

Prognostic factors for malignant pleural mesothelioma

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Abstract The prognosis for patients with malignant pleural mesothelioma (MPM) is generally regarded as poor, although rare cases of long-term survivors are recognised. Previous prognostic scoring systems have been based on clinical trial populations and are not widely utilised. Population studies consistently confirm non-epithelioid histology, advanced age, and male gender as independent risk factors for poor outcome of MPM. Genetic and immunohistochemical studies continue to provide advances in tumour biology, but no clear validated prognostic factors have been identified to date. Nuclear mitotic and atypia grading systems may provide useful prognostic knowledge; further evaluation is needed. Such biomarkers as soluble mesothelin-related protein and osteopontin provide some prognostic information, though with limitations. The baseline serum neutrophil-to-lymphocyte ratio could also provide prognostic information. Modern metabolic imaging techniques, for example PET/CT, can indicate prognosis by use of baseline total glycolytic volumes (TGV). TGV may also be useful in identifying early responders to systemic chemotherapy treatment. Research to identify clinically useful prognostic factors in MPM remains a priority.

Keywords Mesothelioma · Prognosis · Prognostic markers · Prognostic factors

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Introduction

Malignant pleural mesothelioma (MPM) is a heterogeneous disease with reported median survival of 8–12 months. Because the vast majority of mesothelioma cases are caused by exposure to asbestos, the disease can affect a wide age range, and survival and prognosis vary substantially for different populations. There is no cure for MPM; current recognised chemotherapy regimes may improve survival by a few months [1]. Therefore the prognosis for MPM is generally regarded as poor, although 3–5 % of patients may survive more than five years.

Prognostic factors for any disease are important and useful, because they can enable stratification into risk groups which can, in turn, be used to facilitate treatment decisions, for example identifying those most likely to benefit from aggressive therapy and predicting response to therapy, or identifying those for whom early palliation may be more appropriate. In recent years there has been much interest in “personalised medicine” as a result of advancing genetic, molecular, and immunohistochemistry techniques that assist identification and understanding of tumour phenotypes, both in cohorts and on an individual level. This individual approach is rapidly becoming part of standard best practice for lung cancer [2], but for MPM similar advances in understanding, and in relating understanding to prognosis, have been lacking.

Several scoring systems have been developed and validated [3–6] but all of these are based on specific, selected clinical trial populations, which limits how well they can be generalised to the larger population of patients with mesothelioma. Consequently, these scoring systems are not widely used in clinical practice.

This review article will summarise published data on different aspects of prognosis for MPM, including results from population-based studies, clinically relevant laboratory

characteristics and biomarkers, imaging modalities, and genetic and molecular prognostic factors.

Population-based studies

Although mesothelioma is still regarded as a comparatively rare tumour, several relatively large population-based studies and some smaller studies enable analysis of prognostic variables by use of Cox regression. These national and regional reports are useful because they tend to examine all cases of MPM, and not the more selected populations included in many other studies. A summary of findings from the six most relevant studies is presented in Table 1.

With regard to prognostic variables for poor outcome (shorter survival), frequently-recognised independent risk factors are greater age [7–12], male gender [8, 10, 11], and non-epithelioid or sarcomatoid histology [7, 9–12]. The anatomical site of mesothelioma may be important: a recent study from the Netherlands of 1,353 patients with mesothelioma identified non-pleural disease as a significant prognostic factor (hazard ratio (HR) 1.67, 95 % confidence interval (CI) 1.26–2.22) [7]; and a large study from the USA, including 8,128 patients over 33 years, found localised disease to be protective (HR 0.81, 95 % CI 0.75–0.86) [8]. A smaller regional study from Japan also identified poor performance status (PS of 4 HR 3.22, 95 % CI 1.19–8.74)

and high c-reactive protein (CRP) levels (HR 1.8, 95 % CI 1.06–3.06) as significant independent prognostic variables for poor outcome, with high white cell count (HR 1.49, 95 % CI 0.99–2.26) approaching significance [9].

In summary, when a population-based approach is used to establish prognostic data, increased age, non-epithelioid histology, and male gender are the most consistently reported factors associated with poor outcome, with anatomical site (pleural/non-pleural) likely to be important. This finding may complement a recent report describing remarkably large numbers of extrapleural metastases from MPM found at autopsy, although this report was not designed to demonstrate an affect on survival [13]. Although population-based studies have distinct advantages, there is also a need to examine the biology of MPM to further elucidate prognostic factors for survival with MPM.

Histology

One of the strongest predictors of survival in mesothelioma is histological subtype: many groups, all reaching similar conclusions, have reported this [7, 9]. Epithelioid mesothelioma has slower progression than the sarcomatoid variant. In a large Italian study that included 429 patients, those with sarcomatoid subtype had a hazard ratio of 2.96 (95 % CI 1.28–6.81; $p=0.02$) [12]. In another study involving 4,555

Table 1 Recent population-based studies of malignant mesothelioma examining survival

Location	Years	<i>N</i>	Description	Prognostic variables identified—worse survival Hazard ratio (95 % CI)	Ref.
Netherlands	2005–2008	1353	Median age 69 years; 1-year survival 47 %; 91.1 % male	Increasing age, 1.04 (1.03–1.06) Sarcomatoid histology, 2.45 (2.06–2.90) Non-pleural MPM, 1.67 (1.26–2.22)	van der Bij, 2012 [7]
USA	1973–2006	8128	Median age 72 years; 1-year survival 33 %; 81 % male	Increasing age, 1.02 (1.019–1.023) Male gender, 1.23 (1.16–1.31) Local disease, 0.80 (0.75–0.86) Treatment: surgery, 0.66 (0.63–0.70)	Milano, 2010 [8]
Japan	1996–2006	347	Median age 67 years; median survival 308 days; 87 % male	Age > 70 years, 2.17 (1.36–3.46) Non-epithelioid histology, 1.58 (1.15–2.18) ECOG performance score = 4, 3.22 (1.19–8.74) Raised CRP, 1.8 (1.06–3.06)	Nojiri, 2011 [9]
Italy	1990–2001	4100	62 % aged between 55–74 years; median survival 9.8 months; 73 % male	Age > 75 years, 1.9 (1.7–2.1) Non-epithelioid histology, 1.8 (1.6–2.0) Male gender, 1.1 (1.0–1.2)	Montanaro, 2009 [10]
Germany	1987–2000	498	Mean age 63.1 years, 87 % male	Age > 60 years, 1.29 (No CIs presented) Sarcomatoid histology, 1.89 (No CIs presented) Male gender, 1.72 (No CIs presented)	Neuman, 2004 [11]
Italy	1997–2001	429	Median survival 275 days; 71.6 % male	Age > 75, 1.82 (1.16–2.86) MPM “suspected”, 1.85 (1.16–2.94) Sarcomatoid histology, 2.96 (1.28–6.81)	Marinaccio, 2003 [12]

CI confidence interval; MPM malignant pleural mesothelioma; ECOG European Cancer Oncology Group; CRP c-reactive protein

German patients on a national register, favourable prognostic factors for long-term survival included epithelioid tumour subtype [11].

In a report from the USA on 232 archived mesothelioma samples (all epithelioid histology), the histological analysis and grading of nuclear atypia and mitotic count were found, by multivariate logistic regression (MLR; nuclear atypia severe vs. mild: HR 1.89, 95 % CI 1.15–3.10; mitotic count high vs. low: HR 2.79, 95 % CI 1.69–4.59) to be independent prognostic factors [14]. A grading system based on these results was used to stratify clinicopathological factors by nuclear grade; it revealed statistical association of higher grade nuclear atypia and mitotic count with lymphatic and vascular invasion, and tumour, nodal, and overall stage. These results must be validated externally, and must also be tested on non-epithelioid histology specimens before wider interpretation. A smaller, earlier report on 40 surgical MPM cases, using an index of mitotic activity (in association with assessment of tumour necrosis and vascular endothelial growth factor), failed to demonstrate any independent significance [15].

Angiogenesis is essential for tumour growth beyond a few millimetres. Edwards et al. reported the use of microvessel density (MVD) to score angiogenesis for archived, surgically resected MPM samples. They discovered, by MLR, that increased MVD levels are an independent prognostic factor for poor MPM survival in this surgical group [16]. In a similar report from the same group on a larger cohort of 171 radical, debulking, and palliative surgery specimens, a high MVD score (i.e. increased angiogenesis) was statistically associated with tumour necrosis scores, with the degree of tumour necrosis also contributing independently as a risk factor for shorter survival [17]. A similar report on 40 surgical MPM specimens demonstrated univariate statistical significance of tumour necrosis for shorter survival, but this was no longer significant after multivariate analysis [15].

Immunohistochemical markers

Over the last decade there have been significant advances regarding immunohistochemical markers for MPM and many other solid tumours. A summary of eleven relevant reports that examine survival as an independent variable is presented in Table 2. These reports are mostly on relatively small, and frequently highly selected (e.g. surgical), populations, limiting how well they generalise to the wider population. Also, many of these studies require external validation [15, 18–26]—with the exception of the recent work by Kao et al., who present data validated for both surgical and conservatively managed cohorts with a range of histology subtypes [27].

Aside from the clear utility of immunohistochemical markers in diagnosis of MPM, at present there is no immunostain, or combination of immunostains, providing reliable prognostic information for clinicians.

Differential gene expression

The relationship between gene expression and prognosis has only been reported in a small number of studies, which again were limited by sample size and population selection. One of the most common genetic alterations in MPM is homozygous deletion of the 9p21 locus, which contains, among others, the CDKN2A gene that encodes methylthioadenosine phosphorylase, an enzyme involved in the salvage pathway of AMP synthesis [28]. Two papers have reported the effects of the homozygous deletion of P16/CDKN2A on MPM prognosis. Lopez-Rios et al. describe a large series examining 99 MPM samples (75 male) and report that a homozygous deletion of P16/CDKN2A is independently associated with worse overall survival [29]. A later report from Dacic et al. on epithelioid MPM biopsy or pleurectomy samples from a smaller population ($n=48$) seems to confirm this finding [30]. However, this report was examining a highly selected population of long survivors (median 36 months) with epithelioid histology, which limits how well the study generalises, especially as the use of aggressive surgery in management of MPM is increasingly contentious. Nevertheless, this finding may provide insight into some genetic aspects of long survival.

The serine protease HtrA1 is a potential tumour suppressor gene, and a report from Baldi et al. on 70 MPM samples of mixed histology demonstrated, by MLR, that up-regulated HtrA1 was independently associated with *improved* overall survival [31]. These findings from open biopsy and pleurectomy samples should be replicated.

Busacca et al. report a small study of in-vitro and MPM samples with mixed histological subtypes which demonstrated different expression of MicroRNAs (miRNAs) among histological subtypes, and reduced expression of specific miRNA regions and different survival among sarcomatoid subtypes [32]. MicroRNAs may act as tumour suppressors or oncogenes, and this report on only 24 specimens and eight sarcomatoid samples requires further appraisal.

A report by Fischer and co-workers described epigenetic alteration by methylation of specific promoter regions of the *RASSF1A*, *RAR β* , and *DAPK* genes that correlated with overall survival in a small sample of 43 MPM patients [33]. Hypermethylation may increase tumorigenic properties, and some demethylating agents have been found to have antitumor activity in MPM. These findings, and the emerging use of epigenetics in the diagnosis of mesothelioma, and assessment of prognosis, require further evaluation.

Table 2 Molecular and immunohistochemical reports examining survival in malignant mesothelioma

Marker of interest	Population studied		Histology	Findings	Comments	Ref.
	N					
Placenta growth factor (PIGF) expression	27 MPM patients undergoing EPP; comparators: 14 reactive mesothelium and 10 normal mesothelium	24 epithelioid; 3 biphasic	Relative overexpression of PIGF in mesothelioma samples. Inverse relationship between PIGF expression and survival ($r=-0.45$)	Selected surgical study population. PIGF is structurally and functionally similar to VEGF and this may complement similar reports on VEGF in mesothelioma	Pompeo, 2009 [18]	
Estrogen receptor- β (ER β) expression	78 MPM patients (59 male, 19 female). Mixed population (65 % had chemotherapy). 21 controls	57 epithelioid, 14 biphasic, 7 sarcomatoid	Reduced ER β staining in tumour tissue compared with controls. MLR analysis against clinical factors indicates higher ER β expression is independent predictor of improved survival	Attempts to examine why female gender previously identified as a good prognostic factor. The finding of high ER β expression in 8 (13.6 %) males and 4 (21.1 %) females needs further evaluation because of small number of females ($n=19$)	Pinton, 2009 [19]	
Steroid receptor coactivator TIF-2 and estrogen receptor- β (ER β)	89 retrospective MPM samples, 73 male, normal pleura controls.	71 epithelioid, 10 biphasic, 8 sarcomatoid	Reduced expression of TIF-2 and ER β in mesothelioma compared with normal pleura. Low TIF-2 expression correlates with worse survival. No demonstrable statistical effect of ER β expression	No difference in ER β expression between sexes in this cohort	Jennings, 2012 [20]	
Vascular endothelial growth factor (VEGF) expression	40 archived MPM histology samples (23 male). No comparator group.	30 epithelioid, 10 biphasic	Moderate to strong VEGF immunostaining in 19 (47.5 %) and 32 (80 %) overall. MLR analysis suggests presence of VEGF staining is independent prognostic factor for poor survival	Complements findings from other cancer types	Deming, 2005 [15]	
Cyclooxygenase-2 (COX-2) expression	29 MPM cases with retrospective chart review	16 epithelioid, 7 biphasic, 6 sarcomatoid	Median survival with low COX-2 expression 14 months compared with 5 months for high COX-2 expression. MLR analysis reveals statistical association with poor survival	COX-2 over-expression and worse survival demonstrated in other tumour types. May down-regulate cell-mediated immunity, promote angiogenesis and inhibit angiogenesis	Baldi, 2004 [21]	
EGFR expression	168 archived MPM sections. Mixture of primary surgical treatment and palliative samples	98 epithelioid, 37 biphasic, 33 sarcomatoid	EGFR immunostaining in 44 %; correlates with epithelioid subtype, better performance status tumour necrosis. No relationship with prognosis after MLR analysis	No relationship to COX-2 expression (as above). Large, varied study population	Edwards, 2006 [22]	
Matrix metalloproteinases 2 and 9 (MMP-2 and 9) expression	35 Prospective MPM samples; comparators: 12 inflamed pleura, 14 uninflamed pleura. Mixture of primary surgical treatment and palliative samples	Not defined	Pro and total MMP-2 levels independent poor prognostic factors in MLR. MMP-9 activity was not prognostic	MMPs are implicated in tumour growth and metastasis in other solid tumours. Mixed study population at different stages of disease limits interpretation of results	Edwards, 2003 [23]	

Table 2 (continued)

Marker of interest	Population studied		Findings	Comments	Ref.
	<i>N</i>	Histology			
p27 immunostaining	36 archived MPM samples from diagnostic biopsy specimens	19 epithelioid, 9 biphasic, 4 sarcomatoid	Positive correlation between greater p27 expression and improved survival ($r=0.33$). Suggestion that epithelioid tumours express more p27 than sarcomatoid	Complements findings in other solid tumours including recurrence and disease-free survival. Different expression between histological subtypes limited by low numbers	Beer, 2001 [24]
PTEN-PI3K pathway expression	30 retrospective MPM diagnostic samples	22 epithelioid, 7 biphasic, 1 sarcomatoid	Marked over-expression of most proteins in PI3K pathway. Combination of high pS6 and p4E-BP1 proteins correlates with worse survival. No significance of any downstream proteins in MLR analysis for survival	PTEN-PI3K implicated with AKT/mTOR activity. To date, use of an mTOR inhibitor in combination with cisplatin has not proved clinically effective	Cedres, 2012 [25]
Epithelial–mesenchymal transition (EMT)	352 mostly untreated MPM samples	113 epithelioid, 194 biphasic, 45 sarcomatoid	MLR analysis reveals low cytoplasmic periostin and high expression of PTEN are independently associated with improved overall survival. Epithelioid subtype associated with high membranous EGFR and integrin $\beta 1$ expression, and nuclear p27. Sarcomatoid subtype associated with high cytoplasmic tumoural and stromal periostin expression	A large sample with good generalizability. PTEN can abolish PI3K/Akt activation, which may be utilised for EMT	Schramm, 2010 [26]
Aquaporin 1 (AQPI) expression	80 consecutive radical surgery patients and 56 conservatively-managed patients	Surgery group: 61 epithelioid, 19 biphasic; Conservative group: 23 epithelioid, 14 biphasic, 19 sarcomatoid	For both cohorts better survival is associated with expression of AQPI by ≥ 50 % tumour cells. MLR analysis reveals expression of AQPI is independent predictor of improved survival	Strong data as findings further validated on conservative management cohort	Kao, 2012 [27]

MPM malignant pleural mesothelioma; MLR multivariate logistic regression;

The reports summarised above provide interesting information, with the potential to advance our understanding of the genetics of tumour biology and prognosis in MPM, but none yet delivers markers or quantitative data that have direct prognostic relevance for clinicians.

Biomarkers

The use of a specific biomarker, or combination of readily-available quantitative laboratory variables, to aid diagnosis of MPM and to determine prognosis is highly desirable, because of the ease of sampling (e.g. blood or body cavity fluid) and the potential to assess disease expression and response to treatment. We discuss several relevant biomarkers below.

Vascular Endothelial Growth Factor (VEGF)

VEGF is a family of proteins which have been shown to be important in angiogenesis and vascular permeability. Tissue VEGF has been shown to correlate with microvessel density, and mesothelioma patients expressing high levels of this cytokine within the tumour have been shown to have a poor prognosis [15]. VEGF is an autocrine growth factor for MPM, and the predictive value of circulating VEGF was the subject of early study in mesothelioma.

Serum VEGF levels were measured in 51 patients with MPM and in 42 individuals with benign asbestos-related diseases. Patients with MPM had higher levels of VEGF than those with benign disease. Within the MPM study population, high levels of serum VEGF (>460 pg mL⁻¹) were associated with poor survival [34]. In another study, pleural fluid VEGF was measured in 46 MPM patients and in 45 individuals with other causes of pleural effusion. Patients with MPM had higher levels than other pleural effusion patients, and levels >2000 pg mL⁻¹ were associated with poorer survival in the MPM subgroup [35].

Neutrophil-to-lymphocyte ratio (NLR)

Inflammation strongly affects the development and progression of cancers, and release of proinflammatory cytokines in patients with MPM can produce systemic inflammatory symptoms, for example fever, sweating, and weight loss. The neutrophil-to-lymphocyte ratio (NLR) is believed to be a marker of systemic inflammation and can be simply calculated from a full blood count that includes the differential white cell count. A high ratio suggests greater systemic inflammation.

Kao et al. retrospectively studied 173 MPM patients undergoing systemic therapy. Forty-two percent had

elevated NLR at baseline. Epithelioid histology (HR 2.0, 95 % CI 1.3–2.9) and NLR of five or below (HR 2.7, 95 % CI 1.8–3.9) were independently predictive of survival [36]. In another study by the same group, 85 patients with MPM undergoing extra-pleural pneumonectomy (EPP) had baseline variable performed; $NLR \geq 3$ was associated with poor prognosis [37•].

Mesothelin and osteopontin

Mesothelin is a cell-surface glycoprotein expressed on normal mesothelial cells which is involved in cell adhesion. In MPM mesothelin is often over-expressed and may be released from the cell surface in the form of soluble mesothelin (also known as soluble mesothelin-related peptides (SMRPs)). SMRPs can be detected in the blood and pleural fluid by use of commercially available ELISA.

Creaney et al. prospectively studied 97 patients with MPM, measuring baseline and serial serum mesothelin levels. Baseline mesothelin levels >5 nmol L⁻¹ were a significant negative prognostic indicator (HR 2.25, 95 % CI 1.2–4.21) and correlated with tumour stage and volume. For 55 patients receiving chemotherapy, changes in mesothelin correlated with radiological response. Median survival for patients with a decrease in mesothelin after chemotherapy was substantially longer than for those with increased mesothelin (19 months vs. 5 months $p < 0.001$) [38•].

Preliminary data indicate that changes in serum mesothelin level over time could be used to monitor disease progression and response to treatment in mesothelioma. A rising mesothelin level of >10 % despite treatment was associated with a worse outcome than that for patients for whom the serum mesothelin level remained stable [39•].

One of the main limitations of mesothelin is the number of MPM patients with serum mesothelin levels below the limits of detection. This is most common among patients with sarcomatoid mesothelioma.

Osteopontin is another glycoprotein; it modulates cell-matrix interactions and is over-expressed by mesothelioma. In an observational study from Belgium, high osteopontin levels were found to be an independent negative predictor of survival with mesothelioma, although less closely associated with treatment response than serum mesothelin levels were [40]. Grigoriu et al. also found that serum mesothelin >3.5 nmol L⁻¹ and osteopontin >350 ng mL⁻¹ were both associated with poorer prognosis for patients with mesothelioma [41].

Imaging modalities: CT and PET

Accurate evaluation of response to treatment in MPM can be difficult to measure on CT because mesothelioma usually

presents as a multifocal pleural abnormality, rather than a single tumour mass. It is usually crescent-shaped, rather than spherical, and it is, therefore, difficult to measure disease progression using the standard RESIST criteria. Attempts to address these issues by use of modified RESIST criteria have been made by Nowak and colleagues [42]; however, the pleural cavity is often fibrotic (particularly in sarcomatoid cases), and tumour death within fibrotic pleura is difficult to differentiate. This has led researchers to look at integrated 2-deoxy-2-(F-18)fluoro-D-glucose-positron emission tomography/computed tomography (^{18}F -FDG PET/CT), both as a baseline prognostic tool and as a potential metabolic marker of response to chemotherapy.

Nowak et al. studied 89 patients with proved mesothelioma, 28 of whom had had a talc pleurodesis before enrolment. All had ^{18}F -FDG PET/CT at baseline, and PET variables studied included total glycolytic volume (TGV), a composite of tumour volume and glycolytic activity. By use of univariate analysis, significant baseline prognostic factors were: total glycolytic volume, sarcomatoid histology, weight loss, CT stage, and EORTC prognostic score. In multivariate analysis, only histology and TGV remained predictive of survival. In non-sarcomatoid disease TGV was more predictive of survival than CT staging. Pleurodesis induces an intense inflammatory response in the pleura, which may result in false positive uptake on FDG-PET imaging. However, the TGV value obtained remained a strong predictor of survival in both the pleurodesis and non-pleurodesis group [43].

In another study of 46 patients with biopsy-proven mesothelioma, all had ^{18}F -FDG PET/CT at baseline and the SUV_{max} was calculated. ^{18}F -FDG PET/CT was better than CT at detecting metastatic disease (9/46, 20 %) [44]. Most of these (8/9) had not previously been detected by CT imaging. Evaluation of progression-free survival by Kaplan–Meier analysis did not correlate with SUV_{max} . In this study, non-epithelioid histology and presence of metastatic disease were the only markers of poor prognosis.

In a further small study, by the same group, of 13 patients due to undergo extrapleural pneumonectomy (EPP) or palliative chemotherapy, ^{18}F -FDG PET/CT was used to measure SUV_{max} , SUV_{avg} , metabolic tumour volume (MTV), and total lesion glycolysis (TLG). TLG was obtained by multiplying MTV by SUV_{avg} . High TLG ($>1,250$) or MTV (>250) at baseline were associated with poor prognosis and a short time to disease progression (TTP). No such association was observed for SUV_{max} [45].

For patients with mesothelioma the volume-based variable TGV can be easily measured by semiquantitative ^{18}F -FDG PET/CT. It can be used to identify early responders to chemotherapy; Francis et al. showed that a 30 % fall in TGV after one cycle of chemotherapy was strongly predictive of survival [42]. This must be confirmed for other cohorts, but

suggests that ^{18}F -FDG PET is likely to be an important research tool when evaluating response to novel treatment.

In summary, ^{18}F -FDG PET/CT seems to provide useful data for patients with mesothelioma. It is better at detecting metastatic disease and, although SUV_{max} does not correlate with survival, TGV does. In addition, falling TGV levels on repeat imaging after chemotherapy are indicative of “metabolic responders”, who seem to have a better prognosis than those with rising TGV levels.

Conclusions

Population-based studies consistently indicate that increased age, male gender, and non-epithelioid histology are poor prognostic factors. There is interest in the utility of mitotic activity and nuclear atypia for assessment of tumours, although this work needs further validation. There are increasing immunohistochemical and genetic profiling reports of MPM that, at present, do not provide clear, validated prognostic data, although these modalities are likely to provide the most significant advances in our future understanding of tumour biology and prognosis. Biomarkers from serum or pleural fluid do provide some prognostic information, although with limitations, because of different expression by different histological subtypes. Their use in monitoring response to treatment is the subject of a large multi-centre UK-based trial (the SWAMP trial; UKCRN ID 8458), which is due to report its findings shortly. Modern metabolic imaging modalities, for example PET/CT are able to provide prognostic information, particularly regarding response to treatment.

At present much of the tissue biology literature is based on predominantly surgical specimens; this is a selected population and not necessarily representative of the wider population. However, anecdote suggests that long survivors will frequently have similar characteristics to surgical populations, so studies of, and reports on, the latter are still valid. It is possible that better understanding of the phenotype of long survivors may, in turn, lead to better understanding of disease prognosis in mesothelioma in general and this should continue to be a research priority.

Overall, the prognosis for mesothelioma is poor. There is, however, wide variation in survival, response to treatment, and disease progression among individuals, which may in part result from the known biological heterogeneity of this cancer. As immunohistochemical and genetic advances continue it may become more apparent why some characteristics of the disease or patient (for example histology or gender) result in a different prognosis. There remains a need for identification of validated, clinically relevant prognostic factors that can be generalised to the wider population with mesothelioma.

Conflict of Interest Fraser J.H. Brims declares that he has no conflict of interest.

Nick A. Maskell declares that he has no conflict of interest.

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