Increasing Hepatitis C Virus (HCV) Screening and Confirmatory Testing in the Birth Cohort in a Large Integrated Health System Carla V. Rodriguez PhD, MPH¹, Kevin B. Rubenstein¹ MS, Haihong Hu¹ MS, Benjamin Linas^{2,3} MD, MPH, Michael Horberg¹ MD, MAS ¹Kaiser Permanente, Mid-Atlantic Permanente Research Institute, Rockville, MD; ² HIV Epidemiology and Outcomes Research Unit, Section of Infectious Diseases, Boston Medical Center; ³Department of Epidemiology, Boston University School of Public Health, Boston, MA

Objective: Describe trends in screening and confirmatory testing in Kaiser Permanente Mid-Atlantic States (KPMAS) relative to the 2013 release of the U.S. Preventative Services Task Force "birth cohort" (born 1945-1965) screening recommendations.

Methods:

- Cohort study, patients \geq 18 yrs with \geq 8 months of enrollment from 1/1/2003-12/31/2014 and ≥ 1 clinical visit.
- Annual screening rate estimated as the number antibody (Ab) tested per persons enrolled each year.
- Survival methods used to describe factors associated with time to Ab testing.
- Stratification by Service Area, interactions with time and robust standard errors to address nonproportional hazards.
- Among Ab+, we describe the cumulative probability and predictors of confirmatory RNA or genotype testing.

Conclusion:

- High screening prevalence, yet >16% of Ab+ were not confirmed, particularly MSM and those with elevated ALT.
- Higher screening rate in non-birth cohort underscores continued focus on risk-based screening.
- Increased screening rate over time in birth-cohort suggests shift.
- More time is needed to confirm this trend, which may be confounded by the advent of new DAAs and increased screening outreach.

1. We observed 665,345 members over 11 years. The KPMAS population is **ethnically diverse**, slightly **more** female and has above average median income (nationally and regionally*). A higher proportion of the "Birth Cohort" were seen in GI/ID clinic and had elevated ALT compared with non-cohort members.

	Not Birth Cohort	Birth Cohort
	404522	260823
Age at Enrollment,		
mean (sd)	37.5 (17.1)	49.5 (6.53)
Median Household	82559.59	86366.32
Income ¹ , mean (sd)	(36599.65)	(38527.86)
Race ²		
Black,%	35.3%	36.4%
American		
Indian/Alaskan Native	0.2%	0.2%
Asian/Pacific Islander	9.8%	9.5%
Hispanic	12.0%	10.9%
Multi-Racial	1.8%	1.8%
White	40.8%	41.2%
Sex, % (n)		
Female	58.1% (235035)	55.0% (143450)
Male	41.9% (169487)	45.0% (117373)
HBV+, % (n)	0.5%(2080)	0.6% (1521)
HIV+, % (n)	0.5% (2051)	0.8% (2032)
Seen by GI/ID (ever), %		
(n)	11.2% (45400)	16.1% (41920)
MSM, % (n)	1.6% (1101)	1.6% (720)
Illicit Drug Use (ever),		
% (n)	<0.1%(28)	<0.1%(50)
Elevated ALT ³ , % (n)	2.4%(9548)	4.5% (11790)
1 Derived from 2013 US Census: *U.S. Census Bureau, 2013 American		

Community Survey

2 Race from self-report; missing values imputed using the Bayesian Improved Surname Geocoding Algorithm (Elliott, M. et.al. (2008); Health Services Research, 43(5 Pt. 1), 1722-1736.)

3 Alkaline Amino Transferase; elevated = 2 consecutive measures >60 IU/L

2. Screening rates increased over time: 23.6 (2004) to 70.8 (2014) per 1000 person-years; sharpest increase after 2013. In total, 18.6% of KPMAS members were screened for HCV.





3. By 120 months, 66% of the non-birth cohort population was left unscreened; compared to 74% of the birth cohort.





5. Among Ab+, 84% received confirmatory testing. No significant differences by service area, race, history of drug use, HBV/HIV status were observed. MSM (aHR=0.66; 95% CI: 0.47, 0.97) and those with elevated ALT (aHR=0.85; 95% CI: 0.76, 0.95) were less likely to receive confirmatory testing. A total of 2.9% of the screened population had confirmed HCV.

4. Although patients in the birth cohort had a lower screening rate, their risk of screening increased faster over time compared to those in the non-birth cohort. Other important predictors of screening included female sex, MSM, Black, Hispanic and Asian Race, IDU,

Risk of Screening by Demographic, Behavioral and Clinical Characteristic. KPMAS 2003–2014

