

1 LETTER TO THE EDITOR

2 **Title:** Histiocytosis-lymphadenopathy plus syndrome on follow-up of a child with secondary  
3 hemophagocytic lymphohistiocytosis

4 *To the editor,*

5 We report the follow-up of an immunocompetent child who was diagnosed with histiocytosis-  
6 lymphadenopathy plus syndrome with an earlier report of hemophagocytic  
7 lymphohistiocytosis, secondary to Parvovirus infection<sup>1</sup>.

8 The patient first presented in 2019 at three-years of age with complaints of fever and a non-  
9 itchy maculopapular rash. She was first in birth order borne out of a non-consanguineous  
10 marriage. She presented with fever, hepatosplenomegaly and pancytopenia. Her bone marrow  
11 biopsy revealed presence of macro-ovalocytes and histiocytes, and subsequently myelofibrosis  
12 and CD71 positivity. The serum anti-parvoviral immunoglobulins were raised. A diagnosis of  
13 secondary HLH post-parvoviral infection was made which was atypical as she had HLH instead  
14 of transient pure red cell aplasia. She was discharged after the fever and pancytopenia subsided  
15 on conservative treatment. Six weeks later, she presented with multiple, sub-cutaneous  
16 swellings, on her face, neck, and trunk, which consisted of chronic inflammatory dermal  
17 deposits. The swellings resolved on their own. On follow-up, her counts were found to be  
18 within normal limits, and she remained asymptomatic in the intervening 2-year period<sup>1</sup>.

19 She then presented in June 2021, with two months history of intermittent, high-grade fever,  
20 which was associated with chills and rigors. The patient had tested positive for SARS-CoV2 on  
21 the nasal swab sample on PCR in May 2021 which was treated with oral steroids for five-seven  
22 days. She improved subsequently and didn't require any further medical treatment or testing.  
23 She developed fever 3 weeks later and was tested as positive for SARS-CoV2 on PCR  
24 (persistent positivity in the absence of a negative report). She maintained saturation on room

25 air but was hospitalized elsewhere for persistent fever and COVID positivity with a suspicion  
26 of post-COVID multisystem inflammatory syndrome in children (MIS-C), which was treated  
27 with intravenous immunoglobulins (IVIg) as per the documents. However, fever persisted, and  
28 she was referred to our hospital for further management. She did not have any other systemic  
29 complaints. There was no significant family history.

30 At presentation, she had pallor, excessive hair on forehead and face, poor oral hygiene,  
31 enlarged Group I and II cervical lymph nodes (maximum diameter 2.5 cm) and  
32 hepatosplenomegaly. Her weight was 16kgs (-0.9 SDS), height 104cm (-1.14 SDS) and head  
33 circumference 48cm (-1.35 SDS). There was no rash or arthropathy. Rest of the systemic  
34 examination was unremarkable. Repeat SARS-CoV2 RT-PCR was negative. Baseline  
35 investigations revealed bi-cytopenia with hemoglobin of 6g/dL, total leucocyte count 1,780  
36 cells/mm<sup>3</sup> (Neutrophils 25%, Lymphocytes 75%), absolute neutrophil count of 445 cells/mm<sup>3</sup>  
37 and platelets 150,000/mm<sup>3</sup>. The peripheral blood smear showed normochromic normocytic  
38 anemia without any atypical lymphocytes. Serum transaminases were raised (alanine  
39 aminotransferase 111 units/L aspartate aminotransferase 64 units/L), blood urea 24 mg/dL,  
40 serum creatinine 0.3mg/dL and qualitative COVID antibody (IgG) were raised. Further  
41 investigations showed raised inflammatory markers – D-dimer (>5000 ng/mL), interleukin-6  
42 (127.2pg/ml), ferritin (1300 ng/mL) and C-reactive protein (225.66 mg/L). She was started on  
43 antibiotics and workup for other inflammatory conditions was planned. The workup for  
44 tuberculosis, blood culture, urine culture, bone marrow culture, WIDAL test for salmonellosis,  
45 HIV by ELISA, IgM ELISA for dengue virus, parvovirus, scrub typhus and anti-viral capsid  
46 antibody for Epstein Barr virus were found negative. Serum antinuclear-antibodies was  
47 negative. A chest radiograph, electrocardiograph and echocardiography were normal. Contrast  
48 enhanced computed tomography of neck, chest, abdomen revealed hepatosplenomegaly and  
49 fibro-atelactatic opacities in right upper zone, middle zone and left lingular segment. Cervical

50 and bilateral axillary lymphadenopathy with mild cardiomegaly were also reported. A fine-  
51 needle aspiration from cervical lymph nodes showed reactive hyperplasia.

52 At the end of the first week of hospitalization, the child's fever, organomegaly and  
53 lymphadenopathy had not improved. The bi-cytopenia worsened into pancytopenia  
54 (hemoglobin – 5.9g/dL, total leucocyte count- 1,470 cells/mm<sup>3</sup>, (Neutrophils 24%,  
55 Lymphocytes 75%, Monocytes 1%) and platelet count- 50,000/mm<sup>3</sup>. Bone marrow aspirate  
56 and biopsy revealed bi-cytopenia with diluted marrow and decreased cellularity, suggestive of  
57 WHO grade -1 myelofibrosis. There were no blast cells or any other evidence of malignancy.  
58 The repeat blood parameters were suggestive of HLH (ferritin- >2000ng/mL, triglyceride- 310  
59 mg/dL, fibrinogen -138 mg/dL).After excluding likely possible infections and malignancy, a  
60 possibility of SARS-CoV2 induced HLH with persistent MIS-C was kept in view of persistent  
61 fever associated with rash, pancytopenia, SARS-CoV2 antibody positivity and raised  
62 inflammatory markers. Intravenous methylprednisolone pulse therapy at 30mg/kg/day was  
63 initiated.

64 Keeping a possibility of exaggerated post-viral immune mediated bone marrow suppression in  
65 this immunocompetent child who previously manifested as parvovirus induced HLH and now  
66 with SARS-CoV2 induced HLH, a possibility of disorders of immune dysregulation was  
67 considered<sup>2</sup>. Genetic testing with whole exome sequencing was performed (outsourced) which  
68 detected a homozygous 5'splice variation in intron 2 of the SLC29A3 gene, affecting the  
69 invariant GT donor splice site of exon 2, a finding consistent with the diagnosis of histiocytosis-  
70 lymphadenopathy plus syndrome (**Figure 1**).

71 The child completed 5 days of methylprednisolone pulse therapy for 5 days followed by oral  
72 steroids over two weeks considering histiocytosis-lymphadenopathy plus syndrome as a  
73 possible diagnosis. She became afebrile after third dose of methylprednisolone with decrease  
74 in lymphadenopathy and hepatosplenomegaly and, was discharged home after three weeks of

75 hospitalization. Last laboratory parameters showed declining trend in inflammatory markers.  
76 Her last hearing screen was normal before discharge.

77 Histiocytosis-lymphadenopathy plus syndrome, also known as “*SLC29A3* spectrum disorder”<sup>3</sup>,  
78 is a group of conditions characterized by mutations in the *SLC29A3* gene, present on  
79 chromosome 10q23 that codes for a nucleoside transporter (ENT3 transporter)<sup>4</sup>. The variable  
80 expressivity of this gene results in varied phenotypic presentations like H syndrome (cardiac  
81 anomalies, camptodactyly, endocrinal disorders, hepatosplenomegaly), pigmented  
82 hypertrichosis with insulin-dependent diabetes mellitus (PHID), Faisalabad histiocytosis, and  
83 familial Rosai-Dorfman disease. All these conditions have an autosomal recessive mode of  
84 inheritance and are characterised by presence of histiocytosis. Accumulation of histiocytes in  
85 different organs and tissues is responsible for the clinical manifestations associated with  
86 disorders of this spectrum<sup>3</sup>.

87 Patients with histiocytosis-lymphadenopathy plus syndrome typically have cutaneous  
88 hyperpigmentation, hypertrichosis (as seen in this patient), short stature refractory to GH  
89 therapy, cardiac anomalies, hypogonadism, hyperglycemia/IDDM, hepatosplenomegaly,  
90 lymphadenopathy (also present in this patient), and various musculo-skeletal pathologies<sup>4,5</sup>.  
91 The skin rash in index case during the earlier hospitalization<sup>1</sup>, was probably a spectrum of H-  
92 syndrome. An understanding of the transcriptome profile in H syndrome links *SLC29A3*  
93 mutations with mitochondrial dysfunction and oxidative stress which causes immune  
94 dysregulation, as was seen in the index patient<sup>6</sup>. The persistent SARS-CoV 2 antigenemia and  
95 elevated inflammatory markers in the index case were contributory to development of MIS-C,  
96 with the predisposition of the underlying genetic profile. The neutrophilic signatures in MIS-  
97 C were clearly seen to be different from those with acute COVID or in healthy controls, though  
98 testing for any underlying genetic signatures was not performed<sup>7</sup>. Type 1 diabetes and  
99 seronegative arthritis are however, commoner manifestations reported in this

100 spectrum<sup>8</sup>. Rheumatological, endocrinal manifestations and camptodactyly may evolve with  
101 age in affected patients, necessitating the need for a continued follow-up<sup>9</sup>.

102 The management of this condition has essentially been symptomatic. However, auto-  
103 inflammatory manifestations like rheumatological manifestations require immunosuppressant  
104 drugs. A case series of five patients reported partial response to oral steroids with need to use  
105 other immunosuppressive agents<sup>8</sup>. Monoclonal antibodies like adalimumab<sup>8</sup> and tocilizumab  
106 have been used in cases with persistent arthritis and autoinflammation refractory to steroids  
107 and immunosuppression<sup>10</sup>. The index patient responded to steroids in this admission but may  
108 need further drugs with evolution of disease phenotype in the future.

109 In hindsight, our patient, presented with immune dysregulation with H syndrome after mild  
110 viral exposures where an ominous genetic basis could be established. This highlights the need  
111 to suspect children with atypical manifestations of infectious diseases early and offer genetic  
112 diagnosis for better disease prognosis and management.

113 **KEYWORDS:**

114 HLH

115 Histiocytosis-Lymphadenopathy plus syndrome

116 H syndrome

117 Steroid therapy

118 MIS-c

119 Post-viral immune dysregulation

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126 ***Authors declare no conflict of interest***

127 **LEGEND:** Whole exome sequencing confirming pathogenic mutation in *SLC29A3* gene

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