Practical Flow Cytometry in Haematology 100 Worked Examples

Mike Leach Mark Drummond Allyson Doig Pam McKay Bob Jackson Barbara J. Bain

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WILEY Blackwell

This edition first published 2015 © 2015 by John Wiley & Sons, Ltd

Registered office: John Wiley & Sons, Ltd, The Atrium, Southern Gate, Chichester, West Sussex, PO19 8SQ, UK

Editorial offices: 9600 Garsington Road, Oxford, OX4 2DQ, UK

- The Atrium, Southern Gate, Chichester, West Sussex, PO19 8SQ, UK
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Library of Congress Cataloging-in-Publication Data

Leach, Richard M. (Haematologist), author.

Practical flow cytometry in haematology: 100 worked examples / Mike Leach [and 5 others]. p.; cm. Includes index. ISBN 978-1-118-74703-2 (hardback) I. Title. [DNLM: 1. Hematologic Diseases-diagnosis-Case Reports. 2. Hematologic Neoplasms-diagnosis-Case Reports. 3. Flow Cytometry-methods-Case Reports. 4. Hematology-methods-Case Reports. WH 120] RC636

616.1'5075-dc23

2015007734

A catalogue record for this book is available from the British Library.

Wiley also publishes its books in a variety of electronic formats. Some content that appears in print may not be available in electronic books.

Set in 8.5/11pt, MinionPro by Laserwords Private Limited, Chennai, India

1 2015

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Preface

In our first publication 'Practical Flow Cytometry in Haematology Diagnosis' we presented an outline approach to the use and applications of flow cytometric immunophenotyping in the diagnostic haematology laboratory. We showed how this technique could be used to study blood, bone marrow and tissue fluid samples in a variety of clinical scenarios to achieve a diagnosis, taking into account important features from the clinical history and examination alongside haematology, morphology, biochemistry, immunology, cytogenetic, histopathology and molecular data. This text was illustrated with a series of 'worked examples' from real clinical cases presenting to our institution. These cases have proven to be very popular and so a companion publication dedicated to 100 new 'worked examples' seemed justified and is presented here.

The principles used in the approach to each case are exactly the same as used in the first publication and cases are illustrated with tissue pathology and cytogenetic and molecular data, which are integrated to generate, where appropriate, a diagnosis based on the WHO Classification of Tumours of Haematopoietic and Lymphoid Tissues. We present a spectrum of clinical cases encountered in our department from both adult and paediatric patients and of course, if the title is to be justified, flow cytometry plays a role in every case. Furthermore, we present both neoplastic and reactive disorders and the cases appear in no particular order so that the reader should have no pre-conceived idea as to the nature of the diagnosis in any case. May-Grünwald-Giemsa (MGG)-stained films of peripheral blood and bone marrow aspirates are presented with flow cytometry data alongside haematoxylin and eosin (H&E)-stained bone marrow and tissue biopsy sections. Immunohistochemistry is used to further clarify the tissue lineage and cell differentiation. Cytogenetic studies using metaphase preparations are used to identify translocations and chromosome gains and losses whilst interphase fluorescence in situ hybridisation (FISH) studies and polymerase chain reaction (PCR) are used to identify gene fusions, break-aparts and deletions. The presentation is brought to a conclusion and the particular features that are important in making a diagnosis are highlighted and discussed. The cases

are also listed according to disease classification toward the end (page 390) so that the text can also be used as a reference manual.

The analysis of blood, bone marrow and tissue fluid specimens requires a multi-faceted approach with the integration of scientific data from a number of disciplines. No single discipline can operate in isolation or errors will occur. Flow cytometry technology is in a privileged position in that it can provide rapid analysis of specimens; it is often the first definitive investigation to produce results and help formulate a working diagnosis. The results from flow cytometry can help to structure investigative algorithms to ensure that the appropriate histopathological, cytogenetic and molecular studies are performed in each case. Tissue samples are often limited in volume and difficult to acquire so it is important to stratify investigations accordingly and to get the most from the material available. It is not good scientific or economic practice to run a large series of poorly focussed analyses on every case. Appropriate studies need to be executed in defined circumstances and flow cytometry can guide subsequent investigations in a logical fashion. In some situations a rapid succinct diagnosis can be achieved; immunophenotyping excels in the identification of acute leukaemia. Cytogenetic studies and molecular data give important prognostic information in these patients. But of course the recognised genetic aberrations need to be demonstrated if the diagnosis is to be substantiated. Acute promyelocytic leukaemia can often be confidently diagnosed using morphology alongside immunophenotyping data, but a PML translocation to the RARA fusion partner, needs to be shown. Flow cytometry cannot operate in isolation; despite having the 'first bite of the cherry' the differential diagnosis can still be wide open. There are a good number of worked examples illustrated here where immunophenotyping was not able to indicate a specific diagnosis. The disease entities with anaplastic or 'minimalistic' phenotypes frequently cause difficulty. Appropriate histopathology and FISH, performed on the basis of flow cytometric findings, highlighting abnormal protein expression and gene rearrangement respectively, can make a major contribution to diagnosis and disease classification. Only when a specific

diagnosis is made and prognostic parameters are assessed can the optimal management plan be considered for each individual patient. Finally, the goal posts are constantly moving and developments in the molecular basis of disease, refining disease classification, are evolving rapidly. Whether we are considering eosinophilic proliferations, the myriad of myeloproliferative neoplasms, lymphoproliferative disorders or acute leukaemias we are constantly noting developments and adjusting diagnosis and prognosis accordingly. This is an era of evolving diagnostic challenge and rapid molecular evolution where the practising clinician needs to keep abreast of the significant developments in all areas of haematopathology.

The flow cytometric principles applied to each case have been described in detail in 'Practical Flow Cytometry in Haematology Diagnosis' and some working knowledge is required to interpret the cases described. We also anticipate a reasonable ability in morphological assessment and a capacity to identify morphological variations seen in various disease states. In spite of this we do endeavour to describe the diagnostic logic that we have applied to each worked example and demonstrate how cellular immunophenotypes have helped determine the nature of the disorder.

This text will be of interest to all practicing haematologists and to histopathologists with an interest in haematopathology but it is particularly directed at trainee haematologists and scientists preparing for FRCPath examinations.

Acknowledgement

We are grateful for the substantial assistance of Dr Avril Morris DipRCPath, Principal Clinical Scientist, West of Scotland Genetic Services, Southern General Hospital, Glasgow with regard to the provision of the cytogenetic data and images relevant to the clinical cases presented here.

List of Abbreviations

ADP	adenosine diphosphate	CML	chronic myeloid leukaemia
AITL	angioimmunoblastic T-cell lymphoma	CMML	chronic myelomonocytic leukaemia
AL	acute leukaemia	CMV	cytomegalovirus
ALCL	anaplastic large cell lymphoma	CNS	central nervous system
ALL	acute lymphoblastic leukaemia	CODOX M/IVAC	cyclophosphamide, vincristine,
ALP	alkaline phosphatase		doxorubicin, methotrexate/
ALT	alanine transaminase		ifosphamide, mesna, etoposide,
AML	acute myeloid leukaemia		cytarabine
AML-MRC	acute myeloid leukaemia with	CR	complete remission
	myelodysplasia-related changes	CRAB	calcium (elevated), renal failure,
ANA	antinuclear antibody		anaemia, bone lesions
APC	allophycocyanin	CSF	cerebrospinal fluid
APL	acute promyelocytic leukaemia	СТ	computed tomography
APTT	activated partial thromboplastin time	CTCL	cutaneous T-cell lymphoma
ASM	aggressive systemic mastocytosis	CTD	cyclophosphamide, thalidomide and
AST	aspartate transaminase		dexamethasone
ATLL	adult T-cell leukaemia/lymphoma	CXR	chest X-ray
ATRA	all- <i>trans</i> -retinoic acid	cyt, cyto	cytoplasmic
AUL	acute undifferentiated leukaemia	DEXA scanning	dual energy X-ray absorptiometry
B-ALL	B-lineage acute lymphoblastic		scanning
	leukaemia	DIC	disseminated intravascular
BCLU	B-cell lymphoma, unclassifiable, with		coagulation
	features intermediate between diffuse	DKC	dyskeratosis congenita
	large B-cell lymphoma and Burkitt	DLBCL	diffuse large B-cell lymphoma
	lymphoma	DM	double marking
BEAM	carmustine (BCNU), etoposide,	EBER	EBV-encoded small RNAs
	cytarabine (cytosine arabinoside) and	EBV	Epstein-Barr virus
	melphalan	EBV LMP	Epstein-Barr virus latent membrane
BL	Burkitt lymphoma		protein
BP	blast phase	EDTA	ethylene diamine tetra-acetic acid
BPDCN	blastic plasmacytoid dendritic cell	eGFR	estimated glomerular filtration rate
	neoplasm	EMA	eosin-5-maleimide
c	cytoplasmic	EORTC	European Organization for Research
CD	cluster of differentiation		and Treatment of Cancer
СНОР	cyclophosphamide, doxorubicin,	ESHAP	etoposide, methyl prednisolone,
	vincristine and prednisolone		cytarabine, cisplatin
CLL	chronic lymphocytic leukaemia	ESR	erythrocyte sedimentation rate

ET	essential thrombocythaemia	LPD	lymphoproliferative disorder	
ETP-ALL	early T-cell precursor acute	МСН	mean cell haemoglobin	
	lymphoblastic leukaemia	MCL	mantle cell lymphoma	
FAB	French–American–British (leukaemia	MCV	mean cell volume	
	classification)	MDS	myelodysplastic syndrome/s	
FBC	full blood count	MDS/MPN	myelodysplastic/myeloproliferative	
FDG	fluorodeoxyglucose		neoplasm	
FISH	fluorescence in situ hydridisation	MF	mycosis fungoides	
FITC	fluorescein isothocyanate	MGG	May–Grünwald–Giemsa	
FL	follicular lymphoma	MGUS	monoclonal gammopathy of	
FLAER	fluorescein-conjugated proaereolysin		undetermined significance	
FLAG	fludarabine, cytarabine, granulocyte	MM	multiple myeloma	
	colony-stimulating factor	mod	moderate fluorescence	
FLAG-IDA	fludarabine, cytarabine, granulocyte	MPAL	mixed phenotype acute leukaemia	
	colony-stimulating factor, idarubicin	MPN	myeloproliferative neoplasm	
FSC	forward scatter	MPO	myeloperoxidase	
GGT	gamma glutamyl transferase	MRD	minimal residual disease	
GI	gastrointestinal	MRI	magnetic resonance imaging	
Gp	glycoprotein	MZL	marginal zone lymphoma	
GP	general practitioner	NLPHL	nodular lymphocyte-predominant	
GPI	glycosylphosphatidylinositol		Hodgkin lymphoma	
H&E	haematoxylin and eosin	NOS	not otherwise specified	
HD	haemoglobin concentration	NR	normal range	
HCL	hairy cell leukaemia	PAS	periodic acid-Schiff	
HCL-V	hairy cell leukaemia variant	PCR	polymerase chain reaction	
	human herpesvirus	PD-1	an antigen, programmed death	
	Hadakin lumphoma		1(CD279)	
	human leucocyte antigen DP	PE	phycoerythrin	
	hereditary spherocytosis	PEL	primary effusion lymphoma	
HTIV-1	human T-cell lymphotronic virus-1	PET	positron-emission tomography	
	immunocytochemistry	Ph	Philadelphia (chromosome)	
la	immunoglobulin	PMF	primary myelofibrosis	
laA	immunoglobulin A	PNET	primitive neuroectodermal tumour	
laG	immunoglobulin G	PNH	paroxysmal nocturnal haemoglobinuria	
laM	immunoglobulin M	PRCA	pure red cell aplasia	
інс	immunohistochemistry	PT	prothrombin time	
IPSS	International Prognostic Scoring System	PTCL-NOS	peripheral T-cell lymphoma, not	
ISCL	International Society for Cutaneous		otherwise specified	
	Lymphomas	PTLD	post-transplant lymphoproliferative	
ISH	in situ hybridisation		disorder	
ISM	indolent systemic mastocytosis	PV	polycythaemia vera	
ITD	internal tandem duplication	RBC	red blood cell (count)	
ITP	'idiopathic' (autoimmune)	R-CHOP	rituximab, doxorubicin, vincristine and	
	thrombocytopenia purpura		prednisolone	
IVLBCL	intravascular large B-cell lymphoma	R-CVP	rituximab, cyclophosphamide,	
LAP	leukaemia-associated phenotype		vincristine and prednisolone	
LBL	lymphoblastic lymphoma	RNA	ribonucleic acid	
LDH	lactate dehydrogenase	RQ-PCR	real-time quantitative polymerase chain	
LFTs	liver function tests		reaction	
LGL	large granular lymphocyte	RS	Reed-Sternberg	

RT-PCR	reverse transcriptase polymerase chain	TdT	terminal deoxynucleotidyl transferase
SAA Sig SLE	severe aplastic anaemia surface membrane immunoglobulin systemic lupus erythematosus	TKI T-LBL t-MDS	T-cell intracellular antigen tyrosine kinase inhibitor T-lymphoblastic lymphoma therapy-related myelodysplastic
SM-AHNMD SMILE	systemic mastocytosis systemic mastocytosis with associated clonal haematological non-mast cell disease dexamethasone, methotrexate,	TRAP TT TTP	syndrome tartrate-resistant acid phosphatase thrombin time thrombotic thrombocytopenic purpura
SSC T-ALL TBI	ifosfamide, L-asparaginase and etoposide side scatter T-lineage acute lymphoblastic leukaemia total body irradiation	USS WAS WASp WBC WM	urea, electrolytes and creatinine ultrasound Wiskott—Aldrich syndrome Wiskott—Aldrich syndrome protein white blood cell (count) Waldenström macroglobulinaemia

Technical Notes

The patients presented in 100 Worked Examples were all real cases encountered and investigated in a regional flow cytometry laboratory serving a population of approximately 2.5 million over a period of 18 months. These are individually presented with a history that reflects the actual events for each patient, commencing with the presenting clinical features and the initial basic laboratory tests and then proceeding to flow cytometry, bone marrow aspirate morphology, bone marrow trephine biopsy histology with immunohistochemistry studies and other specialised cytogenetic and molecular analyses.

Full blood counts

The full blood counts and marrow counts (for appropriate dilutions in relation to antibody) were performed on a Sysmex XN analyser. The differential leucocyte counts are automated counts from the analyser. It should be noted that sometimes, in an automated count, abnormal cells are misidentified and the leucocyte sub-populations differ from a manual differential performed on a blood film. Such misidentifications are indicated by inverted commas.

Biochemistry and immunology studies

All relevant biochemistry and immunology data is given in relation to the context of each patient presentation and in terms of investigations that were thought to be relevant to the case as the clinical diagnosis evolved. Some retrospectively relevant data may be missing but this reflects the true nature of these actual patient scenarios and the investigations that were considered necessary at that time. Serum calcium values given are all corrected in accordance with serum albumin level.

Flow cytometry analysis

Flow cytometry studies were all performed using a Becton Dickinson FACS Canto II analyser. The findings are presented as a list of positive and negative results in relation to the antigen and target cell population and the gating strategies applied to each case are explained. A series of scatter plots and histograms are presented to illustrate specific informative points. The expression of most membrane antigens is graded as positive when more than 20% of gated events are positive; the exceptions being CD34, CD117 and cytoplasmic antigens where a threshold of 10% has been used. Where the percentage positivity for a given membrane antigen in the gated target population is borderline positive so that some cells appear negative and some positive we have used the term 'partial' to describe antigen expression. Cytoplasmic expression of an antigen is indicated with the prefix 'c' (cytoplasmic expression of CD3 being cCD3) but on some scatter plots 'cyt' or 'cyto' has been used. The intensity of antigen expression in terms of median fluorescence intensity is graded as dim, moderate or bright compared to our laboratory reference ranges for normal cells of each relevant lineage. See Figures 1.1a-g for a schematic representation of these principles.



Figure 1.1 Visual representation of strength of fluorescence in flow cytometry (not actual patient specimens), showing an isotype control and eight CD19-positive samples which show fluorescence intensity with CD20 varying from negative to bright. (a) Isotype control, used to set thresholds. (b) Negative (consistent with a CD19-positive, CD20-negative B-cell precursor neoplasm). (c) Partial positive, indicating that CD20 antigen expression varies from negative to positive (consistent with a precursor B-cell neoplasm). (c) Partial positive, indicating that CD20 antigen expression varies from negative to positive (consistent with a precursor B-cell neoplasm). (c) adjusted) Indicating that the threshold for positivity might be reduced by the cytometrist where a discrete dim positive population is identified. (d) Dim CD20 antigen expression (consistent with chronic lymphocytic leukaemia). (e) Moderate intensity, indicating medium strength of CD20 antigen expression (consistent with B-cell non-Hodgkin lymphoma). (f) bright, indicating strong CD20 antigen expression (consistent with hairy cell leukaemia). (g) Two distinct populations, one partial and dim and one bright (could indicate two unrelated B-lineage neoplasms or transformation of a low grade lymphoma). (h) Contrasting with (g), a heterogeneous single population with fluorescence intensity varying from negative to moderate with a minority being bright.



Figure 1.1 (Continued)

Immunohistochemistry in paraffin-embedded formalin fixed tissue

In the following section a list is presented of the immunohistochemical reagents used in assessing the paraffin embedded material (bone marrow trephine and lymph node biopsies) in the worked examples described. It should be pointed out that specificities and sensitivities may differ from the antibodies used in flow cytometry due to the effects of formalin fixation and decalcification resulting in antigen loss or masking. For example, CD5 may be detected by flow cytometry in a peripheral blood B-cell lymphocytosis but immunocytochemistry may on occasion be negative for the same marker in the trephine specimen. CD56 is aberrantly expressed by plasma cells in myeloma yet immunoreactivity for this antibody within plasma cells in paraffin sections is seen in only a minority of cases. The opposite situation may also occur where an antigen such as TdT is strongly positive by immunohistochemistry on the fixed tissue but is negative on the flow sample. Reticulin fibrosis is reported as per the WHO classification as grade 0, 1, 2 or 3.

These specific features of different techniques need to be appreciated when formulating the combined pathology report and an understanding of the strengths and weaknesses of each approach is essential when establishing a final diagnosis. Cytogenetic and molecular studies have a major influence on disease classification. Specific findings can carry diagnostic significance way in excess of any other single investigative modality e.g. *BCR-ABL1*, *PML-RARA*, *FIP1L1-PDGRFA*. Metaphase cytogenetic studies not infrequently fail, either reflecting the quality of the specimen or the disease entity being studied. Informed FISH and PCR studies can carry great diagnostic importance in certain clinical circumstances and molecular diagnostics will continue to inform disease classification with increasing power and specificity over the decades ahead.

Laboratory Values

Abbreviations and Normal Ranges.

Blood		Blood		
Haematology		– Urate	0.2-0.43 mmol/L	
Haemoglobin concentration (Hb)	130–180 g/L (M)	Lactate	<2.4 mmol/L	
-	125–170 g/L (F)	Lactate dehydrogenase (LDH)	80-240 U/L	
Mean cell volume (MCV)	80–100 fl	Aspartate transaminase (AST)	<40 U/L	
Reticulocyte count	50-100 × 10 ⁹ /L	Alanine transaminase (ALT)	<50 U/L	
White blood cell count (WBC)	$4-11 \times 10^{9}$ /L	Gamma glutamyl transferase (GGT)	<70 U/L	
Neutrophils	2-7×10 ⁹ /L	Alkaline phosphatase (ALP)	40-150 U/L	
Lymphocytes	1.5-4 × 10 ⁹ /L	Calcium adjusted	2.1-2.6 mmol/L	
Monocytes	0.2-0.8 × 10 ⁹ /L	Phosphate	0.7–1.4 mmol/L	
Eosinophils	0.04-0.4 × 10 ⁹ /L	C-reactive protein (CRP)	<10 mg/L	
Basophils	0.01-0.1 × 10 ⁹ /L	Bilirubin	<20 µmol/L	
		Albumin	32–45 g/L	
Haematinics		Globulins	23-38 g/L	
Serum ferritin	10-275 ng/mL			
Serum folate	3.1–20 ng/mL	Serum osmolality	270–295 mmol/kg	
Serum vitamin B ₁₂	200–900 pg/mL			
12		Urine protein/creatinine ratio	0–15 mg/mmol	
Coagulation				
Prothrombin time (PT)	9–13 s	Immunoglobulin G (IgG)	6–16 g/L	
Activated partial thromboplastin time	27–38 s	Immunoglobulin A (IgA)	0.8–4.0 g/L	
(APTT)		Immunoglobulin M (IgM)	0.5-2.0 g/L	
Thrombin time (TT)	11–15 s			
Fibrinogen	1.5–4 g/L	Serum free light chains		
D dimer	0–243 ng/mL	Free kappa	3.3–19.4 mg/L	
	-	Free lambda	5.7–26.3 mg/L	
Biochemistry				
Sodium (Na)	135–145 mmol/L	Cerebrospinal fluid (CSF)		
Potassium (K)	3.5-5.0 mmol/L	Protein	<0.4 g/L	
Urea	2.5–7.5 mmol/L	Cells	<0.001 × 10 ⁹ /L (<10	
Creatinine	40–130 µmol/L		cells/µL)	
Bicarbonate	20–30 mmol/L		-	
		- Glucose	2 mmol/L less than serum glucose	

1

Case 1

An 11-year-old boy was admitted with a short history of fever, sweats, dyspnoea and left chest discomfort. There was no past history of note. Examination identified features of a left pleural effusion. There was also a tender swelling of the left anterior chest in the upper pectoral region and palpable cervical lymphadenopathy. The liver and spleen were not palpable.

Laboratory investigations

FBC and blood film: normal U&Es, LFTs: normal. LDH was 1460 U/L.





Imaging

The CXR showed opacification and loss of aeration of the left hemithorax in keeping with a pleural effusion (Figure 1.1).



Figure 1.1 CXR.

CT imaging confirmed this but in addition identified a left pleural-based mass, abnormal soft tissue in the left pectoral muscles (arrows, Figure 1.2) and cervical lymphadenopathy. In addition, there was collapse/consolidation of the lower left lung, creating the appearance of an air bronchogram. A core biopsy of a cervical node was taken and the pleural effusion was aspirated for analysis.

Flow cytometry

The pleural fluid cell count was 0.98×10^9 /L. A cytospin preparation showed three distinct cell types: a small mature lymphoid population in keeping with reactive lymphocytes, an intermediate sized/large sized lymphoid population and a large cell population with pleomorphic morphology and blue cytoplasm (Figures 1.3–1.6). The cells with the abundant cytoplasm (Figures 1.3 and 1.4) and the single binucleate cell (Figure 1.6) are reactive mesothelial cells. The cells with the cytoplasmic blebs (Figures 1.4–1.6) are the disease cells,

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Figure 1.3 MGG, ×500.



Figure 1.4 MGG, ×500.

which were the subsequent focus for immunophenotyping studies.

By applying a blast gate to the suspected malignant cells in the FSC/SSC analysis (Figure 1.7), they were shown to express CD45^{bright} (Figure 1.8), CD2 (Figure 1.9), cCD3 [whilst surface CD3 was negative apart from a few reactive

Figure 1.5 MGG, ×500.



Figure 1.6 MGG, ×500.

T cells (Figure 1.8)], partial CD7 (Figure 1.10) and CD13. Other T-lineage markers were negative.

This is therefore a T-lymphoid neoplasm, indicated by positivity for cCD3 expression, with limited lineage-specific



Figure 1.7 FSC/SSC.





Figure 1.9 CD2/CD19.



Figure 1.8 CD3/CD45.

markers and an aberrant myeloid marker. The tumour has medium sized/large cell morphology. It was showing aggressive clinical behaviour with extranodal tissue invasion in this 11-year-old patient. An anaplastic large cell lymphoma had to be considered and the medium sized/large cells in the pleural fluid were shown to be strongly expressing CD30 (not shown).

Histopathology

An H&E-stained core biopsy of a cervical node is shown in Figure 1.11. The node is replaced by an infiltrate of

Figure 1.10 CD7/CD16.

undifferentiated pleomorphic large cells with prominent nucleoli.

Immunohistochemistry showed the large cells to express CD45, epithelial membrane antigen (EMA), CD2 focally (Figure 1.12), CD7, granzyme B and CD30 (Figure 1.13). In addition, there was strong nuclear and cytoplasmic staining for anaplastic lymphoma kinase (ALK) protein (Figure 1.14).

The CD30 staining was particularly useful in demonstrating lymphatic invasion within the capsule of the node (Figure 1.15).



Figure 1.11 H&E, ×400.



Figure 1.13 CD30, ×400.



Figure 1.12 CD2, ×400.

FISH studies

A t(2;5)(p23;q35) translocation, rearranging the *ALK* and *NPM1* (nucleophosmin) genes, was shown by FISH studies on paraffin-embedded lymph node tissue. The presence of this specific translocation is highly associated with both nuclear and cytoplasmic positivity for ALK.



Figure 1.14 ALK, ×400.

Discussion

Anaplastic large cell lymphoma (ALCL) is an aggressive mature T-cell neoplasm with pleomorphic, often large cell, morphology. It frequently fails to show surface expression of T-lineage-specific markers and to potentially further mislead may express aberrant myeloid antigens. This is an



Figure 1.15 CD30, ×100.

important condition to recognise; it frequently shows rapid progression with extranodal tissue involvement and it can rarely appear in the blood. Treatment of ALK⁺ ALCL is usually rewarding, particularly in paediatric patients, with prompt response to chemotherapy and frequent durable remissions.

Final diagnosis

Anaplastic large cell lymphoma (ALK⁺)

2

Case 2

A 72-year-old woman presented with a few months' history of fatigue and the more recent onset of breathlessness and night sweats. On clinical examination she had a large right-sided pleural effusion but no palpable lymphadenopathy.

Laboratory results

FBC: Hb 158 g/L, WBC 16.6 \times 10⁹/L (neutrophilia and monocytosis) and platelets 502×10^9 /L.

U&Es: normal. LFTs were mildly deranged (ALT 52 U/L, alkaline phosphatase 173 U/L). Albumin was low at 29 g/L and serum LDH was raised at 584 U/L.

Imaging

A CT scan demonstrated a large right-sided pleural effusion with collapse of the right middle and lower lobes and partial collapse of the upper lobe (Figure 2.1). In addition, there were large volume, confluent, necrotic nodal masses in the right hilar, mediastinal, retrocrural, paracardiac and para-aortic areas (not shown) as well as pleural deposits (arrow, Figure 2.1).

Pleural fluid biochemistry and cytology

The pleural fluid LDH was markedly elevated at 2171 U/L with relatively low glucose at 7.2 mmol/L (patient diabetic) and protein of 43 g/L.



Figure 2.1 CT.

Microscopy of the pleural fluid showed lymphoid cells admixed with neutrophils, histiocytes and mesothelial cells. Most of the lymphoid cells were small but an admixed population of medium-sized cells with slightly irregular nuclei was also present. On morphology alone, the lymphoid cells were thought likely to be reactive but the reporting pathologist suggested that a fresh pleural fluid specimen should be assessed using flow cytometry.

Morphology (pleural fluid)

A specimen of pleural fluid was received by our laboratory. The WBC was found to be 6.3×10^9 /L. A cytospin preparation showed a cellular specimen with notable macrophages, neutrophils and small lymphocytes. In addition, some large blastoid lymphoid cells were seen (Figures 2.2–2.5).

Practical Flow Cytometry in Haematology: 100 Worked Examples, First Edition. Mike Leach,

Mark Drummond, Allyson Doig, Pam McKay, Bob Jackson and Barbara J. Bain.

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Figure 2.2 MGG, ×500.



Figure 2.4 MGG, ×500.



Figure 2.3 MGG, ×500.

Flow cytometry (pleural fluid)

The FSC/SSC plot shows the orientation of the different populations of cells described above. The small lymphoid population (Figure 2.6) comprised mainly reactive T cells (black events with mixture of CD4⁺ and CD8⁺ cells) and normal B cells (blue events). Seventeen per cent of all leucocytes were CD19⁺ B cells (Figure 2.7) showing a mature pan-B phenotype with CD20 positivity and a hint of monoclonality (kappa



Figure 2.5 MGG, ×500.

73%, lambda 18%). The SSC analysis in Figure 2.7, however, defines 2 B-cell populations with different scatter characteristics, populations P1 and P2.

By gating on the higher SSC profile B cells (P2 red events, Figure 2.7, which are larger indicated by high FSC in Figure 2.6), a clear clonal population was demonstrated showing strong CD20 positivity, expression of CD10/HLA-DR (Figure 2.8), CD38, FMC7, CD79b and CD22 with kappa light chain restriction (Figure 2.9).



Figure 2.6 FSC/SSC.



Figure 2.7 CD19/SSC.

Note the phenotype of the blue events, population P1, representing residual small polyclonal reactive B cells.

Lymph node biopsy

A CT-guided core biopsy of a paravertebral node showed lymphoid infiltration by predominantly small centrocytic cells with occasional larger centroblasts. The cells were positive for CD20, CD10, BCL6 and BCL2 and negative for CD3, CD5, cyclin D1, CD23, CD43 and CD21. The proliferation fraction was low (~10%). The histological and immunohistochemical appearances were in keeping with follicular lymphoma, grade 2, with no evidence of high-grade



Figure 2.8 CD10/HLA-DR.



Figure 2.9 Kappa/lambda.

transformation. FISH for the t(14;18) translocation was positive.

Bone marrow aspirate and trephine biopsy

These were both normal.

Discussion

The clinical presentation, imaging, pleural fluid morphology and flow cytometry were most in keeping with a diagnosis of an aggressive, mature CD10⁺ B-cell neoplasm. Diffuse large