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# Fibulin-3 levels in malignant pleural mesothelioma are associated with prognosis but not diagnosis

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**Background:** Fibulin-3 (FBLN3) was recently presented as a promising novel biomarker for malignant pleural mesothelioma (MPM), warranting independent validation studies.

**Methods:** ELISA was used to measure cellular and secreted FBLN3 in cell lines, in plasma of xenograft tumour-bearing mice, in plasma from two independent series of MPM and non-MPM patients and in pleural fluid from a third series. Diagnostic and prognostic potential of FBLN3 was assessed by receiver operating characteristics curve analysis and Kaplan–Meier method, respectively.

**Results:** FBLN3 was expressed in all MPM and benign mesothelial cell lines tested, and a correlation was observed between cellular protein expression and secreted levels. Human FBLN3 was detectable in plasma of tumour-bearing mice, suggesting that MPM cells contribute to levels of circulating FBLN3. Plasma FBLN3 was significantly elevated in MPM patients from the Sydney cohort, but not the Vienna cohort, but the diagnostic accuracy was low (63%, (95% CI: 50.1–76.4) and 56% (95% CI: 41.5–71.0), respectively). Although FBLN3 levels in pleural effusions were not significantly different between cases and controls, FBLN3 levels in pleural effusion fluid were found to be independently associated with prognosis (hazard ratio of 9.92 (95% CI: 2.14–45.93)).

**Conclusions:** These data confirm the potential prognostic value of pleural effusion FBLN3, but question the diagnostic value of this protein in MPM patients.

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Fibulin-3 (FBLN3), encoded by the *EGF-containing fibulin-like extracellular matrix protein 1 (EFEMP1)* gene, is a member of the fibulin family of secreted extracellular glycoproteins that are characterised by a tandem repeat of EGF-like domains and a unique C-terminal fibulin-like module (Timpl *et al*, 2003; Zhang and Marmorstein, 2010; Zhou, 2013). The fibulin family forms part of the extracellular matrix (ECM) where they contribute to the stabilisation of supramolecular structures such as elastic fibres and basement membranes through interaction with various other components of the ECM (Argraes *et al*, 2003; Timpl *et al*, 2003). Expression of FBLN3 in condensing mesenchyme during development, which gives rise to bone and cartilage structures, implies an important role of this family member in skeletal development (Zhang and Marmorstein, 2010). In adult tissue FBLN3 is widely expressed in blood vessel walls, and in basement membranes of epithelial and endothelial cells (Giltay *et al*, 1999; Timpl *et al*, 2003; Zhang and Marmorstein, 2010). FBLN3 has been shown to stimulate expression of the tissue inhibitors of matrix metalloproteinases (TIMP)-1 and -3 (Klenotic *et al*, 2004; Kim *et al*, 2012), while inhibiting the expression of the matrix metalloproteinases MMP-2, -3, -7 and -9 (Albig *et al*, 2006; Kim *et al*, 2012), thus being directly involved in the regulation of tissue remodelling. In addition, FBLN3 has been reported to be involved in cell growth and tumour angiogenesis (Gallagher *et al*, 2005; Albig *et al*, 2006).

Fibulins are a relatively newly characterised group of ECM proteins and a link between FBLN3 and cancer has only been established over the past decade. Interestingly, the expression pattern of FBLN3 differs between tumour types, with up- or downregulation found depending on the cancer investigated (Yue *et al*, 2007; Camaj *et al*, 2009; Hu *et al*, 2009; Sadr-Nabavi *et al*, 2009; Seeliger *et al*, 2009; En-lin *et al*, 2010; Hwang *et al*, 2010; Nomoto *et al*, 2010; Wang *et al*, 2010; Hu *et al*, 2011, 2012, 2013; Kim *et al*, 2011; Song *et al*, 2011; Tong *et al*, 2011; Kim *et al*, 2012; Pass *et al*, 2012; Luo *et al*, 2013; Chen *et al*, 2014). FBLN3 therefore appears to have the potential to act as either tumour suppressor or oncogene depending on the cellular context.

More recently FBLN3 has been identified as a potential diagnostic and prognostic marker for malignant pleural mesothelioma (MPM) (Pass *et al*, 2012). Increased abundance of the FBLN3 protein was found in plasma and pleural effusion fluid of MPM patients, with high levels in effusions negatively correlated with survival. The same study also reported elevated FBLN3 expression in MPM primary tumours (Pass *et al*, 2012); however, a direct link between tumour FBLN3 expression and the secreted form of FBLN3 is yet to be made.

Towards this end, we investigated the expression and secretion of FBLN3 in MPM and benign mesothelial cell lines *in vitro* and *in vivo*. Our data confirm an upregulation of FBLN3 protein in MPM cell lines, which is mirrored by an increased secretion of the protein. In addition, we were able to confirm that levels of FBLN3 in pleural effusion may have prognostic value. However, a diagnostic role for FBLN3 levels in plasma was not supported by our data.

## MATERIALS AND METHODS

**Cell lines and protein lysis.** Human MPM cell lines (H28, H226, H2452 and MSTO) and the immortalised normal cell line MeT-5A were obtained from the American Type Culture Collection (Manassas, VA, USA). The additional MPM lines MM05 (Relan *et al*, 2013), VMC23 (Kryeziu *et al*, 2013) and SPC111 (Schmitter *et al*, 1992) and the normal mesothelial line LP9 (Wu *et al*, 1982) were described previously. Cell lines were grown in the corresponding medium (RPMI-1640 for all MPM lines, DMEM for MeT-5A, M199 supplemented with 3.3 nM EGF, 400 nM

hydrocortisone and 1% insulin-transferrin-selenium for LP9) supplemented with 10% fetal bovine serum (FBS) at 37 °C, 5% CO<sub>2</sub> and 95% humidity. All media and supplements were purchased from Life Technologies (Carlsbad, CA, USA) or Sigma Aldrich (St Louis, MO, USA). All cell lines were authenticated in September (H28, H2452, SPC111 and VMC23) or October (H226, H2052, MSTO, MM05, MeT-5A and LP9) 2014 at the Australian Genome Research Facility Ltd. (St Lucia, QLD, Australia) by short tandem repeat profiling using the GenePrint 10 System (Promega, Madison, WI, USA).

To assess secretion of FBLN3 by MPM cells *in vitro*, cells were first grown to 80% confluence in T25 flasks in medium containing FBS, then medium was replaced with 2 ml serum-free medium. Cell-conditioned medium was collected after 24 h, cell debris removed by 5 min centrifugation at 1200 r.p.m., and supernatants stored at -80 °C. Protein lysates were prepared from the same cells by lysis in 300 µl RIPA buffer as described previously (Linton *et al*, 2014). Protein concentration was determined using the Pierce BCA Protein Assay Kit (Thermo Scientific, Waltham, MA, USA) according to the manufacturer's instructions.

**Mouse plasma and serum samples.** Athymic (*nu/nu*) mice were purchased from the Animal Resources Centre (Perth, WA, Australia) and injected subcutaneously in the left flank with  $5 \times 10^6$  H226 or  $3 \times 10^6$  MSTO cells in 100 µl serum-free medium/growth factor-reduced matrigel mix (1:1, matrigel from BD Biosciences, Franklin Lakes, NJ, USA). Once tumours reached around 300 mm<sup>3</sup>, mice were killed and blood was collected via cardiac puncture into either empty or EDTA-containing tubes. Serum samples were allowed to clot for 60 min at room temperature before being further processed as for plasma samples (centrifugation at 2500 r.p.m. for 20 min) and both were stored at -80 °C. Samples were collected as part of animal studies approved by the Sydney Local Health District Animal Welfare Committee.

**Patient samples.** All samples were collected from patients who had provided written informed consent and the study was approved by the Human Research Ethics Committees at Concord Repatriation General Hospital (CH62/6/2009/078) and Royal Prince Alfred Hospital (X09-0322) in Sydney, the Ethical Committee of the Medical University of Vienna (#904/2009) or the Southern Adelaide Clinical Human Research Ethics Committee (#381.09).

In Sydney and Vienna blood was collected by venous puncture from (suspected) MPM patients 1 day before diagnostic procedure (video-assisted thoracoscopic surgery/biopsy) or radical surgery (extrapleural pneumonectomy) or from patients undergoing cardiac or aortic surgery for severe coronary artery (CAD) or aortic disease at participating centres. In Vienna additional control samples were collected from lung cancer patients. Blood was collected into 10 ml K<sub>3</sub>EDTA Vacutainers (BD Biosciences) which, within 30 min of blood collection, were centrifuged for 20 min at 2500 r.p.m. Plasma supernatant was stored in aliquots at -80 °C. Pleural effusion samples were obtained from SA Pathology, Flinders Medical Centre and Institute of Medical and Veterinary Science SA. All pleural effusions were screened for diagnostic suitability by accessing pathology reports, with cytological diagnosis confirmed from subsequent radiological and/or surgical reports. MPM samples were obtained from consecutive patients presenting at Flinders Medical Centre between August 2011 and June 2014, while control samples were identified from the pathology database and collected until age and gender-matched samples from 30 patients in each group were identified. The median time from sample collection to confirmation of diagnosis was 2 days. Samples were spun at 1250 r.p.m. for 10 min, and the resulting supernatant was stored immediately at -80 °C. A summary of basic patient demographics is provided in Table 1.

**Table 1. Patient demographics**

	Sydney cohort		Vienna cohort		Pleural effusion cohort	
	MPM (N = 37)	Non-MPM (N = 32)	MPM (N = 47)	Non-MPM (N = 24)	MPM (N = 30)	Non-MPM (N = 60)
Median age (range)	71.4 (40–83)	66.3 (49–80)	64 (33–81)	66.5 (56–82)	76 (55–94)	73.5 (49–96)
<b>Gender</b>						
Male	32 (86.5%)	27 (84.4%)	35 (74.5%)	19 (79.2%)	26 (86.7%)	51 (85%)
Female	5 (13.5%)	5 (15.6%)	12 (25.5%)	5 (20.8%)	4 (13.3%)	9 (15%)
<b>Diagnosis</b>						
MPM	37 (100%)		47 (100%)		30 (100%)	
Pleural Plaques/Pleuritis		5 (15.6%)		10 (41.7%)		30 (50%)
<b>Adenocarcinoma</b>				4 (16.7%)		27 (45%)
Lung						1 (1.6%)
Breast						1 (1.6%)
Endometrial						1 (1.6%)
Colon						1 (1.6%)
Other lung cancer				3 (12.5%)		
CAD		27 (84.4%)		7 (29.2%)		
<b>MPM histotype</b>						
Epithelioid	26 (70.3%)		35 (74.5%)		27 (90%)	
Biphasic	8 (21.6%)		7 (14.9%)		2 (6.7%)	
Sarcomatoid	2 (5.4%)		4 (8.5%)		1 (3.3%)	
Desmoplastic	1 (2.7%)		1 (2.1%)		0	

Abbreviations: CAD = coronary artery disease; MPM = malignant pleural mesothelioma.

**FBLN3 ELISA.** FBLN3 levels were determined using the human FBLN3 ELISA (#SEF422Hu, USCN Life Science Inc., Wuhan, China) according to the manufacturer's instructions at the centres at which the respective samples were collected. Cell lysates were diluted 1:2, cell-conditioned medium 1:4, plasma samples were either diluted 1:3 (Sydney cohort and mouse plasma) or used undiluted (Vienna cohort), and pleural effusion samples were diluted 1:10 in phosphate-buffered saline. Pleural effusion samples that showed concentrations higher than the highest standard were diluted further and measured again. All samples were measured in duplicate. FBLN3 levels were calculated and expressed as either  $\text{ng mg}^{-1}$  total protein (cell lysates and cell-conditioned medium) or  $\text{ng ml}^{-1}$  (plasma and pleural effusion). Samples with readings below those of the lowest standard were reported as  $1.56 \text{ ng ml}^{-1}$ .

**Statistical analysis.** FBLN3 levels are reported as mean  $\pm$  s.d., and significant differences in FBLN3 levels were determined using independent samples *t*-test. Receiver operating characteristics curve analysis was used to determine the accuracy of FBLN3 to predict correct diagnosis. Survival was calculated from date of diagnosis until date on which analysis was performed, with patients still alive at this time point being censored. The Kaplan-Meier log rank method was used to assess association between FBLN3 and survival, and if significant in univariate analysis, multivariate cox-regression was performed to investigate the value independent prognostic factor.

## RESULTS

**MPM cells express and secrete FBLN3 *in vitro*.** FBLN3 expression was detected in all benign mesothelial (expression range:  $4.99\text{--}10.51 \text{ ng mg}^{-1}$ ) and MPM lines (expression range:  $8.85\text{--}18.78 \text{ ng mg}^{-1}$ , mean expression  $13.89 \pm 3.94 \text{ ng mg}^{-1}$ ). There was no statistically significant difference between levels in the two groups ( $P=0.093$ ) (Figure 1A). Among MPM cell lines, the average expression was higher in cells derived from tumours of epithelioid histology ( $16.18 \pm 3.46 \text{ ng mg}^{-1}$ ) than in those from biphasic (mixed) histology ( $10.64 \pm 2.04 \text{ ng mg}^{-1}$ ). Although not reaching statistical significance ( $P=0.066$ ), this trend suggests a

possible correlation between histological subtype and expression of FBLN3. Assessment of secreted FBLN3 levels in cell-conditioned medium (Figure 1B) revealed a correlation between cellular FBLN3 expression and secretion of the protein (Spearman  $r=0.78$ ,  $P=0.017$ , Figure 1C), with a non-statistically significant trend towards higher secretion observed in cells derived from epithelioid tumours ( $P=0.26$ ).

**FBLN3 is secreted by MPM cells *in vivo*.** To confirm MPM cells as the origin of circulating FBLN3, protein levels were measured in serum and plasma collected from H226 and MSTO xenograft-bearing mice. Although FBLN3 was not detectable in serum samples, 3/3 plasma samples from H226-xenografts and 2/3 plasma samples from MSTO-xenografts were positive for human FBLN3 (Figure 1D). Although from a small number of samples, the *in vivo* data are consistent with the *in vitro* observation that MPM cells actively secrete FBLN3, and suggest that MPM cells contribute to the FBLN3 levels found in the blood. The lack of detection of FBLN3 in serum samples is in line with previous findings of low abundance of FBLN3 in serum, which may be attributed to thrombin-mediated cleavage of FBLN3 (Pass *et al*, 2012).

**Levels of FBLN3 in MPM patient plasma have little diagnostic value.** Abundance of FBLN3 was assessed in two independent cohorts of MPM patients and controls. The mean levels of FBLN3 in plasma from MPM patients were  $16.10 \pm 1.87 \text{ ng ml}^{-1}$  (Sydney) and  $11.51 \pm 1.73 \text{ ng ml}^{-1}$  (Vienna), and were lower than those previously reported (Pass *et al*, 2012; Creaney *et al*, 2014; Hooper *et al*, 2015). There was no significant difference in the FBLN3 levels observed in the two cohorts ( $P=0.077$  for MPM and  $P=0.77$  for non-MPM samples). Analysis of the Sydney cohort showed a modest but statistically significant increase in circulating FBLN3 in MPM patients compared with patients with pleural plaques or CAD ( $16.10 \pm 1.87$  vs  $10.92 \pm 1.54 \text{ ng ml}^{-1}$ ,  $P=0.039$ , Figure 2A), whereas analysis of the Vienna cohort did not ( $11.51 \pm 1.73$  vs  $11.97 \pm 3.56 \text{ ng ml}^{-1}$ ,  $P=0.897$ ; Figure 2B). Receiver operating characteristics curve analyses showed an overall accuracy of 63.2% (95% CI: 50.1–76.4%,  $P=0.06$ , Figure 2C) for correct classification as MPM for the Sydney cohort and 56.2% (95% CI: 41.5–71.0%,  $P=0.39$ , Figure 2D) for the Vienna cohort. At the lower cut-off of  $29 \text{ ng ml}^{-1}$  (Pass *et al*, 2012), FBLN3 in the Sydney and Vienna

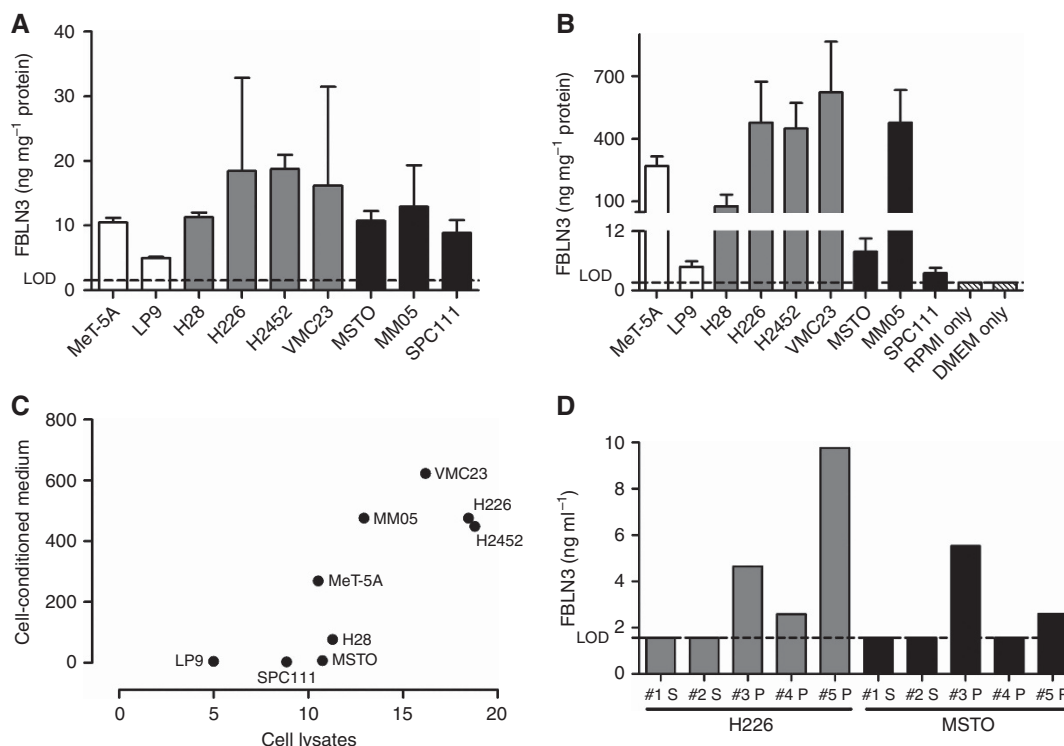


Figure 1. MPM cell lines secrete FBLN3. Cellular (A) and secreted (B) levels of FBLN3 in MPM cell lines (grey bars = epithelioid; black bars = biphasic) and benign mesothelial cells (white bars). (C) Relationship between the amount of secreted FBLN3 and cellular FBLN3 levels. (D) FBLN3 levels in the plasma (P) or serum (S) of mice bearing human MPM xenografts of epithelioid (grey bars) or biphasic (black bars) origin. Lower limit of detection (LOD) of the ELISA is indicated by the dotted line, medium only (RPMI or DMEM) controls are shown in patterned white bars.

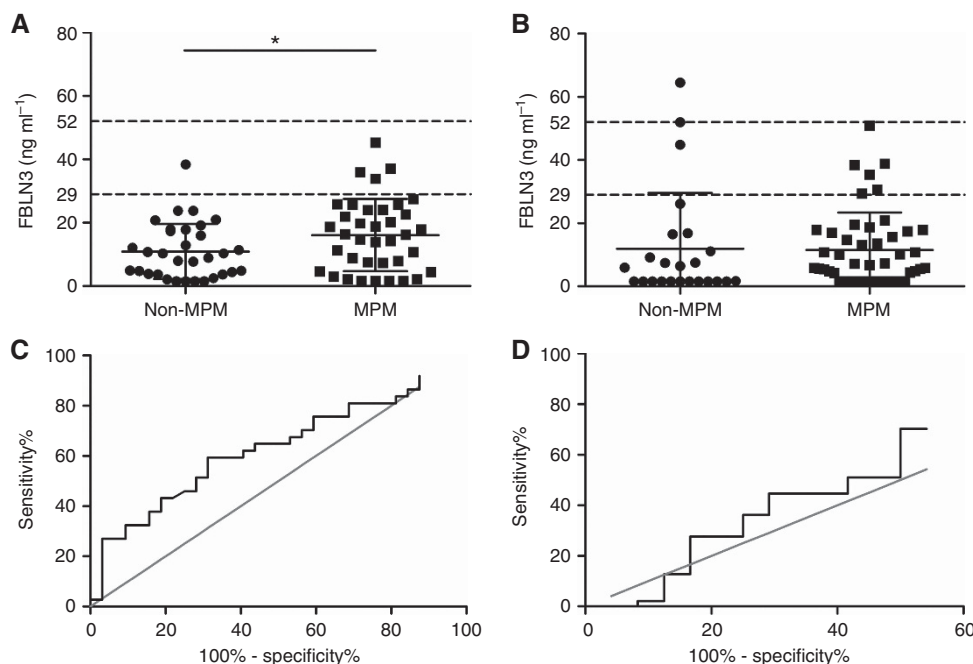
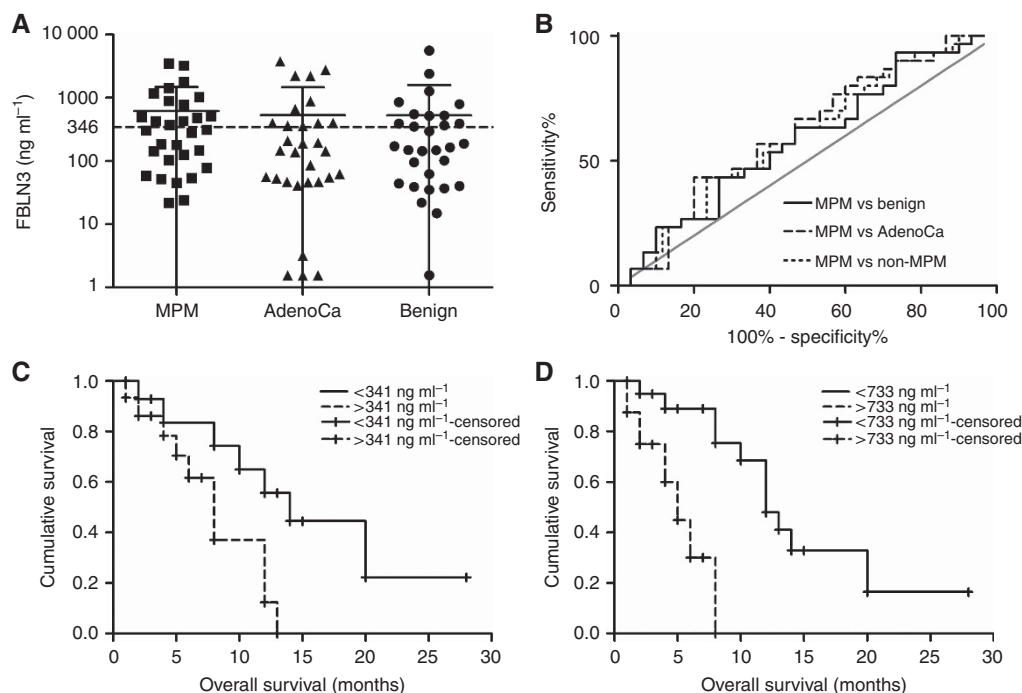


Figure 2. FBLN3 in patient plasma. Plasma FBLN3 protein levels in the Sydney (A) and Vienna cohort (B). Mean levels in both cohorts were below those previously reported. Mean  $\pm$  s.d. are represented by the lines in the scatter plots, and the cut-offs applied in the original study (Pass *et al*, 2012) are indicated by dotted lines. The diagnostic accuracy of plasma FBLN3 was low in both investigated cohorts: (C) Sydney cohort AUC = 0.63 (95% CI: 0.50–0.76); and (D) Vienna cohort AUC = 0.56 (95% CI: 0.41–0.71).

series showed a sensitivity of 13.5% and a specificity of 96.9%, and a sensitivity of 12.7% and a specificity of 87.5%, respectively. Differences in circulating FBLN3 levels between patients with different histological subtypes or associations with survival were not observed (data not shown).

**High FBLN3 levels in pleural effusion fluid are associated with poor prognosis.** Mean FBLN3 levels in MPM effusions were  $628.9 \pm 853.8$  ng ml<sup>-1</sup> and while comparable to those previously published ( $636$  ng ml<sup>-1</sup>) (Pass *et al*, 2012; Creaney *et al*, 2014), significant differences between effusions from MPM patients and





**Figure 3.** Pleural effusion FBLN3. **(A)** FBLN3 was measured in the pleural effusion in MPM and non-MPM conditions. The lines in the scatter plot represent the mean  $\pm$  s.d. **(B)** Receiver operating characteristics analysis of FBLN3 levels in the different physiological conditions. **(C, D)** Kaplan–Meier analysis of FBLN3 in MPM patients using the median FBLN3 level **(C)** or the cut-off applied in the original study (Pass *et al*, 2012) **(D)**.

patients with other conditions were not observed (Figure 3A). Receiver operating characteristics curve analyses showed low accuracy for correct classification of MPM patients in comparisons with all other groups (Figure 3B). Survival analyses in the MPM subset did not show any correlation between clinical prognostic factors (histological subtype, age and gender), but this may be attributed to the relatively small numbers in the various subgroups (Table 1). Low levels of FBLN3 were, however, significantly associated with prolonged survival, and this was independent of whether the median for this group (Figure 3C) or the previously published cut-off (Pass *et al*, 2012) (Figure 3D) was used. In multivariate analyses including histological subtype, age and gender as covariates, FBLN3 remained significant at both cut-offs with hazard ratios (HRs) of 4.03 (95% CI: 1.215–13.36) and 9.92 (95% CI: 2.14–45.93), respectively.

## DISCUSSION

A recent study describing elevated FBLN3 levels in MPM patients reported that FBLN3 is overexpressed in tumour tissue, with higher levels also present in pleural effusion fluid and plasma (Pass *et al*, 2012). These results suggest a possible oncogenic role for FBLN3 in MPM. Our studies with MPM cell lines *in vitro* and xenografted tumours *in vivo* support these previous findings in MPM tissue, and suggest that MPM cells express and secrete FBLN3, and are a source of FBLN3 in patient plasma samples. Interestingly, while FBLN3 was also found to be elevated in pancreatic (Camaj *et al*, 2009; Seeliger *et al*, 2009), cervical (En-lin *et al*, 2010; Song *et al*, 2011) and brain cancer (Hu *et al*, 2009, 2011, 2012), in the majority of cancer types (Yue *et al*, 2007; Sadr-Nabavi *et al*, 2009; Hwang *et al*, 2010; Nomoto *et al*, 2010; Wang *et al*, 2010; Kim *et al*, 2011; Tong *et al*, 2011; Kim *et al*, 2012; Luo *et al*, 2013; Chen *et al*, 2014) the protein is downregulated. These opposing findings are further complicated in malignant glioma where FBLN3 was found to be upregulated compared with normal tissue, but at the same time

positively correlated with prolonged survival (Hu *et al*, 2009, 2011, 2012). Together these data suggest FBLN3 may have varying context- and tissue-dependent roles. This is further highlighted by the involvement of FBLN3 in the regulation of Akt signalling in different types of cancer. Although epigenetic silencing of FBLN3 in nasopharyngeal carcinoma (NPC) leads to activation of the Akt signalling pathway (Hwang *et al*, 2010), in pancreatic cancer overexpression of FBLN3 also results in Akt activation (Camaj *et al*, 2009). These two seemingly opposing effects are thought to be brought about by two different mechanisms. On the one hand, FBLN3 protein has been suggested to interfere with phosphorylation and therefore activation of Akt, meaning that low FBLN3 results in an increase in Akt signalling leading to increased migration and invasion of NPC cells (Hwang *et al*, 2010). On the other hand, in pancreatic cancer, FBLN3 was shown to indirectly stimulate Akt phosphorylation through binding to and activating signalling through the EGF receptor (EGFR) (Camaj *et al*, 2009). In the context of MPM one can speculate that similar to the scenario in pancreatic cancer the elevated expression of FBLN3 together with an overexpression of EGFR (observed in 40–60% of MPM cases (Destro *et al*, 2006; Edwards *et al*, 2006; Gaafar *et al*, 2010; Rena *et al*, 2011)) may act in concert to lead to the increased activation of Akt signalling frequently observed in MPM (Altomare *et al*, 2005). Further investigation is needed to determine whether FBLN3 might therefore act as an oncogene contributing to the invasive nature of MPM.

The initial report describing FBLN3 overexpression in MPM suggested that this can serve as a reliable diagnostic and prognostic biomarker of the disease (Pass *et al*, 2012). However, these promising findings have yet to be fully validated. The original study reported elevated levels of cell-free FBLN3 in both plasma and pleural effusion fluid which were able to separate, with very high sensitivity and specificity, MPM patients from healthy individuals and patients with effusions due to other malignancies (Pass *et al*, 2012). Three validation studies (Corradi *et al*, 2013; Agha *et al*, 2014; Creaney *et al*, 2014) confirmed that

plasma levels of FBLN3 in MPM were elevated, but sensitivity and specificity of the assay were either notably lower or cut-off levels used to determine these were different to those originally applied, making direct comparisons difficult. In line with these data, we found plasma FBLN3 to be significantly elevated in one of our two plasma series. Although overall levels of FBLN3 were much lower than those reported in the first (Pass *et al*, 2012) and one subsequent (Agha *et al*, 2014) report, they were more comparable to those reported in the two other validation studies (Corradi *et al*, 2013; Creaney *et al*, 2014). Average FBLN3 levels were lower than the diagnostic cut-off used by Pass *et al*. In addition, sensitivity and specificity of the assay were lower than those in the original study (Pass *et al*, 2012) and did not reach the cut-off of 80% for both sensitivity and specificity which would classify this protein as a good diagnostic marker according to International Mesothelioma Interest Group guidelines (Husain *et al*, 2013). Most recently, the South West Area Mesothelioma and Pemetrexed Trial also reported on FBLN3 in plasma from MPM patients (Hooper *et al*, 2015). Although this trial was not designed to assess the diagnostic potential of any of the investigated biomarkers, the investigators also found FBLN3 levels to be considerably lower than reported by Pass *et al* (2012), with only 8% of patients having levels higher than the proposed diagnostic cut-off of  $56 \text{ ng ml}^{-1}$  (Hooper *et al*, 2015).

Creaney *et al* (2014) also investigated FBLN3 in pleural effusion samples. Although the authors did not find significant differences in FBLN3 levels between patient groups, high FBLN3 pleural effusion levels were independently associated with shorter survival (HR 2.05). Similarly we did not observe significant differences between the levels of FBLN3 in the various samples of malignant pleural effusions, but confirmed that high effusion FBLN3 levels were independently associated with short survival. Thus FBLN3 in pleural fluid may represent a useful prognostic marker in MPM; however, the initial findings supporting FBLN3 as a reliable diagnostic marker could not be reproduced. The presence of elevated levels of FBLN3 in pleural effusion of MPM patients together with experimental data suggesting that FBLN3 is actively secreted by FBLN3-over-expressing MPM tumour cells clearly asks for additional experimental studies.

The current study has some limitations: FBLN3 assessment in plasma and serum from xenograft-bearing mice, while limited by a relatively small sample size, nevertheless showed that MPM tumour cells actively secrete FBLN3 *in vivo*. The sample sizes of the human material (patient cohorts) were also relatively small; however, the results obtained in the two independent series of plasma samples obtained at different institutions were comparable. In addition, given its retrospective nature, the study is bound by the inherent limitations of such studies. The use of patients undergoing cardiac surgery as controls could also be seen as a weakness of the study, since patients with proven asbestos exposure would be considered the more appropriate control. However, in particular for Australia it is known that due to the large amounts of asbestos in the natural environment and the heavy use of asbestos in the past resulting in large amounts also being present in the built environment the majority of citizens have been exposed to at least low levels of asbestos (knowingly or unknowingly) in a non-occupational setting (LaDou *et al*, 2010; Olsen *et al*, 2011). We therefore assume the phenomenon of (often unknown) low level asbestos exposure also to be reflected in our control series, and this is particularly true for the cardiac patients in the Sydney series. Despite these limitations our data further confirm the prognostic value of pleural effusion FBLN3, but questions the diagnostic value of this protein in plasma of MPM patients. The results of the present study contribute to a growing literature on the value of FBLN3 as a biomarker in MPM, and underline the need for prospective validation.

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## CONFLICT OF INTEREST

The authors declare no conflict of interest.

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