

# The contribution of injecting drug use as a risk factor for hepatitis C virus transmission globally, regionally, and at country level: a modelling study

**Adam Trickey**<sup>1,2</sup>, Hannah Fraser<sup>1</sup>, Aaron G Lim<sup>1</sup>, Amy Peacock<sup>3</sup>,  
Samantha Colledge<sup>3</sup>, Josephine Walker<sup>1</sup>, Janni Leung<sup>3,4</sup>, Jason Grebely<sup>3</sup>,  
Sarah Larney<sup>3</sup>, Natasha K Martin<sup>5,1</sup>, Matthew Hickman<sup>1,2</sup>, Louisa  
Degenhardt<sup>3</sup>, Margaret T May<sup>1,2</sup>, Peter Vickerman<sup>1,2</sup>

<sup>1</sup> Population Health Sciences, University of Bristol, UK

<sup>2</sup> NIHR Health Protection Research Unit in Evaluation of Interventions, UK

<sup>3</sup> UNSW Sydney, Australia

<sup>4</sup> Centre for Youth Substance Abuse Research, University of Queensland, Australia

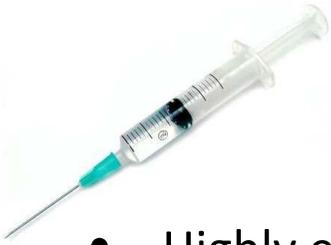
<sup>5</sup> Division of Infectious Diseases and Global Public Health, University of California San Diego, USA



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# Hepatitis C virus epidemic

- Highly effective direct acting antivirals (DAAs) have been developed as treatment for hepatitis C virus (HCV) infection
- An estimated **71 million** people are infected with HCV globally<sup>1</sup>
- The WHO has set targets to eliminate HCV by 2030
  - Includes reducing incidence by 80% from 2015 levels
- HCV is highly prevalent (>30%) among people who inject drugs (PWID)<sup>2</sup>
- Proportion of adults that are PWID in most countries is low (<0.5%)<sup>2</sup>
- Generally assumed role of IDU to HCV transmission important mostly in high-income countries<sup>3</sup>
- In low- and middle-income countries it is thought most HCV transmission is due to other routes<sup>3</sup>
  - Eg. Unsterile medical injections and unscreened blood transfusions

1: Global prevalence and genotype distribution of hepatitis C virus infection in 2015: a modelling study. Blach, 2017

2: Global prevalence of injecting drug use and sociodemographic characteristics and prevalence of HIV, HBV, and HCV in people who inject drugs: a multistage systematic review. Degenhardt, 2017

3: Epidemiology of hepatitis C virus infection. Alter, 2007

## Aim

- We aimed to estimate the percentage of infections prevented if HCV transmission due to IDU were removed from 2018-2030:
  - Country-level
  - Regionally
  - Globally



# Methods

- A dynamic, deterministic HCV transmission model simulated country-level HCV epidemics among:
  - PWID
  - the general population
- Accounted for:
  - population growth
  - ageing
  - demographics
  - disease progression
  - injecting drug use
  - vertical transmission
  - historical treatment numbers

# Model parameterization



- Demographic information: UN datasets
- Key parameters and bounds from various systematic reviews:
  - HCV prevalence among general population<sup>1</sup>
  - HCV prevalence among PWID<sup>2</sup>
  - Proportion of adults that are PWID<sup>2</sup>
- **Countries included if data were available on all three key parameters**
- Model accounted for uncertainty in parameters

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# Key model assumptions

- Decreasing HCV epidemics among general population (around 1%) per year<sup>1</sup>
  - (due to evidence from countries with 2 surveys)
  - Stable HCV epidemics among PWID<sup>2</sup>
  - Stable proportion of adults that are PWID<sup>2</sup>
    - (except in Eastern Europe and Sub-Saharan Africa: increasing)
  - These assumptions were investigated in many sensitivity analyses
- 

1: Global prevalence and genotype distribution of hepatitis C virus infection in 2015: a modelling study. Blach, 2017

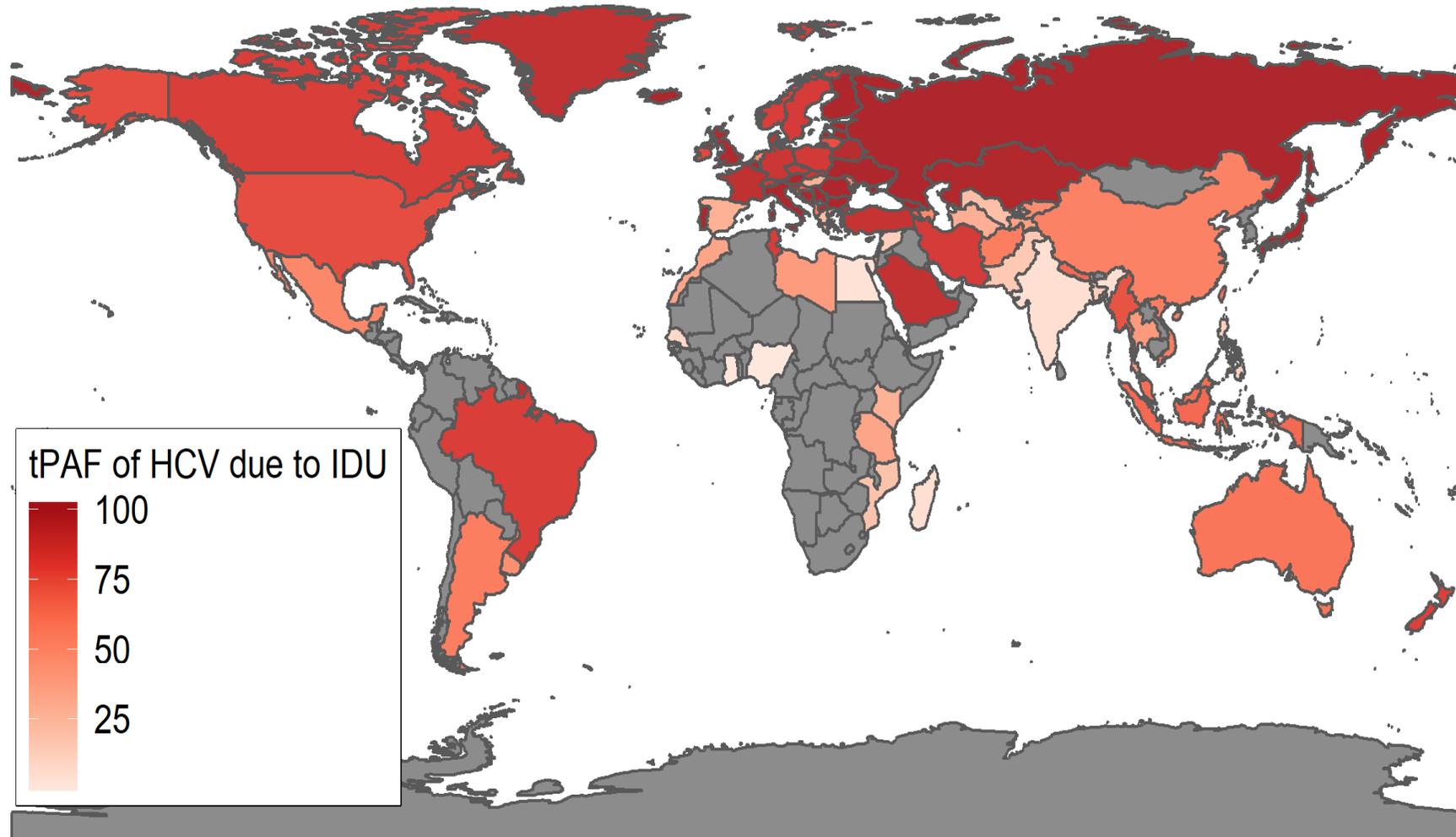
2: Global prevalence of injecting drug use and sociodemographic characteristics and prevalence of HIV, HBV, and HCV in people who inject drugs: a multistage systematic review. Degenhardt, 2017

# Analyses

- Models for each country projected the HCV epidemic to 2030 (baseline projections)
  - 1000 runs for each country
- The population attributable fraction of HCV transmission (incidence) due to IDU in each country, regionally, and globally, was estimated – the **tPAF**
  - This is the percentage of transmission that would be avoided if the extra transmission due to IDU removed
- To do this, the baseline model fits for each country were re-run with the transmission risk due to IDU set to zero from 2018 onwards
  - Compared vs baseline projections
- Generalised linear regression models used to determine what country-level factors are associated with the tPAF of HCV due to IDU

# Contribution of IDU to HCV 2018-2030

- Successfully calibrated for all 88 countries with data (85% of world's population)

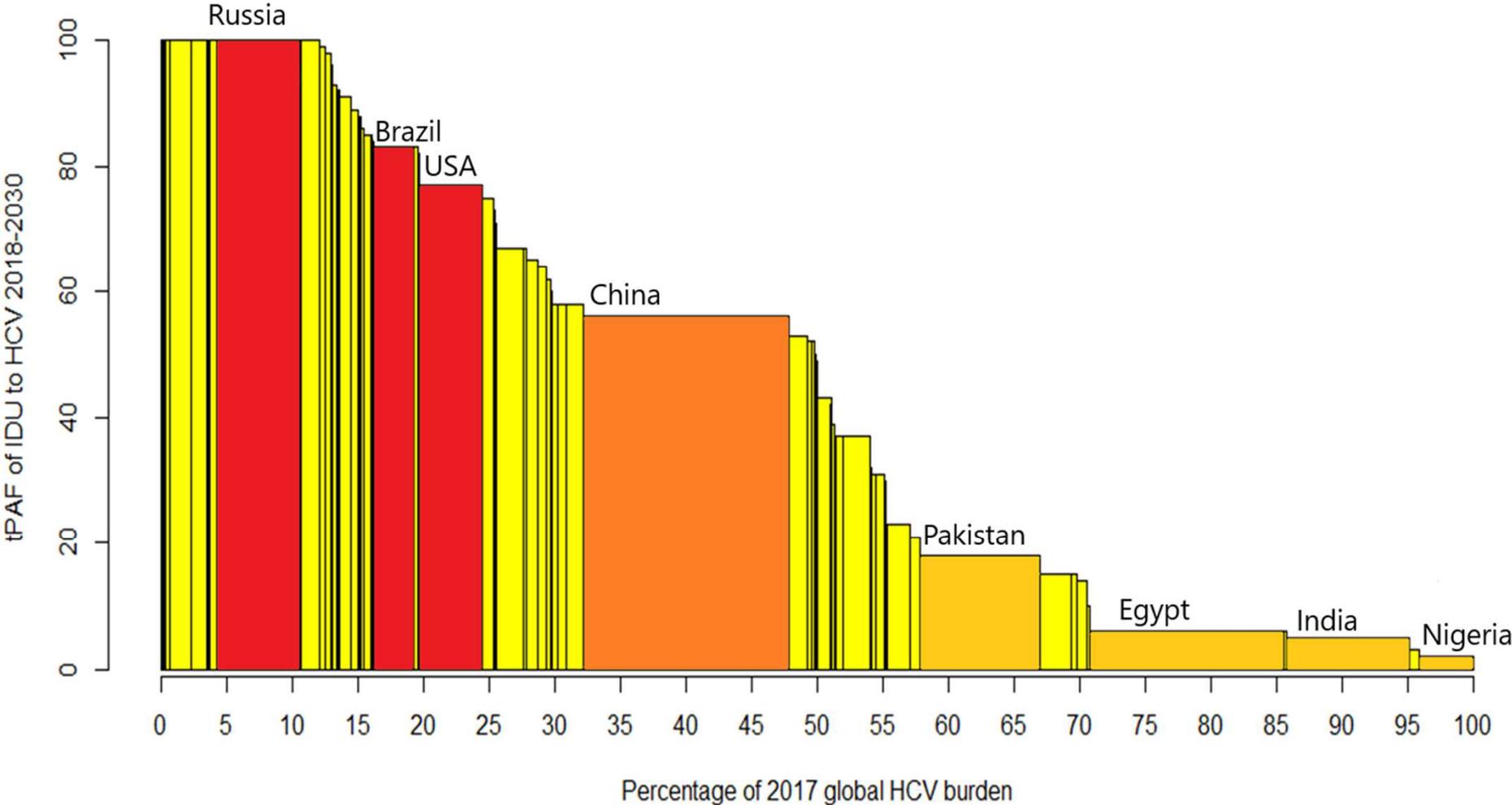


\*Countries in grey were not included in the model due to missing data

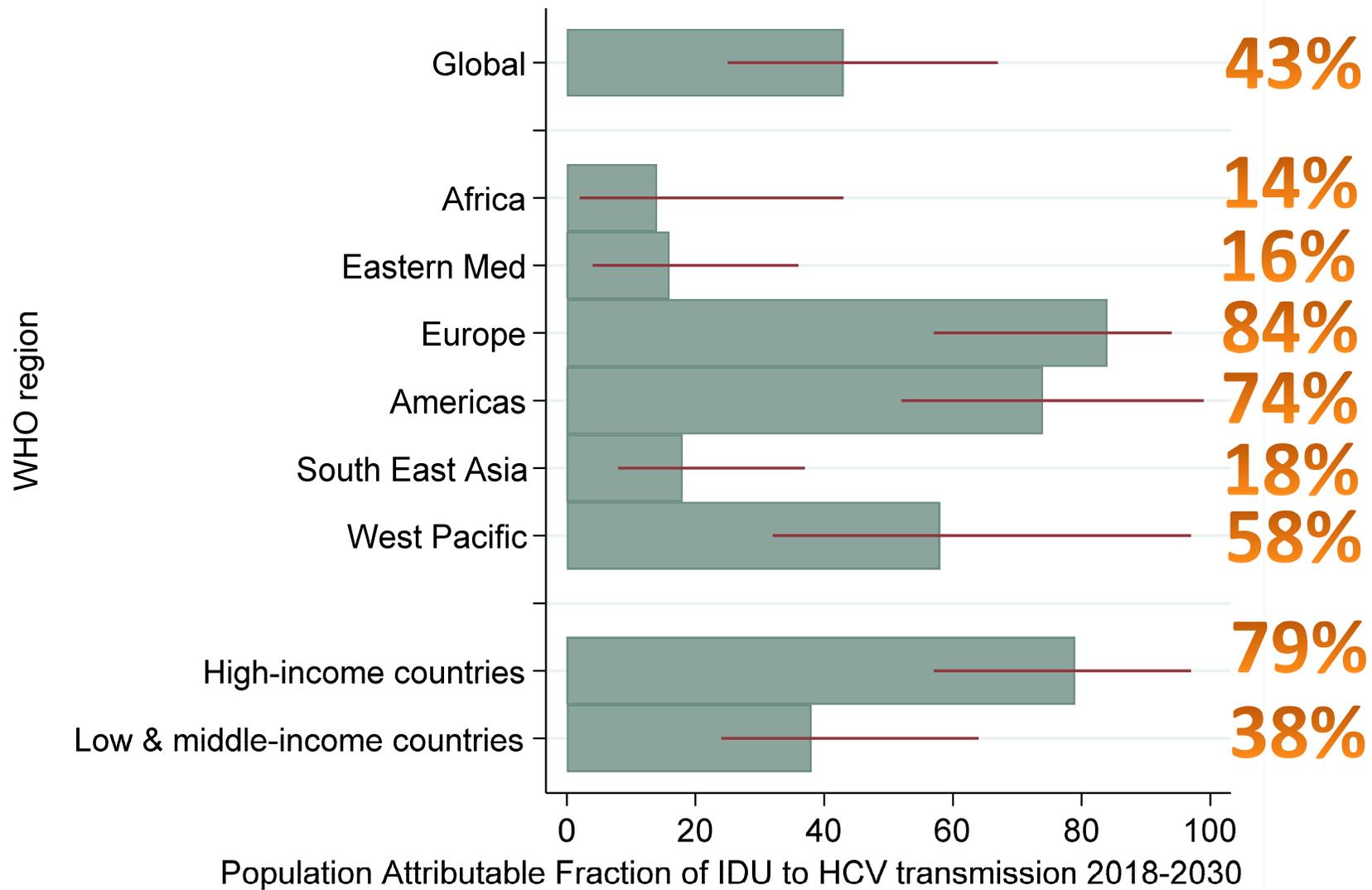
# Each country's percentage of global HCV burden (2017)

vs

# Contribution of IDU to HCV transmission 2018-2030 (tPAF)

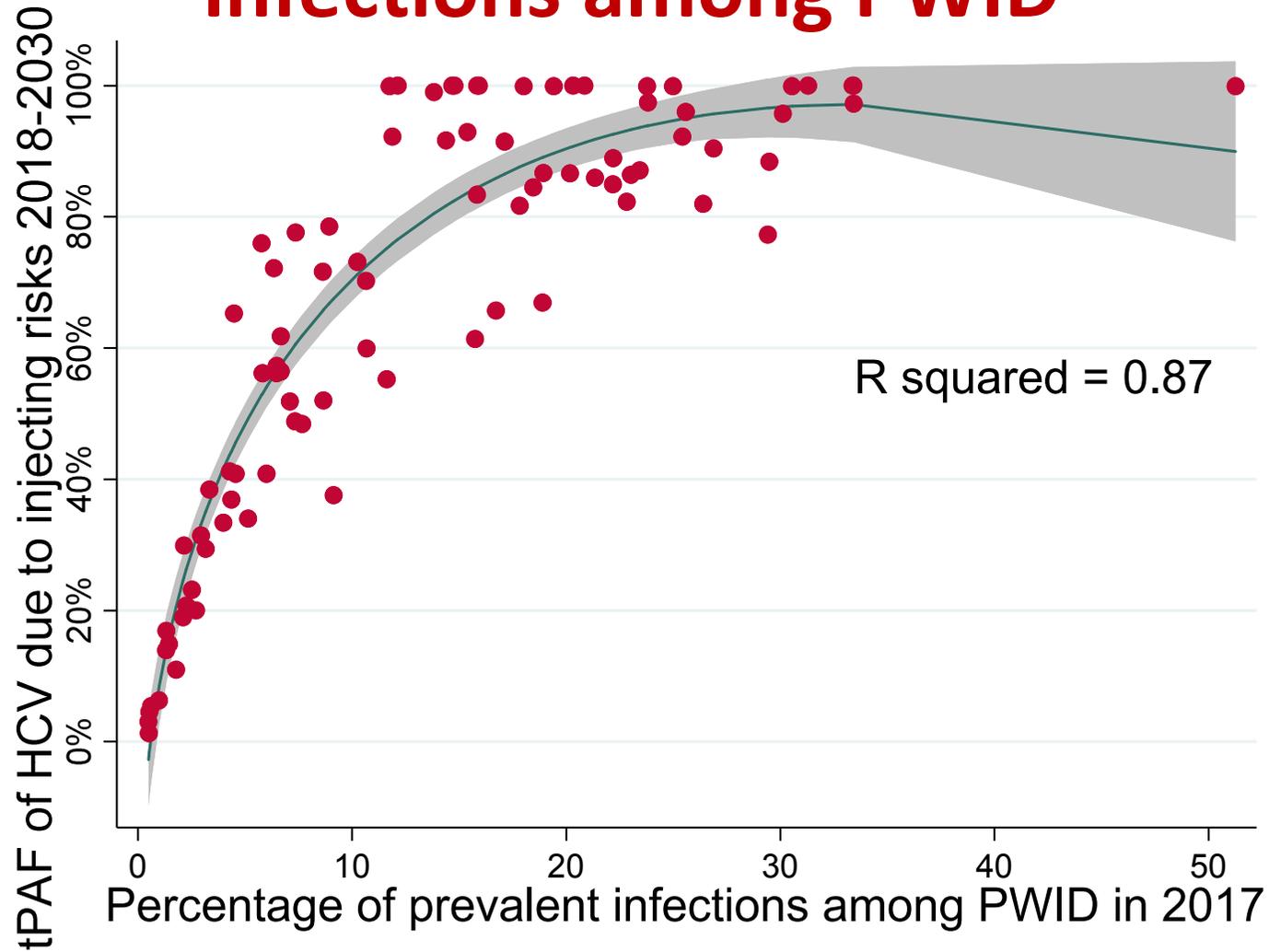


# Regional contribution of IDU to HCV 2018-2030



\*Red lines are 95% credibility intervals

# Association with current percentage of infections among PWID



**No other multivariable associations found with the tPAF**

# Sensitivity analyses

- Various sensitivity analyses tested assumptions
- Most had very little effect on the tPAF
- Largest changes occurred when:
  - Assuming a stable general population HCV prevalence over time (33% tPAF)
  - Varying epidemic trajectories by region (30% tPAF)



## Limitations

- Taking data from disparate sources can create imprecise results but overall trends should be robust
- **Data!**
  - Data unavailable for many countries (particularly Africa)
  - Not necessarily high quality data
- Migration not included – lack of data
- Assumptions about directions of epidemics:
  - Only 3 countries had 2 robust, comparable general population estimates
  - Investigated in sensitivity analyses

# Key results and conclusions

- **43% (25%-67%)** of incident HCV infections would be prevented from 2018-2030, globally, from removing all extra transmission due to IDU
- Varies regionally
- **79%** in high-income countries
- **38%** in low- and middle-income countries
- Unsafe injecting practices among PWID contribute substantially to incident infections globally
- Any intervention that can reduce transmission among PWID will have a pronounced effect on country-level incidence
  - OST, NSP, treatment as prevention



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