk of aggressive behaviour.

MATERIALS AND METHODS

Case selection
Twenty three gastrointestinal tumours of mesenchymal origin that had been reported in the past 10 years were selected for histopathological review from the archives of the department of pathology, Medical University, Pécs, Hungary. Immuno-histochemical examination revealed muscle origin (leiomyosarcoma) in three of the cases showing synchronous c-kit negativity and diffuse, strong smooth muscle actin (SMA) and/or desmin positivity, therefore all have been excluded from the study. Only the cases with evidence of clinically malignant behaviour (metastatic and/or locally recurrent cases) were included for further, detailed examination. Of the seven cases that were analysed, four had multiple liver metastases, two metastasised to the regional lymph nodes, another metastasised to the peritoneal/retroperitoneal regions, and one had multiregional involvement including the omentum, peritoneal layer, mesentery, and retroperitoneum (fig 1A). One of the patients who had liver metastases (case 2) also had recurrent tumour four years after the first presentation and surgery (fig 1B,C). The age of the patients varied from 26 to 77 years (mean, 58). The male to female ratio was 5 : 2.

Abbreviations: FNCLCC, French Federation of Cancer Centers; GIST, gastrointestinal stromal tumour; HPF, high power fields; SMA, smooth muscle actin
Histology and immunohistochemistry

The tissue blocks were fixed in buffered formalin, routinely processed, cut, and stained according to the standard haematoxylin and eosin histological protocol. Selected blocks and 13 different antibodies were used to determine the immunophenotype of the tumours. Table 1 summarises the antibodies used. We used the recommended dilutions and microwave pretreatment (700 W, 3 × 5 minutes) in standard citrate buffer for all the antibodies. The slides were developed with the Vectastain universal ABC kit and 3-amino-9-ethylcarbazole was used as substrate.

The Mib-1 index was determined by counting 1000 tumour cells in the fields showing the highest number of positive nuclei. All the positive cells were counted, regardless of the intensity of staining. The number of mitoses was counted in 30 consecutive high power fields (HPF; ×400 magnification using a Nikon Optiphot 2 microscope).

Statistical analysis

Statistical analysis (t-test and χ² test) was carried out using Microsoft Excel software. Significance was set at p < 0.05.

RESULTS

The primary tumours were between 35 and 140 mm in the largest diameter (mean, 83). Four of the seven tumours were solitary, and three were multinodular. Two of the tumours localised to the stomach, four to the small bowel, and as mentioned earlier one (case 6) showed multiregional involvement. The macroscopic details are summarised in the table 2.

Histologically, two of the tumours had an epithelioid phenotype, four had a spindle cell phenotype, and one showed a mixed phenotype. The FNCLCC (French Federation of Cancer Centers) system (originally created for soft tissue sarcomas) and others, such as the schemes established by Newman et al and de Saint Aubain Somerhausen and Fletcher.
were used to grade the cases. The FNCLCC system uses phenotypical and proliferative parameters, such as the mitotic count, the presence and grade of atypia, and necrosis, creating a scoring system with four categories: G0 (benign, score 3), GI (low grade, score 4–5), GII (intermediate grade, score 6–7), and GIII (high grade, score 8–10). The other two schemes, which are very similar to one another, discriminate three categories: benign, borderline, and malignant, combining the epithelioid/spindle cell features with atypia and the number of the mitoses.

According to the FNCLCC system, the cases were classified as three GI, two GII, and two GIII, but as seven malignant (M) tumours according to the other two systems (table 3; fig 1D,E). When a recently published new system, which is based upon the combination of mitotic count and tumour size, was used all of the cases were graded as high risk tumours.

The mitotic index varied from eight in each 30 HPF to 171 in each 30 HPF, being eight, nine and 10 in each 30 HPF in the GI tumours, 30 and 15 in each 30 HPF in the GII tumours, and 45 and 171 in each 30 HPF in the GIII tumours (table 3). Immunohistochemically, all but one case showed moderate to strong reactivity for c-kit (CD-117). The negative case coexpressed synaptophysin, PGP-9.5, and S-100, thereby demonstrating its neurogenic origin. All the cases were positive for vimentin, five of five for neuromelanin specific enolase, four of seven for CD-34 and synaptophysin; they were all negative for chromogranin A, neurofilament, cytokeratin, and desmin.

There was weak and focal reactivity for SMA in four of seven cases. The recurrent tumour seen in case 2 had the same phenotypical features as the original tumour, although the histological grade changed to GII, and immunophenotypically focal, weak SMA and diffuse, moderate S-100 and chromogranin A positivity appeared. The Ki-67 index varied from 4% to 44% (4%, 10%, and 16% in the GI tumours, 15% in both of the GII tumours, and 30% and 44% in the GIII tumours).

| Case | Age/sex | Localisation | Metastasis | Size (mm) | Type
|------|----------|--------------|------------|-----------|-------
| 2    | 62/M     | Small bowel  | Liver      | 55        | S     |
| 3    | 61/M     | Small bowel  | Lymph node | 50        | S     |
| 5    | 26/F     | Stomach      | Liver      | 50*       | MN    |
| 6    | 61/M     | Multiregional| Multorgan  | 115*      | MN†   |
| 14   | 50/M     | Small bowel  | Liver      | 35        | S     |
| 20   | 77/F     | Stomach      | Liver, lymph node | 140 | S     |
| 22   | 67/M     | Small bowel  | Peri-retroperitoneal | 135 | MN‡   |

Size refers to the largest diameter of the tumour.
F, female; M, male; MN, multinodular tumour; S, solitary tumour.
*The largest nodule; †multiple involvement in the omentum, peritoneum, retroperitoneum, and mesentery; ‡multinodular small bowel tumour.

### Table 3

<table>
<thead>
<tr>
<th>Case</th>
<th>Phenotype</th>
<th>Mitosis</th>
<th>Atypia</th>
<th>Necrosis</th>
<th>Grade G0</th>
<th>Grade GI</th>
<th>Grade GII</th>
<th>Grade GIII</th>
</tr>
</thead>
<tbody>
<tr>
<td>2</td>
<td>Spindle</td>
<td>9/30</td>
<td>+</td>
<td>–</td>
<td>GI [2+1+2]</td>
<td>M</td>
<td>M</td>
<td>H 4%</td>
</tr>
<tr>
<td>3</td>
<td>Spindle</td>
<td>171/30</td>
<td>+</td>
<td>+</td>
<td>GI [2+2+4]</td>
<td>M</td>
<td>M</td>
<td>H 44%</td>
</tr>
<tr>
<td>5</td>
<td>Epithelioid</td>
<td>8/30</td>
<td>+</td>
<td>–</td>
<td>GI [2+2+4]</td>
<td>M</td>
<td>M</td>
<td>H 10%</td>
</tr>
<tr>
<td>6</td>
<td>Epithelioid</td>
<td>30/30</td>
<td>+</td>
<td>+</td>
<td>GI [2+2+4]</td>
<td>M</td>
<td>M</td>
<td>H 15%</td>
</tr>
<tr>
<td>14</td>
<td>Spindle</td>
<td>45/30</td>
<td>+</td>
<td>+</td>
<td>GI [2+2+4]</td>
<td>M</td>
<td>M</td>
<td>H 30%</td>
</tr>
<tr>
<td>22</td>
<td>Mixed</td>
<td>15/30</td>
<td>+</td>
<td>+</td>
<td>GI [2+3+2]</td>
<td>M</td>
<td>M</td>
<td>H 15%</td>
</tr>
</tbody>
</table>

Mitosis refers to the number of mitoses in 30 high power fields (×400 magnification). Atypia: –, no atypia; +, atypia in less than 50% of the tumour cells. Necrosis: –, no necrosis; +, less than 50% of the tumour is necrotic. Grade G0 G1 GII GIII: grading according to the FNCLCC system; GI, GI, and GIII corresponds to grade I, II, and III (scores for atypia, necrosis, and mitosis, respectively). Grading according to Newman and colleagues and de Saint Aubain Somerhausen and colleagues: M, malignant. Risk assessment according to Fletcher and colleagues: H, high risk of malignant behaviour. Ki-67 represents the number of positive nuclei/1000 cells expressed as a percentage.
Twelve of the 23 gastrointestinal tumours of mesenchymal origin were non-metastasising GISTs. Seven localised to the stomach, four to the small intestine, and one to the large intestine. The size of the tumours varied from 10 to 135 mm (mean, 45). Ten of the 12 cases were solitary and the remaining two were multinodular. The mitotic index varied from 0 in each 30 HPF to 48 in each 30 HPF, and the Ki-67 index varied from 0% to 45%. All the tumours showed histological signs of malignancy: eight were G1, three were GII, and one was GIII. The different clinicopathological variables, such as the size of the tumour, the multinodular/solitary appearance, the localisation, the mitotic and Ki-67 indices, the phenotype, and the age of the patient were compared and analysed in the two (metastatic and non-metastatic) groups. Using the t-test, there were no significant differences in the mitotic and Ki-67 indices (p = 0.08 and p = 0.09, respectively) or the age of the patients (p = 0.16) between the two groups, but tumour size was significantly greater in the metastatic group compared with the non-metastatic group (p = 0.05). According to the t-test, there were no significant differences in the occurrence of multinodular tumours (p = 0.21), the localisation (p = 0.46), the phenotype (p = 0.14), or the histological grade (p = 0.45) between the metastatic and non-metastatic groups.

**DISCUSSION**

GISTs account for only a small proportion of all gastrointestinal tumours, but make up most of the mesenchymal stromal tumours. The size, intestinal localisation, high mitotic activity and Ki-67 index, aneuploidy of the tumour cells, point mutations or deletions in the c-kit gene, and high histological grade are the major indicators of poor prognosis. Three of our cases were low grade (GI histomorphology), with low or moderate mitotic activity (eight, nine, and 10 mitoses in each 30 HPF) and Ki-67 index (4%, 10%, and 16%). The remaining cases were intermediate or high grade tumours and had higher proliferative parameters (15–171 mitoses in each 30 HPF, 15–44% Ki-67 index) with an obvious malignant phenotype. The mitotic index is thought to be of more value than the Ki-67 index in discriminating between low and high grade cases. However, we found that it was not only those tumours with a high mitotic count and obvious high grade features that behave in a malignant manner, but that low grade neoplasms are also capable of metastasising.

The aim in most malignancies is to find one or more prognostic clinicopathological factor that can aid in predicting the biological behaviour or metastatic capability of the tumour, and this is the case with GISTs. In our study, several clinicopathological variables were analysed and compared between the metastatic and non-metastatic groups of stromal tumours, but only the size of the primary tumour showed a significant difference (p = 0.05) between the two groups. This means that bigger tumours have a greater chance of disseminating or forming metastases, which fits in with the every day experience of malignant tumours in general. Although the difference between the groups was significant, the results should be interpreted with caution because the number of cases investigated was small.

“We found that it was not only those tumours with a high mitotic count and obvious high grade features that behave in a malignant manner, but that low grade neoplasms are also capable of metastasising”

Recently, Joensuu et al used the tyrosine kinase inhibitor, STI-571 (Glivec), in a patient with a metastatic GIST and saw a significant reduction in the mass of this otherwise chemoresistant metastatic tumour. In another recent phase I study, STI-571 was used successfully (25 of 36 patients had objective tumour regression) and safely with only minor and manageable side effects. These successes have intensified the research in this area and increased the number of clinical trials on c-kit positive GISTs. With regard to this therapeutic possibility, it is important that among the metastatic cases in our study the frequency of low grade, c-kit positive tumours with bland histological features was relatively high. Thus, we stress the importance of a careful histological analysis, the necessity of a widely accepted standard grading or risk assessment system (the Fletcher risk assessment system seems to be good at predicting metastatic potential because all of our cases were of “high” risk tumours), and the importance of clinical follow up (including the regular checking of the liver status) even in “benign looking” GISTs, because these cases are potential candidates for this promising new treatment either in adjuvant, neoadjuvant, or palliative settings. Apart from this, a reliable grading system that would help in predicting the biological behaviour of these tumours would help to avoid overtreatment in the future.

**ACKNOWLEDGEMENTS**

The authors wish to thank Novartis Hungária Kft. Pharma for technical assistance.

**Authors’ affiliations**

T Tornóczky, L Pajor, Department of Pathology, Medical School, University of Pécs, Szegedi út 12, H-7643, Hungary
E Kövér, Department of Oncology, Baranya County Hospital, Rákóczi u.2, H-7623, Pécs, Hungary

**REFERENCES**

Molecular diagnosis is valid for culture negative infective endocarditis

Patients with suspected infective endocarditis can expect a more reliable diagnosis after a clinical validation study has confirmed that molecular methods are suitable for identifying the causal organisms in culture negative endocarditis.

A universal bacterial or fungal PCR test was positive in samples of heart valves from 14 of 15 patients with culture negative endocarditis who had undergone surgery; 11 patients had a positive bacterial and three a positive fungal test. Bacteria were identified in eight patients as species of *Staphylococcus*, *Streptococcus*, *Enterobacter* and—unusually—*Tropheryma whipplei* and *Borrelia burgdorferi* (sensu lato). Background contamination made the others undetectable. Fungi were identified as *Aspergillus* and *Candida* spp. Samples from 13 controls were PCR negative. These results correlated with clinical criteria except in two patients; their condition was asymptomatic and evident only at operation.

The retrospective case-control study included 15 culture negative patients with infective endocarditis according to Duke criteria—clinical measures of the condition—and 13 age and sex matched controls who needed valve replacement for other reasons. Samples were taken aseptically during the operation. These were subjected to universal bacterial and fungal PCR with primers from conserved sequences of rRNA genes. Amplified products were analysed by fragment length or restriction fragment length polymorphism, or were sequenced, as necessary. The results were compared with data from the case notes.

Culture negative infective endocarditis has a high death rate and complication rate. Identifying the culprit quickly and reliably means that correct treatment—with surgery if needed—can be started sooner.

*Heart* 2003;89:263–268.