ANZTCT Registry

Lymphoma Pre-infusion Data

1.PATIE		PART B: LYMPHOMA PRE-INFUSION			
Hospital:	AID (ABMTRR id):			mpleted for a previous co	
JPN:	DOB://	then skip	the diagnosis c	questions and go directly	to Section 4
lame ID:	Infusion date://	Lymphor	Lymphoma histology:		
		Diagnosi	s pathology rep	ort submitted to Registry	y: Y N
		_	(prior to any tr	ansformation)	
PART A: DI	SEASE CLASSIFICATION	1. DIAG	NOSIS (prior to	any transformation)	
		Lymphoma histology at diagnosis:			
		Diagnosi	s pathology rep	ort submitted to Registry	y: Y N
Date of diagnosis://	_ (of this histology)	lan an a l			l tiali
umphama histology at tim	o of infusion	Immunoi		tains performed: Y N	- -
ymphoma nistology at tim	e of infusion:		BCL-2	Pos Neg Unk	% pos
	. .		BCL-6	Pos Neg Unk	% pos
f DLBCL - germinal center E ubtype based on	3-cell type or activated B-cell type,		CD5	Pos Neg Unk	
	ochemistry (e.g. Han's algorithm)		CD10	Pos Neg Unk	
Gene expres			CD30	Pos Neg Unk	
Unknown m	ethod		C-MYC	Pos Neg Unk	% pos
			Cyclin D1	Pos Neg Unk	
Transformation from CLL: Y			EBER ISH	Pos Neg Unk	
If yes: Was 17p abnorm	ality detected: Y N		Ki-67	Pos Neg Unk	% pos
ransformation from diffor	ent lymphoma histology: Y N		MUM1	Pos Neg Unk	
			SOX11	Pos Neg Unk	
		Were cyt	ogenetics perfo	ormed: Y N Unk	
Date of original diagnos	is://	Tested vi	a FISH: Y N	Unk	
		• A	bnormalities ide	entified no abnormaliti	es
				ities:	
		Repo	ort submitted: Y	N	
	latest assessment prior to infusion:				
(N Not done			a karyotyping:		
If yes, Date of sca Deauville score: _		 Abnormalities identified No evaluable metaphas abnormalities 			ietaphases r
Deauville score				ities:	
		Repo	ort submitted: Y	N	
DISEASE STATUS AT INF	USION	2. LABO	RATORY VALU	JES AT DIAGNOSIS	
isease status at time of in	fusion:				Value
Date assessed:		WBC x1	10 ⁹ /L Mantle cel	l, Hodgkins only	
	_	Hb g/L	Follicular, Hod	gkins only	
		Abs lym	phocyte count x1	0 ⁹ /L Hodgkins only	
lumber of treatment lines	between diagnosis and infusion:	Lympho	ocyte % Hodgki	ns only	
	<u> </u>	Serum	albumin g/L Ho	odgkins only	
		LDH U/	ΊL		
		LDH UL	.N U/L		

Lymphoma	Pre-infusion	Data
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3. NODAL AND ORGAN			DISEASE TRANSFORMATION continued	-	
PET (or PET/CT) positive: Known nodal involvement If yes: Total number noda	Y N Not done t: Y N		 Were cytogenetics performed: Y N Unk Tested via FISH: Y N Unk Abnormalities identified no abnormalities Specify abnormalities: 	ies	
excluding follicular: \Box <td colspan="3"> Tested via karyotyping: Y N Unk Abnormalities identified No evaluable metaphases no abnormalities Specify abnormalities: Report(s) submitted: Y N </td>			 Tested via karyotyping: Y N Unk Abnormalities identified No evaluable metaphases no abnormalities Specify abnormalities: Report(s) submitted: Y N 		
Stage of organ involveme B symptoms present: Y		nknown	5. LABORATORY VALUES AT TRANSFORMA	TION	
ECOG score: [Lab parameters at transformationWBC x109/LMantle cell, Hodgkins onlyHb g/LFollicular, Hodgkins onlyAbs lymphocyte count x109/LHodgkins only	Value	
4. DISEASE TRANSFORI If transformed from CLI If not transformed from questions	L, then go directly to S		Lymphocyte % Hodgkins only Serum Albumin g/L Hodgkins only LDH U/L LDH ULN U/L		
Pathology at transformati Transformation date same Yes (concurrent No - Date of tran	-	se Treatment	6. NODAL, ORGAN INVOLVEMENT AT TRAI PET (or PET/CT) positive: Y N Not done Known nodal involvement: Y N	NSFORMATION	
Immunohistochemical sta	ains performed: Y N	Unk	If yes: Total number nodal regions involved		
BCL-2	Pos Neg Unk	% pos	excl follicular: □1 □≥2	□ Unknown 	
BCL-6	Pos Neg Unk	% pos	follicular only: $\Box \ge 5$ $\Box < 5$ Largest nodal mass (max dimensions)	Unknown	
CD5	Pos Neg Unk			_x un	
CD10	Pos Neg Unk		Extranodal / splenic involvement? Y N Unk		
CD30	Pos Neg Unk		Site(s) of involvement:		
C-MYC	Pos Neg Unk	% pos			
Cyclin D1	Pos Neg Unk		Stage of organ involvement: I II III IV	Unknown	
EBER ISH	Pos Neg Unk		B symptoms present 6months prior transform: N	(N Unk	
Ki-67	Pos Neg Unk	% pos		-	

ECOG score: _____ 🗌 Unknown

MUM1

SOX11

Pos | Neg | Unk

Pos | Neg | Unk

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7. DISEASE TREATMENT	8. DLBCL - complete	e if CR not achiev	ved after 1st lir	ne therapy	
Treatment was given after diagnosis: Y N	LDH U/L: LDH ULN, U/L:				
Complete this section as many times as required for each line given Systemic therapy: Y N Date started:/_/ Date stopped:/_/	Stage of organ involvement: I II III IV Unknown				
Number of cycles: Specify regimen/agents:	ECOG score:	🗌 Unknown			
This therapy line given to mobilised cells: Y N	Extranodal or splenic i If yes, site(s) of invo	•	•		
Intrathecal therapy: Y N					
Date started:// Date stopped:// Specify therapy:	9. DISEASE ASSESSMENT AT LAST EVALUATION PRIOR TO PREPARATIVE REGIMEN / INFUSION				
Intraocular therapy : Y N	Were cytogenetics per	rformed: Y N U	Ink		
Prophylaxis Treatment ocular disease Unknown Date started: _/_/_ Date stopped: _/_/ Specify therapy: Date stopped://	 Tested via FISH: Y N Unk Results: Abnormalities identified no abnormalities Specify abnormalities:				
Radiation therapy: Y N Date started: _/_/_ Date stopped: _/_/_ Extent of radiation field: Radiation site(s): Dose per fraction Number fractions: Total dose: □ Gy □ cGy	 Tested via karyotyping: Y N Unk Results: Abnormalities identified No evaluable metaphases abnormalities Specify abnormalities: Report(s) submitted: Y N 				
Technique:				/alue	
Surgery Y N Date of surgery:// Splenectomy: Y N Other site:					
Photopheresis: Y N	Abs lymphocyte count x10°/L Hodgkins only Minimal residual disease:				
Cell therapy: Y N \rightarrow if yes, complete CT form		BM Blood specify other	Pos Neg ND	Date sample	
Best response to line of therapy: (Radiographic criteria) CR PR NR/SD PD Not done Date assessed://	Flow cytometry PCR			_/_/_ _/_/_	
Best response to line of therapy: (Metabolic criteria) CR PR NR/SD PD Not done Date assessed://	NGS, 3 rd gen Pathology report(s) su	hmitted to Registr		_/_/_	
This therapy given as maintenance / consolidation: Y N Relapse/progression occurred after this therapy line: Y N	Known nodal involven (follicular only If yes:	nent: Y N			
If yes, date relapse/progression://	Largest nodal mass (max dimensions) x cm Extranodal or splenic involvement: Y N Unk Site(s) of involvement:				