

# **Defining a New Prognostic index for Stage I Non-seminomatous Germ Cell Tumors using CXCL12 Expression and Proportion of Embryonal Carcinoma**

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**Translational Relevance:** Patients diagnosed with stage I non-seminomatous germ cell tumors face a choice between surveillance with treatment at relapse or up-front adjuvant therapy. Whilst adjuvant therapy is effective, it may be unnecessary and long-term effects of chemotherapy are increasingly recognized. Histological evidence of vascular invasion is currently used to target patients for adjuvant therapy, but improved markers for risk of relapse are required. Embryologically, primordial germ cells use CXCR12/CXCR4 and KITLG/KIT signaling to migrate to the developing gonads. Previously we showed that CXCL12 stimulates migration of germ cell tumor cells in a CXCR4-dependent manner and that tumor cell expression of CXCL12 was associated with reduced risk of metastatic relapse. Here we validate this finding in a large series of samples from patients that underwent surveillance within prospective clinical trials and propose CXCL12 expression and percentage of embryonal carcinoma as clinically useful biomarkers to assist in stratifying patients for adjuvant therapy.

## Abstract

**Purpose:** Up to 50% of patients diagnosed with stage I non-seminomatous germ cell tumors (NSGCT) harbor occult metastases. Patients are managed by surveillance with chemotherapy at relapse or adjuvant treatment up-front. Late toxicities from chemotherapy are increasingly recognised. Based on a potential biological role in germ cells/tumors and pilot data, our aim was to evaluate tumor expression of the chemokine CXCL12 alongside previously proposed markers as clinically useful biomarkers of relapse.

**Experimental design:** Immunohistochemistry for tumor expression of CXCL12 was assessed as a biomarker of relapse alongside vascular invasion, histology (percentage embryonal carcinoma) and MIB1 staining for proliferation in formalin fixed paraffin-embedded orchidectomy samples from patients enrolled in the Medical Research Council's TE08/22 prospective trials of surveillance in stage I NSGCT.

**Results:** TE08/TE22 trial patients had a 76.4% 2-year relapse free rate (RFR) and both CXCL12 expression and percentage embryonal carcinoma provided prognostic value independently of vascular invasion (stratified log rank test  $p=0.006$  for both). There was no additional prognostic value for MIB1 staining. A model using CXCL12, percentage embryonal carcinoma and VI defines 3 prognostic groups that were independently validated.

**Conclusions:** CXCL12 and percentage embryonal carcinoma both stratify patients' relapse risk over and above vascular invasion alone. This is anticipated to improve the stratification of patients and identify high-risk cases to be considered for adjuvant therapy.

## Introduction

Although overall rare cancers, Testicular Germ Cell Tumors (TGCT) are the most common solid malignancy to affect young adult Caucasian males. They are divided into seminomas that resemble primordial germ cells or non-seminomas (NSGCT) that exhibit embryonal or extra-embryonal patterns of differentiation (1). 60% of NSGCT present with stage I disease (confined to the testes) and this proportion is increasing (2). 15-50% of such patients may harbor micro-metastatic disease and will relapse without further treatment (3, 4).

Assuming protocols are closely adhered to (5), immediate adjuvant treatment (1-2 cycles of Bleomycin, Etoposide and Cisplatin – BEP (6)) or surveillance with chemotherapy as salvage both have excellent rates of cure. Chemotherapy may have significant long-term effects including cardiovascular disease (7, 8), second malignancies (9, 10), Reynauds syndrome, neuropathies, fertility and emotional disorders. For this reason the routine use of adjuvant chemotherapy has been criticised in some quarters. Adjuvant retroperitoneal lymph node dissection is an alternative adjuvant strategy but with associated potential complications and morbidities these patients would also benefit from improved risk-stratification (11-13).

Histological evidence of vascular invasion (VI) is the only validated histological prognostic factor currently used to define risk of relapse in clinical stage I NSGCT (3, 4), not-with-standing the distinct role that plasma tumor markers play in assessing disease state. Tumors with VI have relapse rates of up to 50% and patients may be offered adjuvant chemotherapy reducing subsequent relapses to approximately 2% (6). In the absence of VI, around 15% patients relapse and surveillance may be a

more reasonable option. More accurate stratification of patients for likelihood of relapse would improve patient management, decreasing the risk of unnecessary treatment with associated side-effects and reducing risks and costs associated with over-treatment and excess imaging (14).

A systematic review (4) identified the percentage of Embryonal Carcinoma (%EC) within the primary tumor and the proliferation marker MIB1 as promising markers for relapse. Both are continuous variables and studies used differing cut-offs with variable levels of risk prediction. Univariate odds ratios for relapse range from 2.8 to 9 (4, 15-18). However %EC and VI were correlated and in multivariate analysis the prognostic effect of %EC diminished (4). In a subsequent study, 77% of metastasizing tumors showed MIB1 staining in >70% cells, equivalent to an odds ratio of 3.18 (95% CI: 1.51, 6.65). However, these data derive from a series of 195 patients after retroperitoneal lymph node dissections and not strictly a surveillance population (19).

Considering novel molecular markers, TGCT resemble primordial germ cells (PGC) (20) that physiologically utilize KITLG/KIT and CXCL12/CXCR4 for migration and survival during embryological development (21-23) and in the maintenance of the spermatogonial stem cell niche (24). Both signalling pathways are implicated in the malignant counterpart; KIT is strongly implicated in testicular tumorigenesis with both discrete amplification and activating mutations described (20) and TGCT express the receptor CXCR4 that can mediate invasive migration towards its ligand, CXCL12 (25, 26). Importantly, NSGCTs can express CXCL12 in an autocrine fashion and in a pilot study containing a high proportion of Embryonal Carcinoma cases (n=80) of

TGCT this was associated with reduced risk of relapse (25). Here we investigate this feature further for clinical utility alongside VI, %EC and MIB1 staining and in addition to other clinicopathological characteristics.

Two recent clinical trials conducted by the Medical Research Council (MRC) investigated surveillance strategies in clinical stage I NSGCT and provide a unique large cohort of well-characterized stage I NSGCT patients managed on prospective protocols. TE08 (NCT00003420) (27) compared two frequencies of CT scanning during surveillance of stage I NSGCT. TE22 (NCT00045045) (28) investigated the ability of a baseline FDG PET scan to distinguish patients at lower risk of relapse who might safely be managed by surveillance. In total 501 patients were managed by surveillance, with a relapse rate of 21% (103 patients). 130/501 (26%) had VI, although in an unselected population this would be closer to 50%.

To refine treatment stratification in patients diagnosed with stage I NSGCT we set out to investigate and validate CXCL12, %EC and MIB1 as biomarkers prognostic for relapse. Additional evidence of a prognostic effect for CXCL12 over and above the previous pilot data (25) was first sought using a tissue microarray (TMA) comprising representative cores from 59 stage I NSGCT patients managed with surveillance. We then investigated the markers MIB1, CXCL12 and %EC in the samples from stage I NSGCT patients managed by surveillance in the MRC TE08/TE22 clinical trials. Finally, we validated a combined prognostic model using VI, CXCL12 and %EC in the previous cohort of samples (25).

## **Materials and Methods**

### *Patients and tumor samples*

This study has national research ethics committee approval (09/MRE00/30) and complies with the REMARK guidelines for biomarker studies (29). Samples from stage I NSGCT patients managed by surveillance were collected from the Medical Research Council (MRC) trials TE08(27, 28). Specifically these were patients diagnosed with stage I NSGCT (negative tumour markers and CT scan confirming stage I) and enrolled post-operatively into a randomised study of two alternate imaging surveillance protocols (TE08) or in the case of TE22, undergoing FDG-PET imaging followed by surveillance if negative, to assess the negative predictive value of this scan. Formalin fixed paraffin embedded (FFPE) tumor blocks were available for 200/501 (40%) cases; 139 from TE08 and 61 from TE22. 10 of the 61 TE22 patients were not eligible for this study as 7 were PET positive and received adjuvant chemotherapy and 1 was PET negative but received adjuvant chemotherapy at the patient's request. 2 cases were lost to follow up. The final trial samples consisted of material from 190 patients (Table1). Importantly, this cohort was representative of the overall trial sample set with a relapse-free rate of ~78% at 2 years after orchidectomy (Supplementary Material Figure S1). Complete tumor cases were retrieved from each patient, and a full set of haematoxylin and eosin (H&E) sections from tumor for each case were examined by a board certified histopathologist. Representative tumour material, to include all significant areas of pathology, was selected from each case. VI was assessed as previously described (16). Additionally, sections from a TMA containing 0.6mm diameter cores were available from Princess Margaret Hospital, Toronto (JS) representing primary tumors from 59 patients with stage I NSGCT managed by surveillance and a minimum follow-up of 2

years (Supplementary Material Table S1). Finally, TMAs comprising material from 80 patients with stage I non-seminomatous germ cell tumors managed with surveillance at the Royal Marsden Hospital (RMH, previously described in (25)) were re-scored as per the below by a pathologist (DB) blinded to outcomes.

#### *Sectioning, histology review and staining*

Sections were stained with H&E and assessed to ensure adequate tumor material. Sections were deparaffinized prior to staining for CXCL12 (Antibody 79018, 1:100, R&D Systems, Minneapolis MN, USA), including positive (tonsillar crypt) and negative controls as previously described (25), and MIB1 (Antibody M7240, 1:100, Dako). Antibodies were visualized using the Bond Polymer Refine Kit (Leica Biosystems, Newcastle, UK). Immunostaining was performed on a Bond max automated immunostainer.

#### *Scoring and categorization*

H&E slides were scored for %EC (by DB) as a continuous variable, and then additionally grouped as described in previous studies or a new data-derived grouping for subsequent analysis. Immunostaining was scored by two independent histopathologists (KT and IC) recording intensity of staining as 0-3 (absent, weak, medium and high intensity) and % cells staining positive. Samples where scores differed were reviewed and a consensus obtained. Scores were categorized as absent/weak CXCL12 if <1% cells across the whole tumor stained positive for CXCL12. A second exploratory analysis was also performed classifying <10% cells staining for CXCL12 as CXCL12 absent/weak. Analysis for MIB1 was performed separately using both intensity and % cells positive using cut-offs described in the

previous studies i.e.  $\geq 70\%$  and  $\geq 40\%$  (15-18) as well as additional exploratory analyses.

### *Statistical methods*

The primary outcome measure was relapse-free rate, measured from the date of orchidectomy to the date of relapse confirmation, with relapse-free patients censored on the date last known to be alive. Relapse free rates on Kaplan Meier survival curves were compared by the logrank test, with an initial assessment of the independence of %EC, CXCL12 and MIB1 over VI determined by logrank tests stratified for VI. Subsequently, a proportional hazards regression model was fitted to adjust for baseline clinical variables, VI, %EC, MIB1 and CXCL12 staining, using forward and backwards stepwise selection. Chi squared tests were used to investigate the association of %EC and CXCL12/MIB1 staining with clinicopathological variables.

## Results

### *TMA CXCL12 expression and outcome*

To investigate CXCL12 as a marker for relapse prior to application to the clinical trial sample sets, we first studied the TMA representing 59 cases. 25/59 cases (42.4%) demonstrated moderate/strong expression of CXCL12 cells of which 3 had relapsed (RFR 88.0%). Of the 34 patients with absent/weak staining for CXCL12, 11 had relapsed (RFR 67.6%, log rank test  $p=0.68$ ). Although not reaching statistical significance, the rates of CXCL12 expression and subsequent relapse were consistent with previous data [25] and analysis of CXCL12 expression was taken forward to TE08/TE22 samples.

### *TE08/22 – CXCL12 and relapse*

Samples representing 182/190 samples from patients in the TE08 and TE22 trials were assessable for CXCL12 staining (Table 1; Figure 1A,B,C,D). Using  $<1\%$  as the cut off, 37 (20.3%) tumors were classified as absent/weak with the other 145 (79.7%) moderate/strong (as scored by two pathologists,  $\kappa=0.465$ ,  $p<0.001$ ). In an exploratory analysis, a  $<10\%$  cut off was also assessed and produced a similar performance (Supplementary Table S2 and Figure S2). Therefore either cut-off may be used. There was no association between CXCL12 and VI, the presence of seminomatous elements or raised markers pre-orchidectomy. There was however a strong association with the presence of Embryonal Carcinoma which was more prevalent in those with absent/weak staining, (75.7% vs 30.3% of those with moderate/high staining, chi-square  $p < 0.001$ ). The logrank test shows evidence of a prognostic impact, alone and stratified by VI ( $p=0.006$ , Table 2) for CXCL12 with reduced relapse-free rate in the absent/weak group (Figure 2A). In VI+ve patients,

CXCL12 further stratified relapse rates; 56 patients with VI but moderate/strong staining had a 2-year RFR of 62.4% versus 11 patients with VI and absent/weak staining for CXCL12 where a RFR of 27.3% was observed (95% CI 1-53.6%).

#### *TE08/22 - %EC and relapse*

177/190 patients were assessable for %EC (Table 1; Figure 1E,F,G). This showed a bimodal distribution, with clusters at 0% and 100% and a relatively even spread of the remaining values between these levels. %EC was significantly higher in patients with VI (median 70% vs 20%, Mann Whitney test  $p=0.013$ ), and also in those with absent/weak CXCL12 intensity (medians 100% vs 20% for moderate/strong,  $p<0.001$ ) and with presence of MIB1 staining (medians 50% vs 10%,  $p=0.012$ ).

%EC was assessed as in previous reports as a continuous variable, a binary variable (presence/absence) and applying previously reported cut-offs, (<45% 46-70%, >70%; above or below 50%) (4). In addition, to better reflect the unusual distribution of %EC, a data-derived categorization was investigated, formed by dividing the data initially into approximate quintiles (0%, 1-25%, 26-75%, 76-99%, 100%), collapsing groups with similar relapse free rates to create an “optimal” categorisation ( $\leq 25\%$ , 26-99%, 100%). With the exception of presence/absence of EC, higher %EC was associated with higher relapse rates (Figure 2B, Table 2), independent of VI, for all categorizations.

#### *TE08/TE22 - MIB1 and relapse*

179 cases were assessable for MIB1 staining; 45 (25.1%) were MIB1 weak on both intensity and proportion of cells staining (Table 1; Figure 1H). There was a significant

association of both MIB1 intensity and proportion of cells staining for MIB1 with decreasing likelihood of the tumor containing seminomatous elements (Mann-Whitney test  $p < 0.001$ ) and increasing likelihood of VI (Mann-Whitney test  $p = 0.004$  for intensity and  $p < 0.001$  for % cells staining).

There was no evidence of prognostic value for MIB1 staining intensity (logrank test for trend  $p = 0.26$ ) nor for the proportion of cells staining positive for MIB1, either when analysed as per previous reports ( $\geq 70\%$  and  $\geq 40\%$ ), as quartiles or using a log rank test for trend. In contrast to previous studies, only 5/179 patients (3%) had MIB1 staining in  $>70\%$  of cells. The main distinction observed was between the 45 samples with weak vs any staining for MIB1 (Table 2). Analyzing the samples in this binary fashion (MIB1 positive or negative) has a prognostic effect (univariate analysis), which was reduced after stratification for VI (Table 2).

#### *TE08/TE22 multivariate analyses*

Multivariate analyses were performed on the 177 patients with complete data to assess the additional prognostic value of these factors over and above the presence/absence of VI and clinical variables, specifically VI (yes/no), histology type, seminomatous elements present/absent, age (continuous variable), alpha feto protein (AFP) raised pre-orchidectomy (yes/no), human chorionic gonadotrophin (HCG) raised pre-orchidectomy (yes/no), CXCL12 expression, MIB1 staining (high/weak), %EC (as a continuous variable), EC present/absent, and %EC categorised according to previous studies, ( $<45$ ,  $46-79$ ,  $>80\%$ ;  $<50\%$  vs  $\geq 50\%$  (4)) and %EC categorised according to the optimal cut offs for this dataset ( $<25\%$ ,  $26-99\%$  and  $100\%$ ).

Both forward and backwards stepwise model selection procedures were used which all led to a model (Supplementary Table S3) including only VI and %EC (continuous variable). Dropping EC as a continuous variable but keeping all the other variations, the model includes only VI and %EC, using the “optimal” categorization (Figure 2B). However, if %EC is used as previously reported (4), then VI and CXCL12, but not %EC, are retained as independent variables (Figure 2A).

Components of both groupings have potential clinical utility in defining subsequent relapse risk, and as such we also present a combined model (Figure 3A) using 3 prognostic groups. Specifically these are a) VI negative, <100% EC and any CXCL12, b) VI negative, 100% EC any CXCL12 or VI positive, any EC and mod/high CXCL12 and c) VI +ve, any EC, absent/weak CXCL12 with 2 year RFRs of 94.3% (95% CI 89.4%, 99.2%), 63.9% (52.9%, 74.9%) and 30% (1.6%, 58.4%) respectively (Logrank  $X^2=38.6$  on 2.d.f,  $p<0.001$ ).

#### *Validation of model combining VI, CXCL12 and %EC on RMH cohort.*

Staining characteristics of the cohort of 80 RMH patients managed with surveillance mirror those of the TE08/TE22 patients and are detailed in Table 3. Overall this cohort experienced a 2 year relapse-free rate (RFR) of 60% (95% CI 59%-71%). For patients with VI this was 47% (31-63%) as opposed to 69% without (54-83%).

CXCL12 expression was prognostic as previously described with tumors absent/weak for CXCL12 having a 2 year RFR a relapse rate of 34% (18-50%) whereas tumors with moderate/high staining for CXCL12 having an RFR of 80% (68-92%), HR 0.24 (95% CI 0.11-0.49)  $p<0.001$ . CXCL12 expression retained additional prognostic value over and above VI (stratified log rank  $p<0.001$ ). Finally

%EC was also prognostic in this cohort, with the 2year RFR ranging from 79%(61-87) for patients with  $\leq 25\%$ EC, through 64% (47-82%) with 26-99% EC and 48% (31-66%) in cases that were 100%EC(  $p=0.04$ ). Using the combined model developed above, the three risk groupings have 2 RFR of 81.5%(66.8-96.2) for patients that were VI negative and %EC<100, 64.5%(47.6-81.4) for those that were VI negative and 100%EC or VI positive and moderate/high expression of CXCL12 and a 2 year RFR of just 27.3%(8.7-45.9) for those that had evidence of VI but absent/weak expression of CXCL12 (Figure 3B, log rank Mantel-Cox <0.001).

#### 4. Discussion

Using samples from stage I NSGCT patients managed by surveillance on prospective clinical trials, we have demonstrated that CXCL12 expression and %EC in the primary tumor are both predictors of relapse independently of VI. Importantly, samples obtained for this study were representative of the total trial populations (~78% relapse free rate at 2 years, by which point most relapses have usually occurred) and hence represent patients with stage I NSGCT in general. This provides prognostic information of potential clinical relevance over and above the presence/absence of histological evidence of VI being able to identify patients at low, moderate or very high risk of relapse (Tables 2, S2 and 3). However, %EC and CXCL12 are strongly inversely correlated and in multivariate analyses one or the other – but not both - are selected in addition to VI, depending on how %EC is analyzed. This gives two alternative prognostic models, as illustrated in figure 2A and B. While the multivariate analysis does not support a model containing VI with both %EC and CXCL12, there are unique characteristics of each combination that an ideal model would combine, specifically identification of a very high risk group (VI+, CXCL12 absent) and identification of VI-ve patients who have a prognosis closer to that of VI+ve (VI-ve, 100% EC). We therefore derived an exploratory model that combines these elements into 3 prognostic groups (Figure 3A). Using a cohort of cases previously characterised for CXCL12 (25), through the additional inclusion of %EC scores and VI we demonstrate the validity of this model (Figure 3B).

As shown previously (4), %EC is a subjective analysis. CXCL12 expression (<1% or <10% of cells staining positive as thresholds to classify as CXCL12 absent) together

with VI is a potentially more reproducible approach to determining prognosis, and one that identifies a small group of patients with distinctly poorer prognosis.

Adjuvant chemotherapy is highly effective treatment and even 1 cycle, currently being tested in a UK phase II single arm study (BEP 111), can reduce the risk of recurrence to <4% (11, 13, 30, 31). This use of adjuvant BEP has been criticized for exposing a high proportion of patients who would not go on to relapse to intensive chemotherapy. Refining the current risk model (utilizing lymphovascular invasion alone) would therefore be of clinical utility, allowing adjuvant therapy to be focused on those at highest risk. Our analysis in this cohort supports our previous study in showing that CXCL12 immunohistochemistry can add valuable additional prognostic information to the model based on VI, particularly in identifying a small cohort of patients with a very high risk of relapse. It also identifies groups for whom surveillance, potentially using reduced intensity follow-up (at least to the reduced frequency arm used in TE08 (27)) may be most appropriate, and those suitable for either surveillance or adjuvant therapy depending on personal preferences. Identifying high-risk patients for minimal effective adjuvant therapy will ultimately reduce long-term side-effects for the stage I NSGCT population as a whole.

The prognostic value of MIB1 was analysed in a number of ways looking at intensity and percentage of cells both separately and combined and including previously used categorizations (15-18, 32). Although patients with weak staining have a better prognosis than those with high MIB1 staining, MIB1 staining was associated with VI and in multivariate analysis does not add clear independent prognostic value.

The prognostic value of CXCL12 expression demonstrated here is consistent with a growing body of evidence for the CXCL12/CXCR4 axis supporting the male germ cell niche and the metastatic spread of cancers, including this possibility in germ cell tumors (20, 22, 24, 25, 26, 33). CXCR4 expression is associated with invasion and metastases in a range of tumor types where lower levels of CXCL12 in tumors predict a reduced risk of metastatic dissemination (34, 35). Furthermore in breast cancer, lower levels of plasma CXCL12 appear to be associated with increased risk of metastatic relapse (36). That autocrine expression of CXCL12 consistently reduces the subsequent risk of relapse in stage I NSGCT (independent from histological VI) suggests the abrogation of a chemokine gradient (towards CXCR4) might prevent extravasation into the vascular compartment and/or invasion at metastatic sites. To this end, assessment of stromal and/or plasma CXCL12 might provide additional prognostic information. The apparent association with histological subtypes of TGCT requires further work, aligned with a better understanding of how these tumours develop from the *in situ* counterpart. Further investigations will inform on these and other potential mechanisms of dissemination and relapse (37) in NSGCT patients.

As a paradigm of protecting future quality of life in an essentially curable disease a prospective study is recommended (potentially also investigating novel imaging strategies in an effort to minimize radiation exposure e.g. MRI) to further validate the prognostic value of CXCL12 and/or %EC expression in addition to the presence/absence of histological VI. This is anticipated to lead to the ability to identify patients diagnosed with stage I NSGCT at high risk of relapse to be considered for adjuvant therapy whilst others are safely surveilled.

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**Table 1.** Histological subtype, % Embryonal Carcinoma, tumour marker status and immunohistochemistry for CXCL12 and MIB1 for 190 stage I NSGCT primary tumours from patients treated in the TE08/TE22 clinical trials. EC - embryonal carcinoma; CC - choriocarcinoma; AFP - alpha fetoprotein; HCG - human chorionic gonadotrophin.

		Vascular invasion				Total			
		No		Yes		N		%	
		N	%	N	%				
<b>Histology</b>	Pure EC	23	19%	18	27%	41	22%		
	Mixed NSGCT	79	65%	39	57%	118	62%		
	Yolk sac	8	7%	3	4%	11	6%		
	Differentiated Teratoma	9	7%	1	2%	10	5%		
	Other type	3	3%	7	10%	10	5%		
<b>EC presence</b>									
	EC absent	29	26%	9	14%	38	22%		
	EC present	83	74%	56	86%	139	79%		
	<i>Not known</i>	10		3		13			
<b>% EC Optimal categories</b>									
	<=25%	60	54%	20	31%	80	45%		
	26-99%	29	26%	27	42%	56	32%		
	100%	23	21%	18	28%	41	23%		
	<i>Not known</i>	10		3		13			
<b>AFP pre-orchidectomy</b>									
	normal	53	43%	25	37%	78	41%		
	raised	69	57%	43	63%	112	59%		
<b>HCG pre-orchidectomy</b>									
	normal	65	53%	29	43%	94	50%		
	raised	57	47%	39	57%	96	50%		
<b>CXCL12 (≤1%= weak/absent)</b>									
	Insufficient tumor	7	6%	1	2%	8	4%		
	absent/weak	26	21%	11	16%	37	20%		
	moderate/strong	89	73%	56	82%	145	76%		
<b>MIB1 staining</b>									
	Weak	37	33%	8	12%	45	25%		
	High	77	68%	57	88%	134	75%		
	<i>Not assessable</i>	8		3		11			
<b>Total</b>		122	64%	68	36%	190	100%		

**Table 2.** Univariate and stratified logrank test results for factors of interest in 177 cases with complete data.

	No. Pts	2 yr RFR	95% CI	Log rank p-value	Stratified (by VI) log rank p-value
<b>Vascular invasion</b>					
absent	112	88.3	(82.2, 94.4)	<0.001	n/a
present	65	58.3	(46.0, 70.6)		
<b>CXCL12</b>					
mod/high, no VI	87	90.7	(84.6, 96.8)	0.078	0.009
mod/high, VI	55	63.7	(50.6, 76.8)		
<i>Sub-total mod/high</i>	142	80.4	(73.7, 87.1)		
absent /weak, no VI	25	80.0	(64.3, 95.7)		
absent /weak, VI	10	30.0	(1.6, 58.4)		
<i>Sub-total absent</i>	35	65.7	(50.0, 81.4)		
<b>MIB1</b>					
Weak, no VI	35	91.4	(82.2, 99.9)	0.007	0.045
Weak, VI	7	100.0	(39.8, 99.9)		
<i>Sub-total weak</i>	42	92.8	(85.0, 99.9)		
High, no VI	76	86.7	(79.1, 94.3)		
High, VI	56	53.7	(40.4, 67.0)		
<i>Sub-total high</i>	132	73.0	(65.4, 80.6)		
<b>EC (Present/absent)</b>					
Absent, no VI	29	96.3	(89.2, 99.9)	0.096	0.243
Absent, VI	9	66.7	(35.9, 97.5)		
<i>Sub-total absent</i>	38	89.2	(79.2, 99.2)		
Present, no VI	83	85.4	(77.8, 93.0)		
Present, VI	56	56.8	(43.5, 70.1)		
<i>Sub-total present</i>	139	74.3	(66.9, 81.7)		
<b>EC (optimal categories)</b>					
≤25%EC, no VI	60	94.9	(89.2, 99.9)	<0.001	0.006
≤25%EC, VI	20	68.1	(46.9, 89.3)		
<i>Sub-total ≤25%</i>	80	88.4	(81.3, 95.5)		
26-99%EC, no VI	29	93.1	(83.9, 99.9)		
26-99%EC, VI	27	57.3	(38.1, 76.5)		
<i>Sub-total 26-99%</i>	56	76.4	(65.2, 87.6)		
100% EC, no VI	23	63.6	(43.4, 83.8)		
100%EC, VI	18	50.0	(26.9, 73.1)		
<i>Sub-total 100%</i>	41	57.5	(42.2, 72.8)		

**Table 3.** Tumor characteristics (VI, CXCL12 and %EC) for 80 stage I NSGCT RMH patients managed by surveillance

		No Vascular Invasion		Vascular invasion +		Total	
		N	%	N	%	N	%
<b>EC%</b>	<=25%	14	33.3%	5	13.9%	19	24.4%
	26-99%	13	31.0%	15	41.7%	28	35.9%
	100%	15	35.7%	16	44.4%	31	39.7%
<b>CXCL12 Intensity</b>	0	11	26.2%	19	50.0%	30	37.5%
	1	2	4.8%	3	7.9%	5	6.3%
	2	9	21.4%	5	13.2%	14	17.5%
	3	20	47.6%	11	28.9%	31	38.8%
<b>CXCL12</b>	Absent/low	13	31.0%	22	57.9%	35	43.8%
	Moderate/high	29	69.0%	16	42.1%	45	56.3%
<b>Proposed grouping</b>	No VI, %EC <100	27	64.3%	0	0.0%	27	33.8%
	No VI and 100% EC or VI and mod/high CXCL12	15	35.7%	16	42.1%	31	38.8%
	VI and absent/low CXCL12	0	0.0%	22	57.9%	22	27.5%
<b>TOTALS</b>		42	100%	38	100%	80	100%

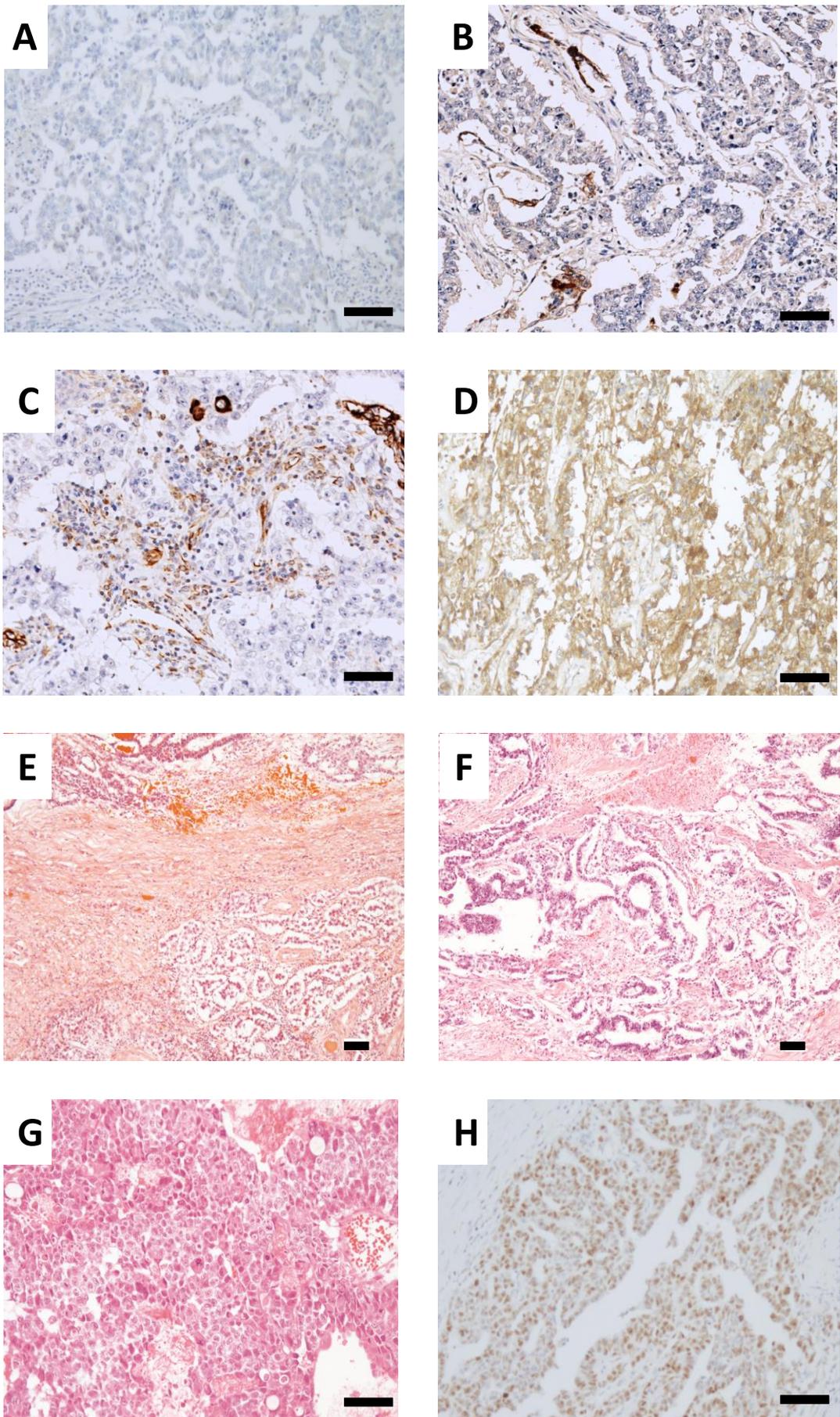
## Figure Legends:

**Figure 1. Representative staining of stage I non-seminomatous germ cell tumor samples.** Immunohistochemistry for CXCL12 staining **A** negative **B** <10% **C** ~30% and **D** 100% positive. **E** Haematoxylin and Eosin staining showing a combined seminomatous (bottom of photomicrograph) and non-seminomatous tumour (top) composed of less than 10% embryonal carcinoma. **F** Tumor composed of 25% embryonal carcinoma and 75% yolk sac tumor. The yolk sac and embryonal carcinoma are intermingled in a polyembryomatous fashion, mimicking the earliest stages of embryonic development. **G** Tumor entirely composed of embryonal carcinoma. **H** Embryonal carcinoma showing 75% positivity for MIB1 immunohistochemistry. (Scale bar, 100 microns).

**Figure 2. Relapse-free survival data for:****A** 182 patients with stage I NSGCT managed by surveillance with the MRC TE08 and TE22 clinical trials, stratified by vascular invasion (VI) and immunohistochemistry for CXCL12 in combination with the presence/absence of histological VI (stratified log rank test  $p=0.006$ ). **B** 177 patients with stage I NSGCT managed with surveillance stratified by %EC and VI, illustrated using optimal cut-offs for %EC.

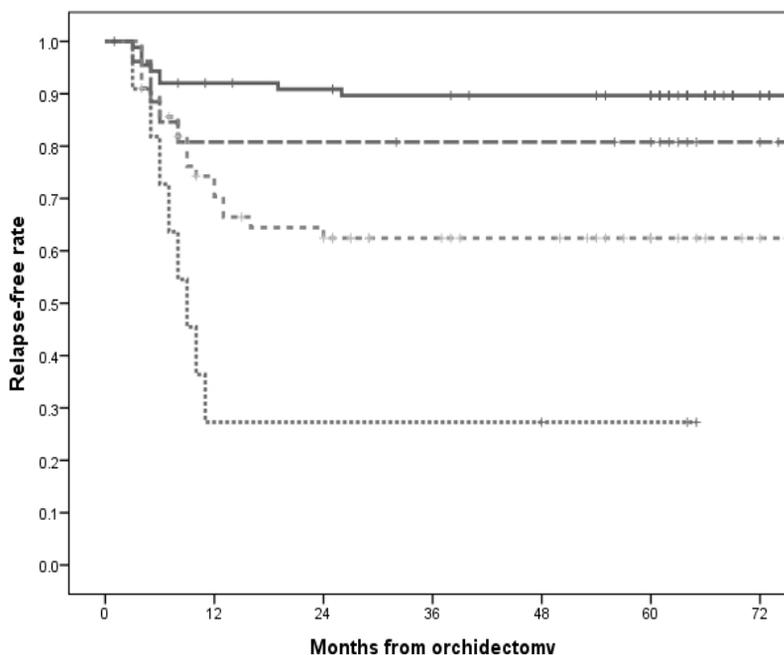
**Figure 3. Relapse free survival data for:****A** 177 patients with stage I NSGCT managed with surveillance according to data-derived risk groups. **B** An independent cohort of 80 patients with stage I NSGCT managed with surveillance stratified by the risk grouping derived in A.

Figure 1.



**Figure 2**

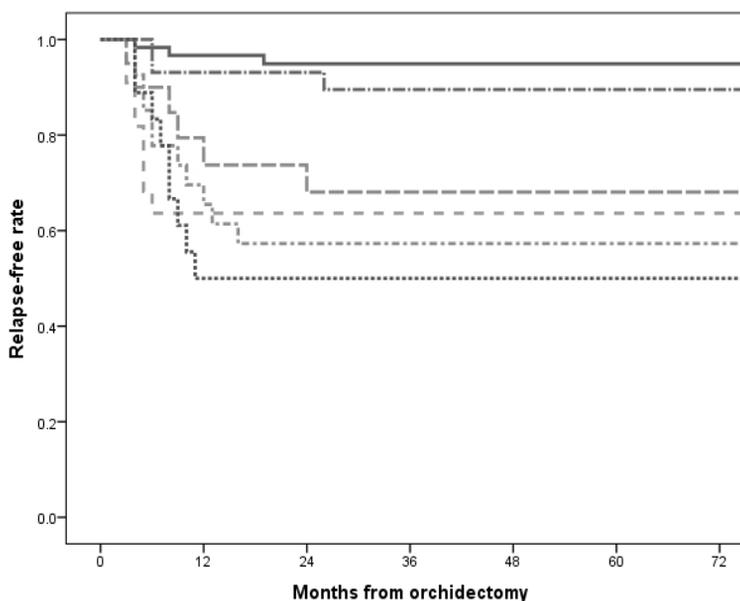
**A**



Numbers at risk	0	12	24	36	48	60
High CXCL12, no VI	89	79	77	75	73	67
Low CXCL12, no VI	26	21	21	20	20	18
High CXCL12, VI+ve	56	36	30	23	20	11
Low CXCL12, VI+ve	11	3	3	3	2	2

Logrank  $\chi^2=34.8$  on 3.d.f,  $p<0.001$

**B**



Numbers at risk	0	12	24	36	48	60
No VI $\leq 25\%$ EC	60	56	54	54	53	50
No VI 26-99% EC	29	27	27	24	23	20
No VI 100% EC VI	23	14	14	14	14	13
+ve, $\leq 25\%$ EC VI	20	13	11	8	7	4
+ve, 26-99% EC VI	27	16	14	11	10	6
+ve, 100% EC	18	9	8	7	5	3

Logrank  $\chi^2=34.8$  on 3.d.f,  $p<0.001$



# Clinical Cancer Research

## Defining a New Prognostic index for Stage I Non-seminomatous Germ Cell Tumors using CXCL12 Expression and Proportion of Embryonal Carcinoma

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