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## The impact of fertility preservation on treatment delay and progression-free survival in women with lymphoma: a single-centre experience

Lymphoma affects many young women of childbearing age. The American Society of Clinical Oncology recommends early discussion of the reproductive risks of treatment, and referral to fertility preservation (FP) specialists when appropriate (Loren *et al*, 2013). Women referred for FP undergo a several-step process including ovarian stimulation, oocyte retrieval, and oocyte or embryo storage (De Vos *et al*, 2014). Barriers to FP include poor access to reproductive specialists and concerns for treatment delay (Quinn *et al*, 2009, 2015). The real-world treatment delay and outcomes among female lymphoma patients attempting FP with modern techniques have not been reported previously.

At our institution, practitioners are required to address fertility in all newly diagnosed cancer patients through the use of automated prompts in the electronic medical record. We have a dedicated in-house fertility preservation patient

© 2016 John Wiley & Sons Ltd British Journal of Haematology, 2018, **180**, 889–918 navigator (FPPN) to educate patients and expedite referrals to the reproductive specialists. We performed a retrospective chart review of lymphoma patients that contacted any fertility specialist prior to treatment at Northwestern University from 1 May 2006 until 31 August 2015. Patients who underwent FP were compared to women that contacted a FPPN but did not undergo preservation. The Northwestern University institutional review board approved the use of the clinical database for this project.

Our primary objective was to assess differences in time to treatment (TTT) associated with FP. In newly diagnosed patients, TTT was defined as the time from the initial haematology consultation until the initiation of therapy. In patients with relapsed disease, TTT was defined as the time from the date of biopsy or haematology consultation until treatment initiation. Our secondary objective was to assess



progression-free survival (PFS), which was defined as time from date of treatment until progression or death. The Wilcoxon rank sum test was used to compare age, TTT and follow-up time between groups. Fisher's exact test was used to compare stage, planned treatment setting, and Eastern Cooperative Oncology Group performance score (ECOG PS). Kaplan–Meier curves with the log rank test were used to compare PFS between the two groups, using a two-tailed significance level of 0.05.

A total of 128 patients were identified from a fertility patient log, including 40 who underwent FP. Thirty-three of 40 patients undergoing FP and 50 of 93 patients who chose not to undergo FP were available for analysis. Reasons for exclusion included patients seen by reproductive endocrinology only and no haematologist consulted at Northwestern, lack of chemotherapy treatment records, or no treatment received following fertility contact. Pertinent baseline

Table 1	I.	Patient	characteristics.
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characteristics are outlined in Table I. Comparing the two groups, there was a significant difference in age (P = 0.01), but not in stage (P = 0.05), planned treatment setting (P = 0.99) or ECOG PS (P = 0.99). Median follow-up was 39.3 (1.5–103.4) months, and did not differ between controls and those undergoing FP (P = 0.16).

Median TTT among FP patients was 28 days overall (range: 18–76) versus 15.5 days (range: 0–74) for controls (P < 0.001; Fig 1A). Factors other than FP led to treatment delays prior to and after FP. The median time to first contact with a fertility specialist was 0 days (range -15 to +11) from haematology consultation, with several patients having contact prior to their haematology visit. The median time from oocyte retrieval until treatment initiation was 5 days (range 0-21). Seven patients had greater than 8 days from oocyte collection to treatment. The reasons were variable: 3 had delays in diagnostic work-up; 1 deferred for a trip out of

Patient characteristics Median age (range), years Disease	Patients under n (%)* 26 (20–35)	going fertility prese	rvation	Controls n (%)* 29 (17–45)		
	All	HL	NHL	All	HL	NHL
Number of patients	33 (100)	21 (67)	12 (33)	50 (100)	31 (62)	19 (38)
Planned treatment setting						
Frontline	25 (76)	14 (67)	11 (92)	37 (74)	24 (77)	13 (68)
R/R	8 (24)	7 (33)	1 (8)	13 (26)	7 (23)	6 (32)
Treatment						
ABVD	12 (36)	12 (57)	_	23 (46)	23 (74)	
R-CHOP-like†	8 (24)	_	8 (64)	6 (12)	_	6 (32)
escBEACOPP	2 (6)	2 (10)	_	1 (2)	1 (3)	
DA-EPOCH-R	2 (6)	_	2 (18)	4 (8)	_	4 (21)
R-Hyper-CVAD	1 (3)	_	1 (9)	4 (8)	_	4 (21)
HDCT + SCT	7 (21)	7 (33)	_	10 (20)	7 (23)	3 (16)
Other‡	1 (3)		1 (9)	2 (4)		2 (11)
Ann Arbor Stage§						
I/II	28 (85)	17 (81)	11 (92)	32 (64)	23 (74)	9 (47)
III/IV	5 (15)	4 (19)	1 (8)	17 (34)	8 (26)	9 (47)
Bulky (>10 cm)	9 (27)	6 (29)	3 (25)	10 (20)	4 (13)	6 (32)
ECOG PS¶						
0	24 (73)	17 (81)	7 (58)	38 (76)	22 (71)	16 (84)
$\geq 1$	5 (12)	2 (10)	3 (25)	9 (18)	6 (19)	3 (16)
LDH > ULN	9 (27)	4 (19)	5 (42)	17 (34)	7 (23)	10 (53)
$\text{ESR} \ge 50$	9 (27)	9 (43)	_	11 (22)	11 (35)	_
B symptoms	10 (30)	8 (38)	2 (17)	19 (38)	13 (42)	6 (32)

HL, Hodgkin lymphoma; NHL, non-hodgkin lymphoma; R/R, relapsed/refractory; ABVD, Adriamycin, bleomycin, vinblastine, dacarbazine; R-CHOP, rituximab, cyclophosphamide, Adriamycin, vincristine, prednisone; escBEACOPP, escalated bleomycin, etoposide, Adriamycin, cyclophosphamide, vincristine, procarbazine, prednisone; DA-EPOCH-R, dose adjusted etoposide prednisone, vincristine, cycophosphamide, Adriamycin, rituximab; R-HyperCVAD, Course A- cyclophosphamide, vincristine, Adriamycin, dexamethasone, cytarabine, mesna, methotrexate; Course B- methotrexate, leucovorin, cytarabine; HDCT + SCT, high dose chemotherapy plus stem cell transplant; ECOG PS, Eastern Cooperative Oncology Group performance status; LDH, lactate dehydrogenase; ULN, upper limit of normal; ESR, erythrocyte sedimentation rate.

†R-CHOP, BR (bendamustine rituximab), R-CVP (rituximab, cyclophosphamide, vincristine, prednisone).

‡Involved field radiotherapy, radio-immunotherapy, romidepsin.

§One patient with NHL seen at relapse, initial staging information not available.

PECOG PS missing for 3 HL patients in control group, and 2 HL and 2 NHL in the fertility preservation group.

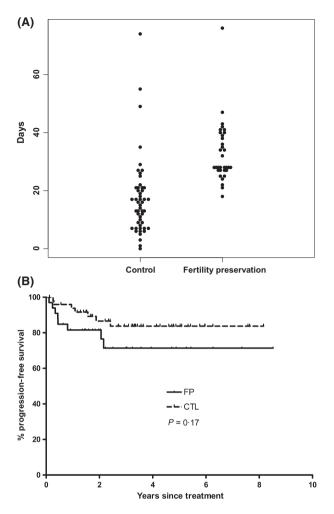


Fig 1. (A) Time to treatment among patients undergoing fertility preservation *versus* controls. Patients undergoing fertility preservation had longer time to treatment (P < 0.001). The median times to treatment were 28 and 15.5 days in patients undergoing and not undergoing fertility preservation respectively. (B) There was no difference in 5-year progression-free survival between patients undergoing or not undergoing fertility preservation (P = 0.11). CTL, control; FP, fertility preservation.

town; 2 were treated subsequently at an outside institution; and 1 had follicular lymphoma with no urgency to treat. The median number of days to complete stimulation protocol was 11 (range: 5–14). A median of 14 oocytes (range: 0–37) were retrieved per patient. In 2 women, no oocytes could be successfully retrieved. Five women achieved pregnancy following FP compared to 6 controls. Of these, 3 were spontaneous and 2 required reproductive assistance, one from frozen embryos and one from frozen oocytes. Of 3 women returning to use their frozen gametes, 2 were successful and 1 was unsuccessful. Ovarian stimulation did not result in any known complications.

In total, 15 patients relapsed after contacting a fertility specialist, including 7 patients in the control group and 8 in the FP arm. Patients who subsequently relapsed in both arms had high-risk features prior to fertility. There was no difference in 1-year and 5-year PFS between FP patients compared to controls (FP: 1-year PFS= 81.6%, 5-year PFS = 71.4%; Controls: 1-year PFS = 93.8%, 5-year PFS = 83.7%, P = 0.17; Fig 1B).

Our study has some limitations, including those associated with a retrospective analysis. Only patients who contacted a fertility specialist were included and therefore our analysis is subject to selection bias. Our population was heterogeneous with a wide range of lymphoma subtypes, stages and treatments received. Additionally, the relatively small number of patients available for analysis limited our ability to match patients based on age, disease or prognosis, and thus there were some baseline differences between groups. However, our uniform method of referral is a major strength of our study. To our knowledge, our institution is one of the few major universities with a FPPN to centralize the collection of data on women undergoing FP, and provides a bridge between the fields of reproductive endocrinology and malignant haematology. Overall, this analysis provides important information regarding the expected delays and outcomes associated with FP. Our study demonstrates that if referral is prompt, FP contributes minimal delay to treatment and is not associated with adverse outcomes. Furthermore, it underscores the importance of access to specialists in FP and the role of the fertility navigator.

## Author contributions

Drs. Allen and Winter had full access to all of the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis. *Study concept and design:* Winter, Allen, Pavone, Moravek. *Acquisition, analysis, or interpretation of data:* Allen, Pavone, Smith, Rafael Confino, Winter. *Drafting of the manuscript:* Winter, Allen. *Critical revision of the manuscript for important intellectual content:* Allen, Winter, Pavone, Smith, Gordon, Kazer, Lawson. *Statistical analysis:* Rademaker. *Obtained funding:* N/A. *Administrative, technical, or material support:* Confino, Smith. *Study supervision:* Winter, Pavone.

### **Conflict of interest disclosures**

The authors have no relevant conflicts of interest to disclose.

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# Pathology findings in patients with cutaneous T-cell lymphomas treated with allogeneic haematopoietic stem cell transplantation

Cutaneous T-cell lymphomas (CTCL) include mycosis fungoides (MF), Sézary syndrome (SS) as well as CD30 positive T-cell lymphoproliferative disorders (Bradford et al, 2009). One recently emerging therapeutic option in advanced stages of CTCL is allogeneic haematopoietic stem cell transplantation (HSCT) (Lechowicz et al, 2014). The rationale behind this strategy is the beneficial graft-versus-lymphoma effect, first described for CTCL by Burt et al (2000). It has been documented that this treatment achieves good response rates in patient groups which previously had a dismal prognosis (Hosing et al, 2015). Despite its highly beneficial impact on patient survival (Wingard et al, 2011), allogeneic HSCT poses several risks to the patient. Potential hazards include acute and chronic graft-versus-host disease (GvHD), infection with a variety of organisms and disease relapse (Hilgendorf et al, 2015). In this report, we describe the histopathology and immunohistochemistry findings in various biopsies performed following allogeneic HSCT in patients with CTCL at the Hammersmith Hospital.

All patients who had been treated with allogeneic HSCT for CTCL at the Hammersmith Hospital between 2004 and 2015 were included in the study. After exclusion of patients on whom no biopsies had been performed after HSCT, a total of 28 patients were included. Patients' characteristics and CTCL entities are summarized in Table SI.

A recurrence of the CTCL was seen in 14 patients, and could be classified as recurrent SS (n = 10), transformed MF (n = 3) and CD8<sup>+</sup> aggressive CTCL (n = 1). The median time to recurrence was 77 days (range 26–242 days).

Relapsed SS cases showed the classical morphology of medium sized atypical lymphoid cells with cerebriform nuclei (immunophenotype CD2<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup>CD5<sup>+</sup>/CD7<sup>-</sup>CD8<sup>-</sup>CD30<sup>-</sup>). Transformed MF cases showed large atypical lymphoid cells that expressed CD30 and had a high Ki67 index (>60%) along with other immunophenotypic features of MF (see Fig 1G–L). The case of CD8-positive aggressive CTCL showed medium to large sized atypical lymphoid cells with moderate amounts of cytoplasm and multiple small basophilic nucleoli (immuno-