

Novel somatic alterations in unicentric and idiopathic multicentric Castleman disease

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Objectives: Castleman disease (CD) is a heterogeneous group of disorders involving systemic inflammation and lymphoproliferation. Recently, clonal mutations have been identified in unicentric CD (UCD) and idiopathic multicentric CD (iMCD), suggesting a potential underlying neoplastic process.

Methods: Patients with UCD or iMCD with next generation sequencing (NGS) data on tissue DNA and/or circulating tumor DNA (ctDNA) were included.

Results: Five patients were included, 4 with iMCD and 1 with UCD. Four patients (80%) were women; median age was 40 years. Three of five patients (60%) had ≥ 1 clonal mutation detected on biopsy among the genes included in the panel. One patient with iMCD had a 14q32-1p35 rearrangement and a der(1)dup(1)(q42q21)del(1)(q42) (1q21 being IL-6R locus) on karyotype. This patient also had a *NF1* K2459fs alteration on ctDNA (0.3%). Another patient with iMCD had a *KDM5C* Q836* mutation, and one patient with UCD had a *TNS3-ALK* fusion but no ALK expression by immunohistochemistry.

Conclusions: We report 4 novel somatic alterations found in patients with UCD or iMCD. The 1q21 locus contains IL-6R, and duplication of this locus may increase IL-6 expression. These findings suggest that a clonal process may be responsible for the inflammatory phenotype in some patients with UCD and iMCD.

Supplemental Table 1. Patient information and fulfillment of the diagnostic criteria¹

Patient	1	2	3	4	5	Reference range
Age [†] /Sex	28.7/F	30.4/F	58.8/F	39.7/M	39.8/F	
Diagnosis	iMCD	iMCD	iMCD	iMCD	UCD	
Pathology	Hyaline vascular	Plasmacytic	Plasmacytic	Plasmacytic	Hyaline vascular	
Slides Reviewed at UCSD [‡]	No (Stanford)	Yes	Yes	No	No	
Biopsy site	Right axillary LN	Left inguinal LN	Submental LN	Inguinal LN	Supraclavicular LN	
HHV-8 IHC/PCR	Neg	Neg	Neg	Neg	Neg	Neg
Imaging	PET/CT with cervical, axillary, pericardiac, retroperitoneal, pelvic, and mesentery LAD	CT with mediastinal, abdominal, and pelvic LAD	PET/CT with cervical, mediastinal, and abdominal LAD	CT with bilateral axillary and inguinal LAD	PET/CT with no LAD	
CRP (mg/L) [†]	ND	88	ND	ND	0.1	<0.5
ESR (mm/hr)	100	N/A	84	106	6	0-20
Hgb (g/dL)	9.5	10.2	12.9	11.9	13	13.6-17.0
Platelet (1000/mm ³)	387	468	487	273	295	140-400
Albumin (g/dL)	2.2	4.3	3.9	2.8	4.6	4.0-
eGFR (mL/min)	ESRD	ESRD	>60	>60	>60	
IgG (mg/dL)	1542	1140	2969	4880	1078	700-1600
B symptoms	N/A	N/A	No	No	No	
Hepatomegaly splenomegaly	Yes	No	No	No	No	
Edema	Yes	Yes	No	No	No	
Cherry hemangioma /violaceous rash	No	No	Yes	No	No	
Lymphocytic interstitial pneumonitis	No	No	No	Possibly [§]	No	
IL-6 (pg/mL)	41	ND	20	13	ND	≤5

Abbreviations: CRP: C-reactive protein; CT: computed tomography; eGFR; estimated glomerular filtration rate; ESR: erythrocyte sedimentation rate; ESRD: end stage renal disease; F: female; Hgb: hemoglobin; HHV-8: Human herpesvirus-8; IHC: immunohistochemistry; IL-6: interleukin-6; iMCD: idiopathic multicentric Castleman disease; LAD: lymphadenopathy; LN: lymph node; N/A: not available; Neg: negative; ND: not done; PCR: polymerase chain reaction; PET: positron emission tomography; UCD: unicentric Castleman disease; UCSD: University of California San Diego

[†]At diagnosis

[‡]Pathology reports were reviewed by a hematopathologist (HYW) when slides were unavailable for review

[§]Not biopsy confirmed

Supplemental Table 2. Summary of reported genetic alterations in iMCD and UCD

Study	N [†]	Method	Genetic alteration [‡]	Somatic vs germline	VAF	Proposed mechanism
iMCD						
Nakamura et al.(24) 1993	1	Karyotype	t(7;14)(p22;q22)	Somatic	NR	<i>IL6</i> gene is located on 7p21-22 and translocation potentially resulted in high level of IL-6
Kone-Paut et al.(21) 2009	1	Single gene targeted sequencing	<i>MEFV</i>	Germline	NR	Reduced activity of pyrin in white blood cells by <i>MEFV</i> mutation causing uncontrolled inflammation
Patel et al.(25) 2017 [¶]	1	Targeted NGS	<i>JAK1</i> V310I	Somatic	NR	Predicted that <i>JAK1</i> is more readily activated through <i>IL-6R</i> due to loss of inhibitory interactions.
You et al.(30) 2019	22	WES	<i>NCOA4</i> L261F [§] (4)	Somatic	NR	NCOA4 may influence the MAPK/AR-ARA signaling pathway
			<i>DARS2</i> [§]			
			<i>MTCL1</i> [§]			
			<i>RABEP1</i> [§]			
		<i>DNAH11</i> [§]				
Yoshimi et al.(29) 2020	3	Targeted NGS	<i>MEK2</i> P128L	Somatic	10%	Activate MAPK signaling. In vitro, cells demonstrated hypersensitive growth with low IL-3. These cells were also sensitive to MEK inhibitors.
			<i>RUNX1</i> G60C	Germline	49%	Predicted to affect heterodimerization of <i>RUNX1</i> with <i>CBFβ</i> , which is essential for its function. In vitro, bone marrow progenitors had enhanced self-renewal capacity.
UCD						
Pauwels et al.(26) 2000	1	Karyotype	t(1;16)(p11;p11), del(7)(q21q22), del(8)(q12q22) [15/20]	Somatic	NR	Clonal proliferation of stromal elements contribute to pathophysiology.
Cokalaere et al.(20) 2002	1	Karyotype	add(1)(q21), der(6)t(6;12)(q23;q15), add(7)(p22), -9,inv(9)(p11q13), del(12)q(15), +mar	Somatic	NR	Results in rearrangement of <i>HMGIC</i> gene which was found in anti-CD21 reactive follicular dendritic cells.
Chen et al.(19) 2006	2	Karyotype	t(1;22)(p22;q13) [18/20]	Somatic	NR	Genetic changes attributed to stromal cells.
			t(7;8)(q37.3;q12) [1/20]			
Reichard et al.(27) 2011	1	Karyotype	add(6)(p23),add(7)(p15), del(7)(p15),add9(q22) [4] inv(9)(p13q22) [2] -3,+tr [2]	Somatic	NR	Genetic changes were attributed to non-lymphoid cells, most likely stromal, dendritic, or endothelial origin.
Chang et al.(18) 2014	29	Single gene PCR	<i>HUMARA</i> (22)	Somatic	NR	Stromal cell clonal proliferation
Legras et al.(22) 2017	9	Targeted NGS	<i>VHL</i> F119L	Somatic	35%	Loss of tumor suppressor activity
			<i>JAK3</i> V718L	Germline	53%	Activation of STAT, AKT, ERK pathways which lead to proliferation

iMCD and UCD						
Stone et al.(28) 2012	58	PCR of selected SNPs of <i>IL6</i> promoter, <i>IL6R</i> , and <i>gp130</i>	<i>IL6R</i> rs4537545 [°] <i>IL6R</i> rs2228145 [°]	Germline	NR	Increased level of soluble IL6R in the serum leads to increased level of IL-6 signaling
Baker et al.(17) 2018	2	WGS	<i>FAS</i> R68G (2)	Germline	NR	Defective Fas-induced lymphocyte apoptosis
Nagy et al.(23) 2018	18	Targeted NGS and WES for FDSC patients	<i>DNMT3A</i> L295Q [*]	Somatic	9%	Disrupts PWWP domain, the DNA binding site, of DNMT3A, resulting in increased proliferation of these cells that produce IL-6.
Li et al.(6) 2018	75	WES Targeted sequencing of <i>PDGFRB</i> p.N666S	<i>PDGFRB</i> [^] (7)	Somatic	NR	Activate KIT kinase through ligand-independent autophosphorylation

Abbreviations: AKT: AKT serine/threonine kinase; AR: androgen receptor; ARA: androgen-associated receptor; DNMT3A: DNA methyltransferase 3 alpha; ERK: extracellular signal-regulated kinases; FAS: Fas cell surface death receptor; FDSC: follicular dendritic cell sarcoma; HMGIC: high-mobility group protein isoform C; HUMARA: human androgen receptor α ; IL-3: interleukin-3; IL-6: interleukin-6; IL-6R: interleukin-6 receptor; iMCD: idiopathic multicentric Castleman disease; JAK: Janus kinase; KIT: KIT proto-oncogene receptor tyrosine kinase; MAPK: mitogen activated protein kinase; MEK: mitogen activated protein kinase kinase; NCOA4: nuclear receptor coactivator 4; NGS: next generation sequencing; NR: not reported; PCR: polymerase chain reaction; PDGFRB: platelet-derived growth factor receptor b; RUNX1: runt related transcription factor 1; SNP: single nucleotide polymorphism; STAT: signal transducer and activator of transcription; UCD: unicentric Castleman disease; VAF: variant allele frequency; VHL: von Hippel-Lindau; WES: whole exome sequencing; WGS: whole genome sequencing; FDSC:

[†]Number of patients with CD who were sequenced

[‡]Reported genetic alterations are single patients among those who are tested except for indicated in parentheses

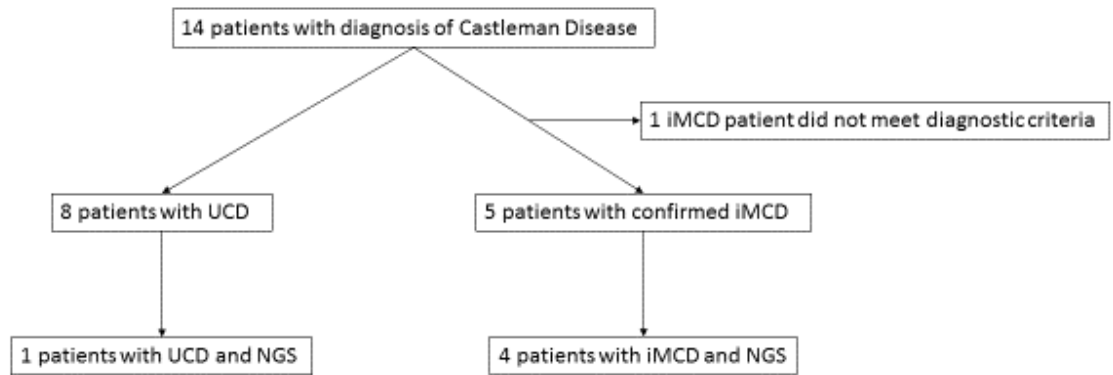
[§]Representative list mutations that are claimed to be significantly associated with adverse prognosis

[¶]Patient did not meet strict diagnostic criteria for iMCD upon further review

[°]Allele frequency for *IL6R* rs4537545 was 0.49 and *IL6R* rs2228145 was 0.42, statistically significantly higher compared to healthy controls

^{*}Several copy number variants were also found. *DNMT3A* mutation was found in patient with iMCD

[^]Other mutations were also identified as part of NGS. *PDGFRB* mutation was seen in 7 of 41 UCD patients and none of the iMCD patients



Supplemental Figure 1. CONSORT diagram

Abbreviations: iMCD: idiopathic multicentric Castleman disease; NGS: next generation sequencing; UCD: unicentric Castleman disease.