

Table S1. Number of co- infections identified for novel HAsVs.

	HAsV-MLB	HAsV-VA
Double infections		
Adenovirus	4	4
Classic HAsV	1	5
Rotavirus	3	0
Sapovirus	1	0
Norovirus	0	1
Triple infections		
Adenovirus and Rotavirus	1	1
Adenovirus and Sapovirus	2	0
Classic HAsV and Norovirus	0	1
Quadruple infections		
Adenovirus, Rotavirus and Sapovirus	1	0