

Background

CD8+ T cells may transiently suppress viral replication in acute persisting hepatitis C virus (HCV) infections, but eventually lose control due to T cell exhaustion or selection of viral variants with escape mutations. With the failure of cellular immunity, high-level viremia is a hallmark of chronic infection.

Following pregnancy however, some chronically infected women experience a substantial drop in viremia.

CD4+ T

Fig 1: Model of adaptive immune responses in acute persisting infection and with pregnancy



Postpartum control of viral replication is due to recovery of the HCV-specific CD8 T cell response and will thus be associated with selection of de novo escape mutations as seen in acute persisting infection.

HCV genomic evolution was compared in 8 women with different patterns of postpartum viral replication. Viral sequence data were obtained in the 3rd trimester or perinatal period (T3), 3 months postpartum (3P), and 12-24 months postpartum follow-up (FU) by Illumina or clonal Sanger sequencing of PCR amplicons spanning the HCV coding region. Analysis focused on nonstructural region to limit confounding effects of antibody pressure.

Comparison:

- Subjects were categorized based on change in HCV-RNA level from (T3) to 3P: elite controllers (>2 \log_{10} drop) and noncontrollers (stable viral load)
- Nucleic acid changes were tabulated if frequency <20% at baseline and >80% of variants at follow-up. Non-synonymous changes were compared to genotype consensus sequence and assessed for location with described and predicted class I epitopes.

Cohort included:

- 4 women with viral control
- 4 women without viral control



Fig 2: Viral load changes between T3 and 3P

Evolution of Hepatitis C Virus Points to Postpartum Recovery of CD8+ T-cell Selection Pressure

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Results

Change in Viral Load 3P-T3 (log₁₀) Fig 5A, 5B, 5C: Correlation of mutations with degree of viral control A: All mutations from 3rd trimester to follow-up B: All mutations from 3rd trimester to 3 months postpartum









6A

6B



pre E1 E2	P7 NS2 NS3 N	IS4A NS4B NS5A	NS5B	Core E1 E2	P7 NS2 NS3	NS4A NS4B NS5A	NS5B
M027				M067			
T3		2000	3000	<u>T3</u>			3000
	1000		3000	, , , , , , , , , , , , , , , , , , ,		2000	
12P				12P			
,	1000	<u> Ц</u>		0	1000	2000	3000
M062]			M073			
Г3	-			T3			
	1000	2000	3000	0	1000	2000	3000
2P	· · · · · · · · · · · · · · · · · · ·			12P			
	1000	2000	3000	0	1000	2000	3000

Fig 6: Visual representation of non-synonymous viral evolution after pregnancy in elite controllers (A) and noncontrollers (B) comparing 3rd trimester sequence to follow-up.

- period among women with viral control.
- women after childbirth



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THE OHIO STATE UNIVERSITY

LEGE OF MEDICINE

Elite Controllers

NS5A NS5B Core E1 E2 P7 NS2 NS3 NS4A NS4B NS5A NS5B MO16-2 T3 0 1000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 T3 0 1000 AA Pesition 2000 3000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 18P 18P 0 1000 AA Pesition 2000 3000 18P 18P 0 1000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 18P 18P 0 1000 AA Pesition 2000 3000 18P 18P 0 1000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 18P 18P 0 1000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 18P 0 1000 AA Pesition 2000 3000 18P 18P 0 1000 AA Pesition 2000 3000 18P 18P 0 1000 AA Pesition 2000 3000 18P 18P 18P 18P 18P 18P 18P 18P							_
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bo 3000 AA Position 2000 AA Position 2000 3000 T3			18P				
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T3 T3 AA Position 1000 AA Position 18P 1000 AA Position AA Position 1000 AA Position			M03	3			
AA Position AA Position AA Position AA Position AA Position AA Position AA Position AA Position			Т3				
18P 100 AA Position 1000 1	00	3000		1000	AA Position	3000	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$			18P				
	······································	3000	o	1000	AA Position	3000	

Green: toward consensus Purple: tangential to consensus Red: away from consensus

Noncontrollers

Conclusions

Significantly more viral mutations emerged in elite controllers after pregnancy compared to non-controllers (p=0.0286), with a similar trend noted when limited to nonsynonymous mutations (p=0.0857)

The number of mutations accumulated at follow-up correlated with degree of viral control. These changes may have been enriched in the early postpartum

Most nonsynonymous mutations occurred in predicted class I epitopes

Together, these findings suggest that re-exertion of HCV-specific CD8 T cell pressure may contribute to the unique control of HCV observed in some