centrifugation. We continued to see suspected erroneous results. We repeated the test in EDTA tubes. 16/19 had HIV VL < 20, the remainder ranged from 27-41 copies/ml. We then implemented the use of EDTA tubes in all samples. No further cases of falsely elevated VL have been seen since the change.

Table 1

PT#	Undetectable for the past 6 moths	Referred absolute adherence	Baseline VL	PPT VL	EDTA VL	days from PPT to EDTA
1	yes	yes	<20	296	<20	8
2	yes	yes	<20	595	<20	15
3	yes	yes	<20	233	<20	6
4	yes	yes	<20	1050	<20	6
5	yes	yes	<20	270	<20	5
6	yes	yes	<20	490	<20	20
7	yes	yes	<20	169	<20	6
8	yes	yes	<20	121	<20	22
9	yes	yes	<20	357	27	13
10	yes	yes	<20	453	41	6
11	yes	no *	<20	610	<20	6
12	yes	yes	<20	1030	<20	10
13	yes	yes	<20	875	<20	7
14	yes	yes	<20	246	<20	6
15	no	yes	160,000	397	<20	6
16	yes	yes	<20	539	30	30
17	yes	yes	<20	368	<20	30
18	yes	yes	<20	158	<20	41
19	no	yes	356	2530	<20	90
20	yes	yes	<20	588	not done	
* missing o	one dose per mo	onth				

Conclusion. We describe potential steps that can interfere in the accuracy of HIV VL results. Detailed review of past history and patient's adherence to treatment is essential to identify systemic problems that may lead to errors. Close communication with staff and laboratory provider will be key to overcome such challenges.

Disclosures. All Authors: No reported disclosures

972. Infection and Overdose Prevention for Persons with Injection Drug Use-Related Infections: Evaluation of an Inpatient Quality Improvement Program

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Session: P-46. HIV: Prevention

Background. Hospitalizations for injection drug use-related infections (IDU-I) are increasing in North Carolina and nationally. Many IDU-I, such as endocarditis, bone, joint, and spine infections, require long antimicrobial courses and extended inpatient stays. These hospitalizations are opportunities to engage patients in overdose and infection prevention.

Methods. A quality improvement (QI) program was piloted for inpatients with IDU-I. Eligible patients admitted to the inpatient pulmonary or infectious disease teams from 11/2019 to 01/2020 were referred to the QI team if they reported or were suspected to have injected drugs over the past year, or felt to benefit from drug-related infection prevention and overdose services. A checklist of recommendations to the care teams included: (1) screening for HIV, Hepatitis B (HBV) and C (HCV), (2) immunization for Hepatitis A (HAV), HBV, and tetanus, (3) prescription of naloxone at discharge, and (4) information on a syringe services program in or near their county. After review of the medical record, the QI team made recommendations on the appropriate taks from the checklist. The number of QI checklist tasks performed on the two inpatient teams during a 9-week pilot period (the above period excepting a two-week break) was reviewed. Baseline comparison data was not incorporated, owing to the challenges in retrospective identification of IDU-I.

Results. 20 patients were included in the intervention. The median age was 32 years (IQR 27-38) and 70% were female. The most common diagnosis was endocarditis (40%) and the median length of stay was 11 days (IQR 5-42). HIV and HCV tests were each conducted in 95% of patients (Table). Screenings for HAV and HBV immunity were done in 90% of patients. HAV, HBV, and Tdap immunizations were given to 20%, 35%, and 50%, respectively. Naloxone was provided to 60% of patients at discharge and half of patients were referred to syringe programs. HCV was detected in 8 patients and HBV in 2 patients. No patients were diagnosed with HIV.

Percentage of infection and overdose prevention services provided to eligible IDU-I patients during hospitalization.

Intervention	Done	% Receiving Intervention	
intervention	n = 20		
HIV test	19	95%	
Hep A Immunity	18	90%	
Hep B Screening	18	90%	
Hep C Screening	19	95%	
Hep A Immunization	4	20%	
Hep B Immunization	7	35%	
Tdap Immunization	10	50%	
Naloxone prescription	12	60%	
Syringe program referral	10	50%	
Hep C Diagnoses	8		
Hep B Diagnoses	2		
HIV Diagnoses	0		

Conclusion. In a setting without comprehensive addiction consultation, a simple intervention provided guideline-concordant infection and overdose prevention services for persons hospitalized for IDU-I.

Disclosures. All Authors: No reported disclosures

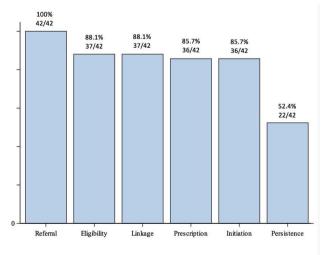
973. PrEP Care Continuum among Transgender Women at a Patient-Centered PrEP Program in Atlanta, GA

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Session: P-46. HIV: Prevention

Background. HIV disproportionally affects transgender women (TGW) of color, with a prevalence of 26% and 44% among Latinx and Black TGW, respectively. Low medication adherence likely contributed to suboptimal pre-exposure prophylaxis (PrEP) efficacy among TGW in clinical trials, but real-world PrEP outcome data for TGW is limited. In this study, we developed the PrEP care continuum for TGW referred to a PrEP program at a large, safety-net urban hospital in the Southeast.

Figure 1. PrEP care continuum of TGW referred a PrEP program. Referrals include all TGW referred to PrEP clinic, eligibility includes all those referred who were deemed eligible for PrEP, linkage refers to those eligible who had ongoing care at the PrEP clinic, prescription refers to those who received their first prescription of PrEP, initiation includes those who started taking the PrEP they were prescribed, and persistence includes those who had a visit within 6 months of study end.



Methods. We analyzed data for those referred to the PrEP program from 3/2018 to 2/2020. We determined the proportion of TGW who were linked to the program, provided a PrEP prescription, started PrEP, and persisted in PrEP care, defined as having at least one follow-up visit within 6 months. Using a multivariate regression model, including age, race, ethnicity, mental health co-morbidities, and substance use, we determined factors associated with persistence in PrEP care. We calculated rates of sexually transmitted infections (STIs) and HIV incidence.

Results. Of the 321 total referrals for PrEP, 42 (13%) were TGW. 81% of TGW were referred from a co-located gender clinic. Median age was 28.5 years (IQR 23-34), 62% were Black, 21% had mental health co-morbidities, 45% used substances, and 35% engaged in transactional sex. Of all TGW who were referred, 37 (88%) were eligible for PrEP and linked to care, 36 (85.7%) were prescribed and initiated PrEP, and 22 (52.4%) persisted in care at the end of the study period. There were no factors associated with persistence in PrEP care. The most common STIs during the first visit were pharyngeal gonorrhea (22.7%) and syphilis (16.7%). STI incidence was highest for rectal chlamydia (12.5%) and pharyngeal gonorrhea (6.5%). There was one HIV seroconversion during the study period.

Conclusion. In a public hospital-based PrEP clinic in Atlanta with a co-located gender clinic, TGW had high rates of linkage to care and PrEP prescription and initiation, despite high rates of mental health diagnoses and substance use. However, there was a significant drop-off in persistence. STI prevalence and incidence were high, but there was only one HIV seroconversion, highlighting the potential benefits of PrEP. Future studies are needed to assess interventions to optimize persistence in PrEP care among TGW.

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974. The Effects of Changes in State-Level Policies Affecting Eligibility for the Supplemental Nutrition Assistance Program (SNAP) on Annual HIV Diagnoses in the United States

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Session: P-46. HIV: Prevention

Background. The connection between food insecurity and HIV outcomes is well-established. The Supplementary Nutrition Assistance Program (SNAP), the primary program in the United States that addresses food insecurity, may have collateral impacts on HIV incidence, but the extent to which it does is unknown. "Broad-based categorical eligibility" for SNAP is a federal policy that provides a mechanism for states to increase the income or asset limits for SNAP eligibility. The Department of Agriculture under the Trump Administration has proposed eliminating this policy.

Methods. We estimated the association between the number of new HIV diagnoses from 2010 to 2014 for each state and (1) state income limits for SNAP eligibility as a percentage of the federal poverty level and (2) state asset limits for SNAP eligibility (increased/eliminated vs. unchanged). We fitted multivariable negative binomial regression models with annual incidence of HIV diagnoses specified as the outcome; SNAP policies as the primary explanatory variable of interest; state and year fixed effects; and time-varying covariates related to the costs of food, health care, housing, employment, SNAP outreach, and total spending on Temporary Assistance for Needy Families (TANF) programs.

Results. From 2010 to 2014, 204,034 new HIV diagnoses occurred in the United States. HIV diagnoses within states had a statistically significant inverse association with state income limits for SNAP eligibility (IRR 0.94 per increase in the income limit by 35% of federal poverty level, 95% CI 0.91-0.98), but no statistically significant association with state asset limits (increased asset limit vs. no change, IRR 1.02, 95% CI 0.94-1.10; eliminated asset limit vs. no change, IRR 1.04, 95% CI 0.99-1.10) (Table). Table

Table – The relationship between state-level policies affecting Supplementary Nutrition

Assistance Program (SNAP) eligibility – income limit as a percentage of the federal poverty level (FPL) and asset limit, either increased or eliminated compared to unchanged – and the annual number of new HIV diagnoses from 2010 to 2014 using negative binomial regression models.

		Unadjusted		State/year fixed effects		Both policies, fixed effects, and time- varying covariates	
		IRR	95% CI	IRR	95% CI	IRR	95% CI
Income limit (per increase of 35% FPL)		1.11	1.01-1.23	0.95	0.92-0.98	0.94	0.91-0.98
Asset limit	No change	Ref		Ref		Ref	
	Increased	0.92	0.61-1.39	0.96	0.89-1.04	1.02	0.94-1.10
	Eliminated	1.23	0.99-1.53	0.98	0.94-1.03	1.04	0.99-1.10

Abbreviations: CI, confidence interval; FPL, federal poverty level; IRR, incidence rate ratio; SNAP, Supplementary Nutrition Assistance Program

Notes: Time-varying covariates include average meal cost in a food secure household, health expenditure per capita, house price index, high school graduation rate, unemployment rate, uninsured rate, state spending on SNAP outreach, and total Temporary Assistance for Needy Families (TANF) spending.

Conclusion. State income limits for SNAP eligibility were inversely associated with the number of new HIV diagnoses for states between 2010-2014. Proposals to eliminate the use of broad-based categorical eligibility to increase the income limit for SNAP may undercut efforts to end the HIV epidemic in the United States.

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975. Why Sex Make a Difference in HIV Clinical Course? Bioinformatics Analysis of Differential Expressed Gene in Females and Males with HIV Disease

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Session: P-46. HIV: Prevention

Background. Human immunodeficiency virus (HIV) disease progression are different among genders, in which women usually progress to acquired immunodeficiency syndrome (AIDS) faster than men. The mechanisms resulting in the gender biases of HIV progression are unclear. We conducted a bioinformatics analysis of differentially expressed genes (DEGs) in women and men with HIV disease to understand the sex-based differences in HIV pathogenesis.

Methods. We obtained microarray data from the Gene Expression Omnibus (GEO) database using our pre-defined search strategy and analyzed data using the GEO2R platform. The t-test was done to compare DEGs between females and males with HIV diseases. The Database for Annotation, Visualization, and Integrated Discovery (DAVID) was implemented to systematically extract biological features and processes of retrieving DEGs via gene ontology (GO) analysis. A Systemic search was performed to evaluate each DEG function and its possible association with HIV.

Results. One gene expression profiling data were retrieved: GSE 140713, composed of 40 males and 10 females with HIV1 infected samples. A GEO2R analysis yielded 19 DEGs (Table 1). The GO analysis result was demonstrated in Tables 2 and 3. Following a systemic search, we found two DEGs, which have previous studies reported an association with HIV: DDX3X (20 studies) and PDS5 (1 study). We proposed DDX3X (t 5.3, p 0.0037) is responsible for gender inequalities of HIV progression because of: 1. DDX3X is needed in the HIV1 life cycle. 2. Several studies confirmed a positive correlation between DDX3X expression and HIV1 replication. 3. Our study found an up-regulated DDX3X expression in women corresponded to the fact that women progress to AIDS faster than men. 4. Our GO analysis showed female up-regulated genes were enriched in positive regulation of the gene expression pathway, which can be explained by DDX3X and its underlying mechanism.

Table 1: DEGs in women and men with HIV1 disease

Down-regu	lated genes in femal	e compare t	o male with HI	V1 disease
Genes	Adjusted P value	t	В	logFC
RPS4Y2	1.74e-47	-69.61	58.56	-9.96
RPS4Y1	1.26e-44	-60.22	57.58	-9.21
UTY	7.62e-39	-45.75	54.97	-5.27
TXLNGY	2.84e-31	-31.75	49.4858	-5.073
DDX3Y	2.08e-25	-31.75	49.4858	-5.073
EIF1AY	5.47e-25	-23.75	43.24326	-7.96
KDM5D	5.04e-21	-23.19	42.66616	-6.036
ZFY	1.90e-20	-18.85	37.21756	-4.085
TTTY10	7.00e-08	-8.43	14.21921	-2.185
TTTY15	6.75e-07	-7.78	12.24742	-2.613
PRKY	4.29e-05	-6.6	8.54374	-1.449
TTTY14	8.23e-04	-5.73	5.82023	-1.038
Up-regulat	ed genes in female o	ompare to n	nale with HIV1	disease
XIST	1.06e-31	32.55	49.9428	11.789
EPB41L4B	6.71e-04	5.8	6.03312	1.507
DDX3X	3.73e-03	5.3	4.46087	0.798
PDS5B	5.62e-03	5.17	4.06903	1.663
HEPH	7.98e-03	5.06	3.72446	1.55
LANCL2	7.98e-03	5.05	3.70198	1.553
EIF1AX	1.48e-02	4.86	3.12847	0.562

Table 2: GO functional enrichment pathway analyses of overall retrieving DEGs

GO enrichment analyses of the DEGs	
GO pathway: Biological process (BP)	
 Translational initiation and Regulation of gene expression 	
 RNA secondary structure unwinding 	
- Chromosome segregation	
- Oxidation-reduction	
GO pathway: Cellular component (CC)	
 Cytosolic small ribosomal subunit and Nucleus 	
GO pathway: Molecular function (MF)	
Histone demethylase activity	
 ATP and rRNA binding 	
 Translation initiation factor activity 	
 ATP-dependent RNA helicase activity 	