Influenza Surveillance of Families in an Observational Household Study 2019-2021

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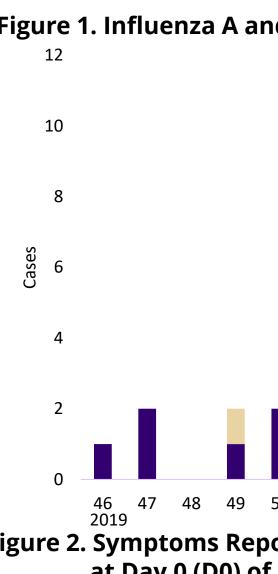
Background

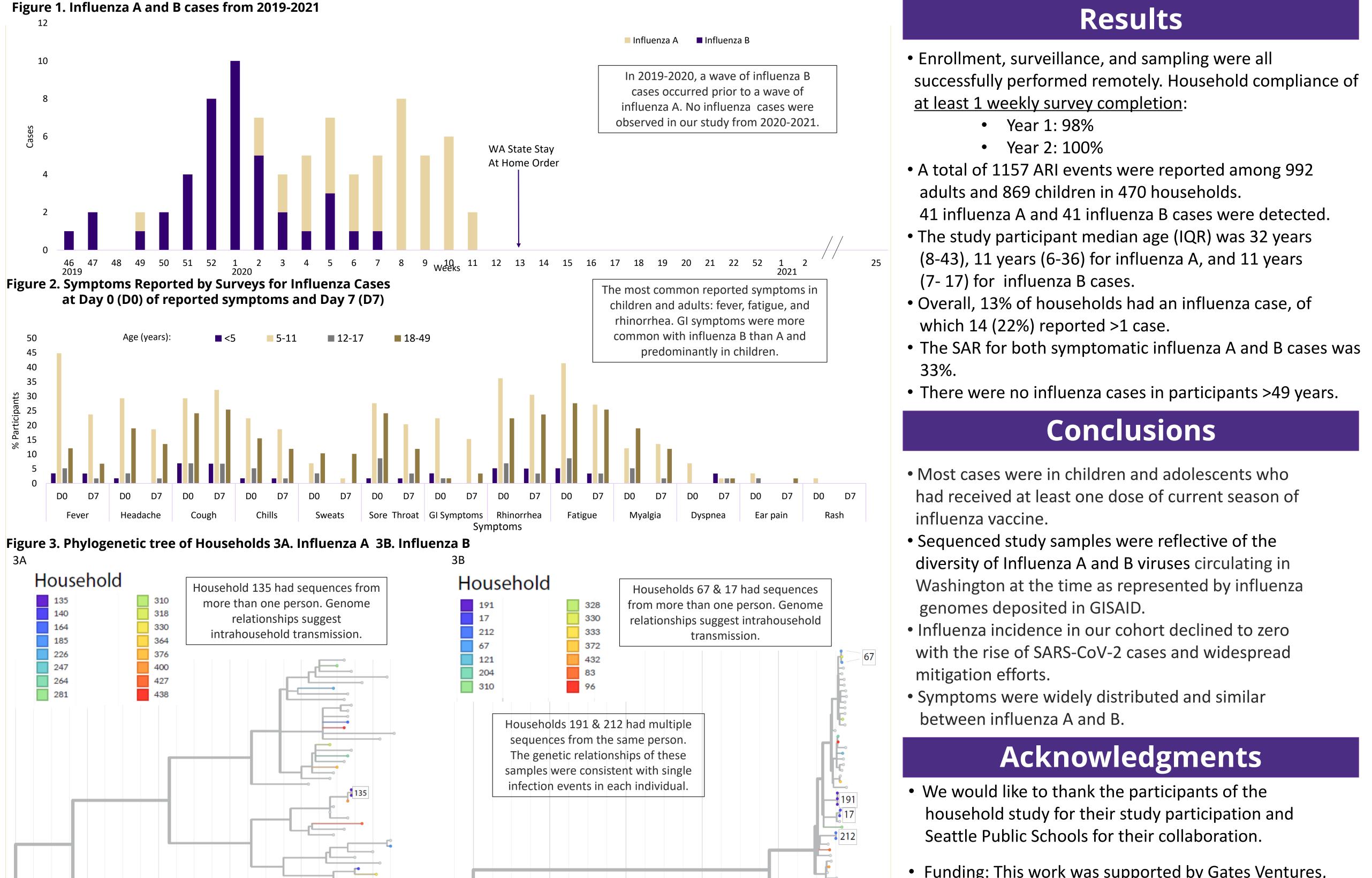
- Families with children may be at higher risk for influenza virus infection.
- The burden of influenza disease may be underestimated without community surveillance.

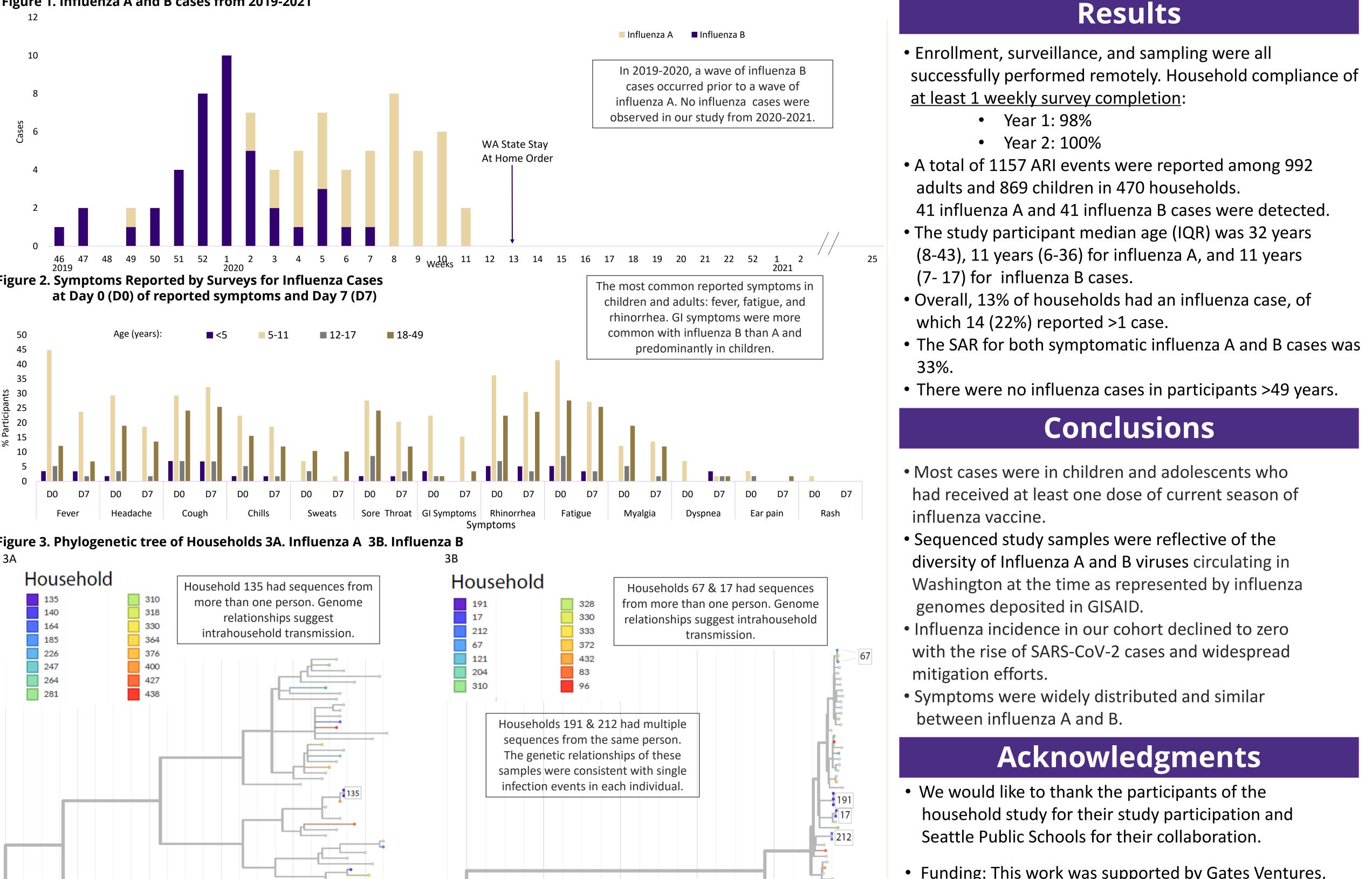
Methods

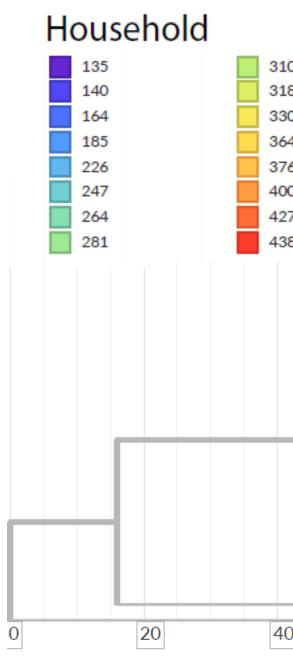
- Households with school-aged children were enrolled, mailed home specimen collection kits, and asked to self-assess for weekly acute respiratory illness (ARI).
- Participants with cough or ≥ 2 ARI pre-determined symptoms were prompted to complete serial illness surveys with parental or self-collected midturbinate nasal swabs.
- Samples were sent to a University of Washington study laboratory for RT-PCR influenza testing.
- Whole genomic sequencing was attempted on samples with Ct<30 and phylogenetic trees created using publicly available influenza sequences in WA state.
- Secondary attack rates (SAR) were calculated

Table 1	Total participants	Influenza	
Characteristics	n=1861 (%)	A n=41 (%)	B n=41 (%)
Sex			
Female, n (%)	951 (51)	20 (49)	22 (53)
Hispanic/Latinx	110 (6)	1 (2)	6 (14)
Race*			
White	1668 (89)	38 (92)	35 (85)
Black	27 (1)	1(2)	2 (5)
Asian	244 (13)	12 (29)	9 (22)
Other	55 (3)	0	3 (7)
Prefer not to say	42 (2)	0	2 (5)
Age (years)			
<5	196 (11)	5 (12)	3(7)
5-17	673 (36)	23 (56)	28 (68)
18-64	967 (52)	13 (32)	10 (24)
>64	25 (1)	0	0
Seasonal Flu Vaccine 2018-2019	1152/1379 (83)	34/41 (83)	38/41 (93)
Seasonal Flu Vaccine 2019-2020	1525/1806 (84)	36/38 (95)	38/41 (93)

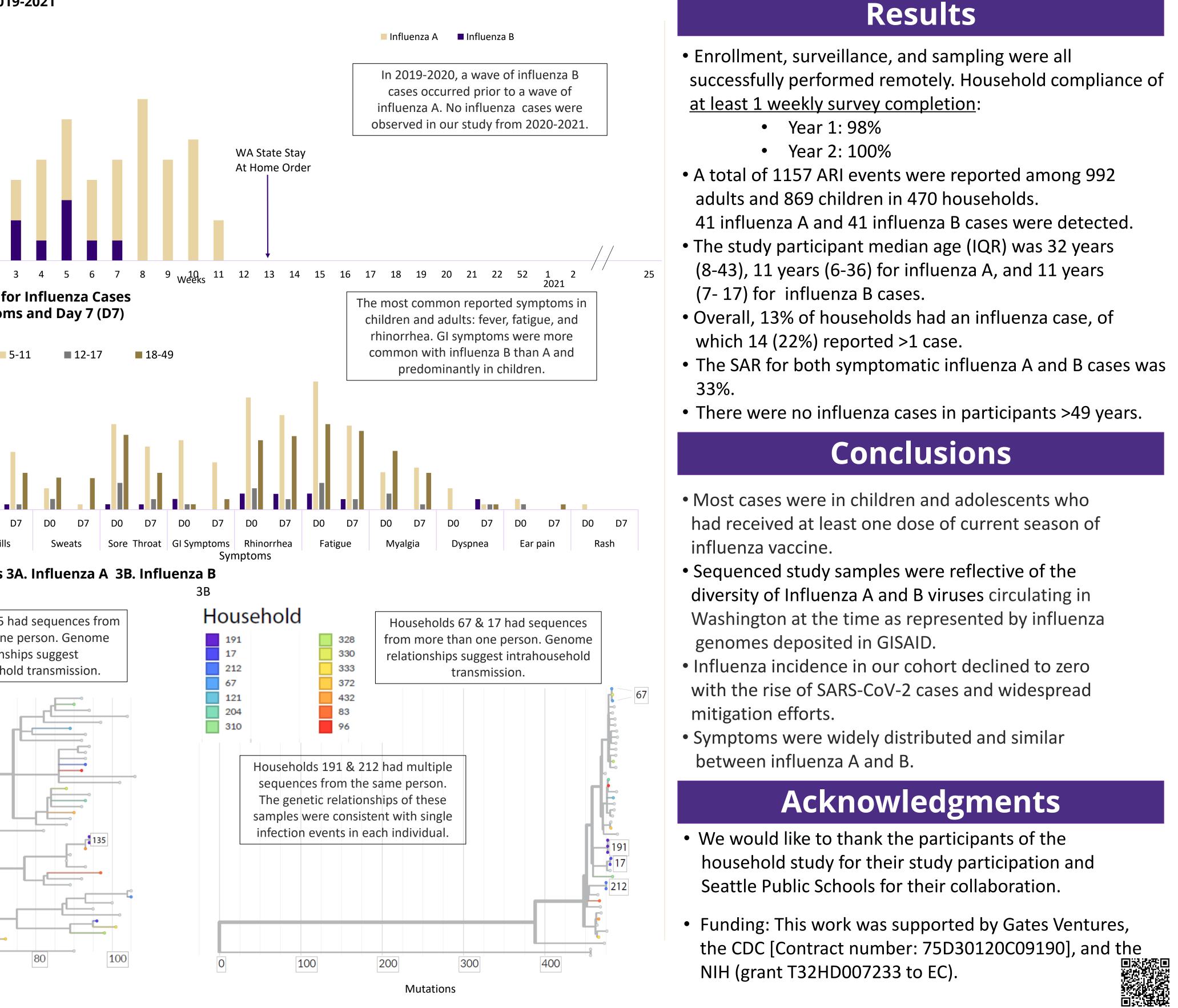








*multiple answers were included



Mutations

60



