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Immune responses and therapeutic challenges in paediatric patients with new-onset acute myeloid leukaemia and concomitant COVID-19

Acute myeloid leukaemia (AML) is a medical emergency often presenting with hyperleucocytosis, coagulopathy and pulmonary infiltration necessitating emergent initiation of therapy. AML with concomitant severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2) infection presents a unique challenge given the lack of evidence-based guidelines or historical experience. While cohort studies have shown early serological responses to SARS-CoV-2 in healthy adults,^{1,2} little is known about the serological responses to infection in patients with AML and the impact of chemotherapy on this response. In the present study, we detail the clinical presentations, treatments, serological and virological responses, and outcomes of two adolescents who presented with AML and concurrent coronavirus disease 2019 (COVID-19).

Two adolescents presenting with AML and COVID-19 were enrolled on an Institutional Review Board-approved protocol to collect prospective/residual specimens used for SARS-CoV-2 serological and virological testing. Nasopharyngeal (NP) real-time reverse transcription polymerase chain reaction (RT-PCR), antibody testing by enzyme-linked immunosorbent assay (ELISA), live-virus focus reduction neutralisation assay-mNG, surface plasmon resonance (SPR) assay and viral genetic sequencing were performed (Data S1). Patient details, treatment and outcome data were abstracted from medical records.

Patient 1 was a 16-year-old Caucasian male with a history of classical Hodgkin lymphoma who presented with fever, cough and hyperleucocytosis [white blood count (WBC) $176 \times 10^9/l$]. His NP SARS-CoV-2 RT-PCR test was positive and peripheral blood flow cytometry (PBFC) confirmed a diagnosis of therapy-related AML. He received hydroxyurea, followed by cytarabine starting on hospital day (HD) 3. Treatment for COVID-19 included hydroxychloroquine, remdesivir and supplemental oxygen (Fig 1).

On HD4, the patient had detectable immunoglobulin (Ig) M and IgG antibodies to SARS-CoV-2 (Table I). The patient did not become lymphopenic throughout his COVID-19 course and maintained detectable binding and neutralising antibodies to SARS-CoV-2. SPR demonstrated that binding antibodies to the pre-fusion conformation of S, the receptor-binding domain (RBD) and S2 subunits all peaked by HD26. The patient had detectable IgM, IgA and IgG1 titres in the final sample on HD70 (Figure S1). He cleared the virus on HD16. Bone marrow (BM) on HD20 showed rare blasts on morphology in the setting of pancytopenia, with no disease detected by flow cytometry. He received additional chemotherapy with azacitidine and gemtuzumab starting on HD26. His treatment complications included bacteraemia and perirectal abscess with *Pseudomonas aeruginosa* and Epstein-Barr virus (EBV) viraemia resulting in multi-organ failure and death on HD74.

Patient 2 was a 19-year-old obese (body mass index of 32 kg/m²), Hispanic male who presented with fever, cough, dyspnoea and hyperleucocytosis (WBC count 67 × 10⁹/l). His NP SARS-CoV-2 RT-PCR test upon admission was positive and PBFC confirmed a diagnosis of AML. He began induction chemotherapy with cytarabine, daunorubicin and etoposide (ADE) on HD2 and he clinically deteriorated on HD4 requiring intubation and mechanical ventilation. For treatment of COVID-19, he received convalescent plasma (CP), remdesivir, tocilizumab and therapeutic plasma exchange (Fig 1).

The patient was lymphopenic by HD5 and showed no immune response to SARS-CoV-2 with absence of IgM antibodies, waning IgG (post CP) and undetectable neutralisation titres (Table I). Concurrently, the patient's NP RT-PCR remained positive with low cycle-threshold (Ct) values. Coinciding with haematological recovery on HD25, the patient demonstrated a serological response with rising IgM, IgG, neutralising antibody titres and SARS-CoV-2 RT-PCR Ct values until his first negative NP RT-PCR result on HD48. His SPR antibody profiling demonstrated class switching (Figure S1) and coincided with robust increases in binding and neutralising titres with haematological recovery. Genetic sequencing and phylogenetic analysis of his SARS-CoV-2 virus from saliva on HD23 indicated that the sequence clustered with clade 20B,³ which was relatively uncommon in Georgia at the time (Figure S2). A total of 40 intra-host single nucleotide variants (iSNVs) were identified with little variability in S and envelope genes (Figure S3 and Table SI). On HD25 his bone marrow evaluation demonstrated complete remission, but he remained critically ill and on HD48 he suffered a fatal sudden cardiac arrest secondary to sepsis.

In the present report, we highlight both the therapeutic challenges in treating COVID-19 and AML concomitantly and describe the immune response in the setting of myelo-suppressive chemotherapy. The viral clearance of the two

patients negatively correlated with the intensity of chemotherapy given for each and likely contributed to the overall severity of COVID-19. Paediatric recommendations for the management of AML and COVID-19 are not well defined. Use of a 'mild' induction regimen (MAG) in nine patients with AML and COVID-19 in Brazil was recently described to have excellent outcomes.⁴ While reduced-intensity regimens are a valid approach, they preclude response-based risk assignment in AML but warrant further investigation in the setting of concomitant COVID-19.

SARS-CoV-2 evolution within immunocompromised patients with prolonged virus replication has been described.^{5,6} Although our investigation was limited to one time point, we did observe within-sample SARS-CoV-2 variants for Patient 2. The functional importance of these variants is unclear; however, future studies into the intra-host variation of SARS-CoV-2 genotypes are needed in the immunocompromised population. Furthermore, Ct values associated with RT-PCR testing for SARS-CoV-2 could provide indirect assessment of viral load.⁷ Lower median Ct values in adults with cancer correlated with higher rates of mortality.⁸ Patient 2's Ct values remained <25 until HD17 suggesting a high viral load. While PCR Ct values have limitations based on sample collection and instrumentation, serial measurements could be used for clinical correlation.⁹

Limitations of our present study include the small number of patients, which limits generalisability of the data. Co-existing factors such as underlying obesity and race may have contributed to a worse outcome in Patient 2.^{10,11} In addition, absence of serological immunity at presentation and administration of CP in Patient 2 confounds interpretation of serological results. Investigation into early immune responses is warranted in this population as they may serve as prognostic indicators at the time of presentation.

In conclusion, our experience demonstrates that patients with AML and COVID-19 can mount immune responses to

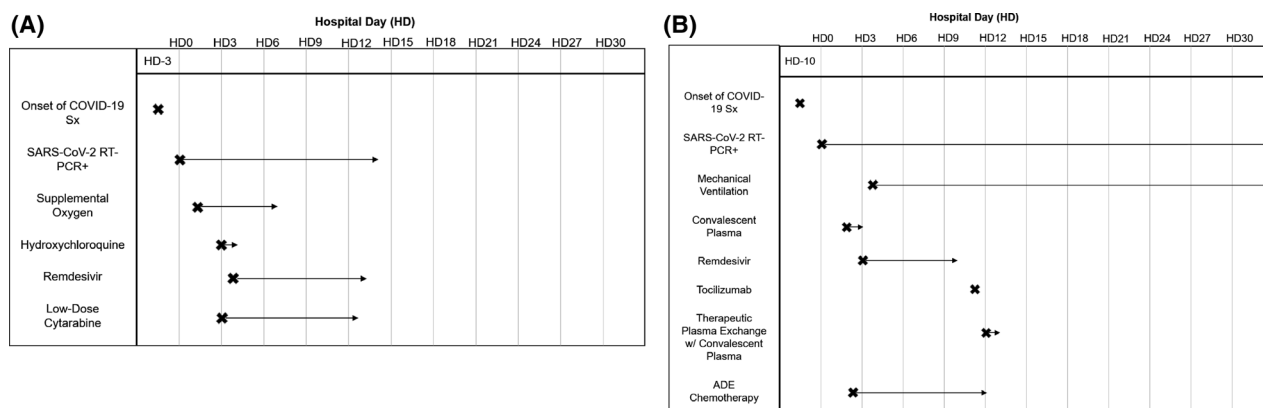


Fig 1. Significant clinical events in first 30 days of hospitalisation for Patient 1 (A) and Patient 2 (B). X, day of first positive sample or first administration; Arrow, duration of positivity or administration; RT-PCR, reverse transcription polymerase chain reaction; Sx, symptoms; ADE, cytarabine, daunorubicin, etoposide.

Table 1. Laboratory results for Patient 1 (A) and Patient 2 (B).

Days from COVID-19 symptom onset		Hospital day	Absolute lymphocyte count, / μ l	IgM antibody titre	IgG antibody titre	SARS-CoV-2 NeutAb	SARS-CoV-2 RT-PCR		
(A)									
3	0	0	1763*	N/A	N/A	N/A	Positive		
7	4	4	696*	804	1327	236	N/A		
19	16	16	590	720	3436	193	Negative		
29	26	26	880	467	4790	91	Negative		
37	34	34	510	312	5621	85	N/A		
45	42	42	770	179	3781	37	N/A		
Days from COVID-19 symptom onset		Hospital day	Absolute lymphocyte count, / μ l	IgM antibody titre	IgG antibody titre	SARS-CoV-2 NeutAb	SARS-CoV-2 RT-PCR	S protein Ct value	ORFIab Ct value
(B)									
10	0	4496†	–	–	–	–	Positive	13.1	13.9
14	4	1330†	85‡	619§	10‡	10‡	Positive	17.7	18.2
22	12	0	85	487	10	10	Positive	N/A	N/A
27	17	0	85	200¶	10	10	Positive	14.6	15.5
35	25	770	3263	1349	559	559	Positive	29.2	36.9
41	31	1535	33562	26804	2763	2763	Positive**	24.8	24.7
45	35	1506	33326	41520	4379	4379	Positive††	33.4	32.4
58	48	1310	N/A	N/A	N/A	N/A	Negative	N/A	N/A

NeutAb, neutralising antibody; RT-PCR, reverse transcription polymerase chain reaction; Ct, cycle threshold; N/A, not available.

*->95% peripheral myeloblasts.

†->85% peripheral myeloblasts 90%.

‡Negative titre.

§Received convalescent plasma on hospital day 2 and 3.

¶Received convalescent plasma on hospital day 12 and 13.

**RT-PCR done on hospital day 33.

††RT-PCR done on hospital day 40.

SARS-CoV-2 even in the face of immune suppression by chemotherapy; however, the intensity of chemotherapy may play a role in the response. Longitudinal research is needed in this vulnerable population to better understand response to SARS-CoV-2 infection and now vaccination.

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Author contributions

Pratik A. Patel designed the research study, performed the research, analysed the data and helped write the paper. Gabrielle Grubbs, Stacey A. Lapp and Venkata V. Edara performed the research and analysed the data. Christina A. Rostad, Evan J. Anderson and Surender Khurana designed the research study, analysed the data and helped write the paper. Claire L. Stokes and Melinda G. Pauly helped write the paper. Anne Piantadosi and Mehul S. Suthar performed the research, analysed the data and helped write the paper. Himalee S. Sabnis analysed the data and helped write the paper. All authors made substantial contributions to research design, or acquisition, analysis or interpretation of data, drafting the paper or revising it critically and approval of the submitted and final versions of the paper.


Conflict of interest

Christina A. Rostad's institution has received funds to conduct clinical research unrelated to this manuscript from BioFire Inc., MedImmune, Regeneron, PaxVax, Pfizer, GSK, Merck, Novavax, Sanofi-Pasteur, Micron, Janssen and Moderna. Christina A. Rostad is co-inventor on patented respiratory syncytial virus (RSV) vaccine technology unrelated to this manuscript, which has been licensed to Meissa Vaccines, Inc. Evan J. Anderson has received personal fees from AbbVie, Pfizer and Sanofi-Pasteur for consulting, and his institution receives funds to conduct clinical research unrelated to this manuscript from MedImmune, Regeneron, PaxVax, Pfizer, GSK, Merck, Novavax, Sanofi-Pasteur, Micron, Janssen and Moderna. He also serves on a safety monitoring board for

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Supporting Information

Additional supporting information may be found online in the Supporting Information section at the end of the article.

Fig S1. Antibody kinetics in serum following SARS-CoV-2 infection.

Fig S2. Phylogenetic tree of SARS-CoV-2 sequence for Patient 2.

Fig S3. Scatter plot indicates the frequency of intra-host single nucleotide variants (iSNVs) across the SARS-CoV-2 genome.

Table S1. Intra-host single nucleotide variants (iSNV) of patient 2's SARS-CoV-2 sequence.

Data S1. Supplementary Information.

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AZD1222 vaccine-related coagulopathy and thrombocytopenia without thrombosis in a young female

The rollout of a vaccination programme against severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2) has been one of the biggest steps towards controlling the global COVID-19 pandemic, with four new vaccines currently authorised for use by the European Medicines Agency (EMA). Since February 2021, a preliminary report of cases of unusual thrombosis in the setting of concomitant thrombocytopenia following vaccination has emerged.¹ The authors describe nine patients in Germany and Austria who presented with clinical and serological features which mimic heparin-induced thrombocytopenia (HIT) and the name vaccine-induced prothrombotic immune thrombocytopenia (VIPIT) was initially proposed.¹ The term vaccine induced immune thrombotic thrombocytopenia (VITT) has since been also commonly used, with the condition being referred to as thrombosis with thrombocytopenia syndrome (TTS) by the CDC and FDA. Patients presented with symptomatic thrombosis, primarily cerebral sinus venous thrombosis (CSVT), and were predominantly young females. The EMA's Pharmacovigilance Risk Assessment Committee (PRAC) noted disproportionality for rare events such as disseminated intravascular coagulopathy (DIC), CSVT and haemorrhagic stroke, especially in younger patients aged 18–54.² They have since acknowledged a causal association between AZD1222

(COVID-19 Vaccine AstraZeneca) and acquired coagulopathy as of 7 April 2021.

We describe the first reported Irish case of VITT. A 35-year-old Caucasian woman presented 14 days after vaccination with AZD1222. She presented with a medical history of migraine, with no known risk factors for thrombosis.

Following vaccination, she described general myalgia and extreme fatigue, and noticed the onset of bruising and petechiae 10 days later. She attended the Emergency Department 14 days following vaccination. Her petechiae had resolved but she had persistent bruising. She described a headache which was slightly different to her known migraine headaches. She had no recent exposure to heparin.

Laboratory investigation revealed thrombocytopenia with a platelet count of $50 \times 10^9/l$, with a normal haemoglobin and white cell count. A coagulation screen showed a mildly prolonged PT and APTT at 12.7 s (normal: 9.6–11.8 s) and 31.3 s (normal 20.8–30.8 s), respectively. Fibrinogen was reduced at 1.16 g/l (normal 1.5–4 g/l) and the D-dimer was markedly elevated at 9.83 $\mu\text{g/ml}$ (normal $< 0.42 \mu\text{g/ml}$). Examination of the peripheral blood smear showed no red cell fragmentation, no blasts and some atypical lymphocytes. A MR cerebral venogram was carried out which excluded underlying CSVT. Marrow aspirate revealed some reactive