



## **HIV in young women in KwaZulu-Natal: Imperative for their Inclusion in HIV Biomedical research studies**

Presented by: **Ayesha BM Kharsany**



# **HIV in young women in KwaZulu-Natal: Imperative for their Inclusion in HIV Biomedical research studies**

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# Overview of Presentation

- Research priorities, studies and Rationale
- Surveillance
- Phylogenetic studies
- HIV Transmission Networks
- Genital Inflammation
- Vaginal flora
- Conclusions

# CAPRISA Research Priorities

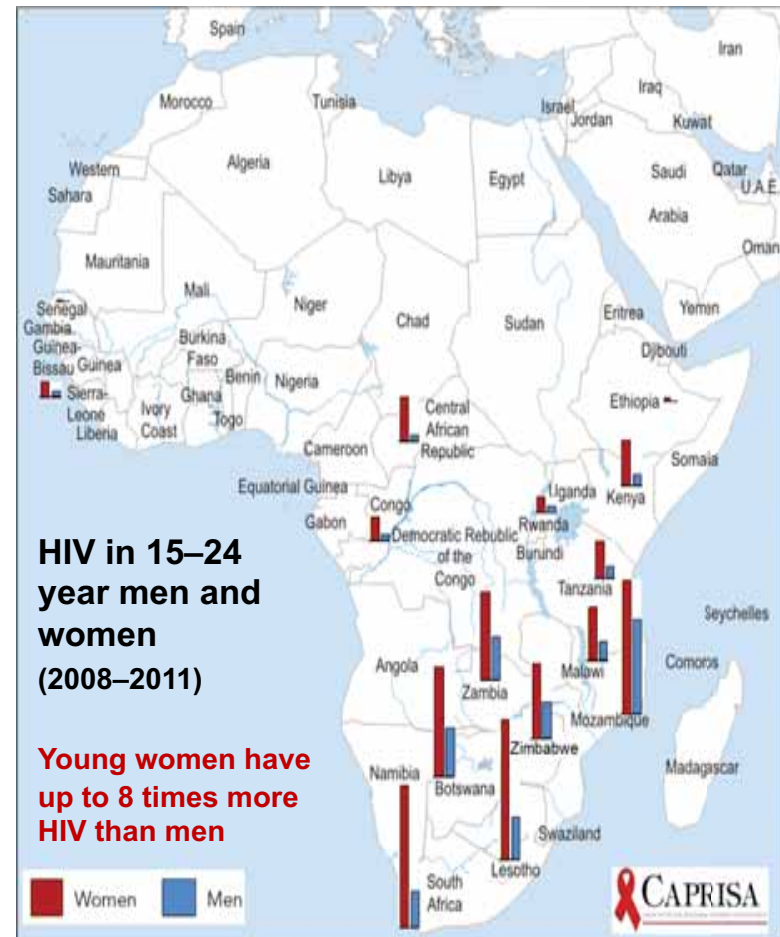
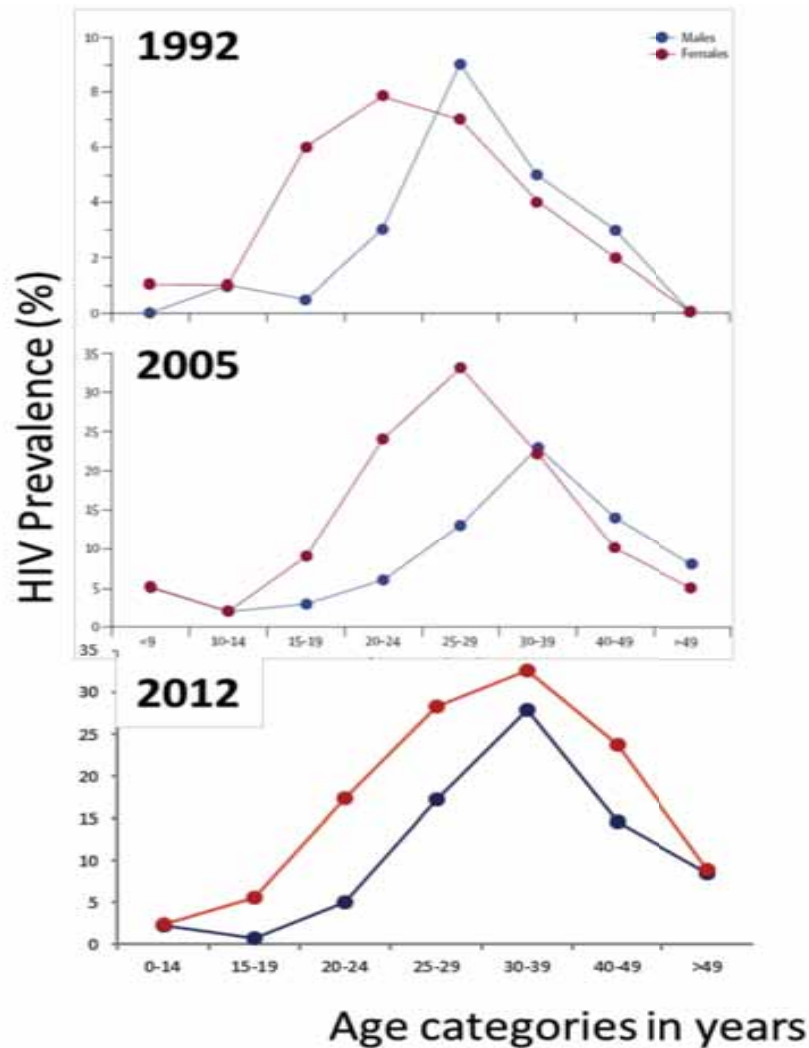
To undertake *globally relevant & locally responsive* research that contributes to understanding HIV pathogenesis, prevention & epidemiology as well as TB-HIV treatment

Understand the epidemiology of HIV in young women


# HIV in South Africa

## Disproportionate burden of HIV in young women

Southern & eastern Africa  
have 70% of the world's HIV  
burden



# Trends in HIV in pregnant women in rural KZN 2001-2013



Age Group (Yrs)	HIV Prevalence (n=4818)
≤16	11.5%
17-18	21.3%
19-20	30.4%
21-22	39.4%
23-24	49.5%
>25	51.9%

Age group (Yrs)	HIV Prevalence %		
	Pre-ART 2001-2003	Early ART 2004-2008	Current ART 2009-2013
<20	22.5	20.7	17.2
20-24	45.5	44.2	37.9
25-29	47.9	58.8	57.6
30-34	26.7	55.5	59.9
≥35	27.4	30.8	53.4
<b>Total</b>	<b>35.3</b>	<b>39.0</b>	<b>39.3</b>

## Key findings

- Pregnant women <20 years old  
>30%
- HIV prevalence consistently high  
30% and 40%  
by age 24; 1 in 2 HIV positive
- HIV prevalence in the <20 year age group  
>20%
- Increase in partner's age  
Increase in HIV prevalence in young women
- HIV incidence  
11.2%

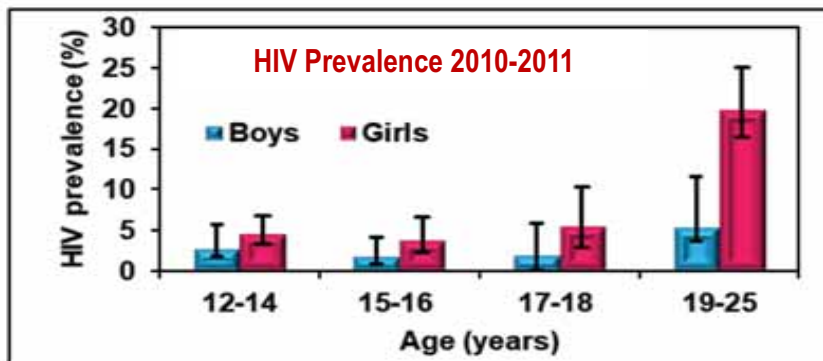
Source: Kharsany AB et al JAIDS 2015,70:289-295.



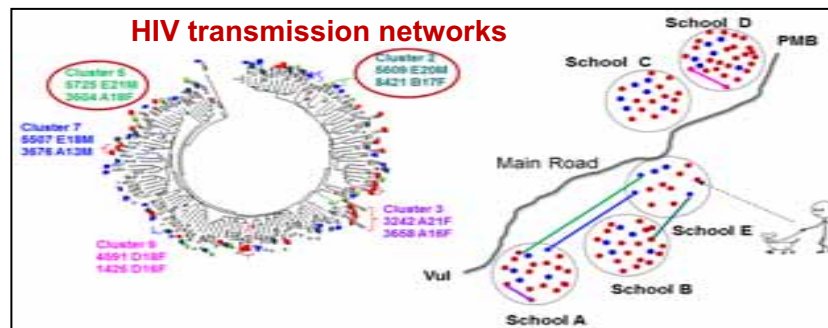
# Trends in HIV in high school students in rural KZN

Age Group (Yrs)	HIV Prevalence 2010 (% 95% CI)	
	Boys (n=1262)	Girls (n=1423)
≤15	1.0 (0.0 - 2.2)	2.6 (1.2 - 4.0)
16-17	1.1 (0.2 - 2.0)	6.1 (2.6 - 9.6)
18-19	1.5 (0 - 3.7)	13.6 (9.0 - 18.1)
≥20	1.8 (0 - 3.9)	24.7 (6.3 - 43.1)
<b>Overall</b>	<b>1.4 (0.9 - 1.9)</b>	<b>6.4 (4.6 - 8.3)</b>

Source: Abdool Karim Q, et al. Sexually Transmitted Infections 2014, 90:620-626.



Source: Kharsany AB et al BMC Public Health 2012,12:231



Source: Kharsany AB. et al AIDS Research and Human Retroviruses 2014,30:956-965

## Key findings

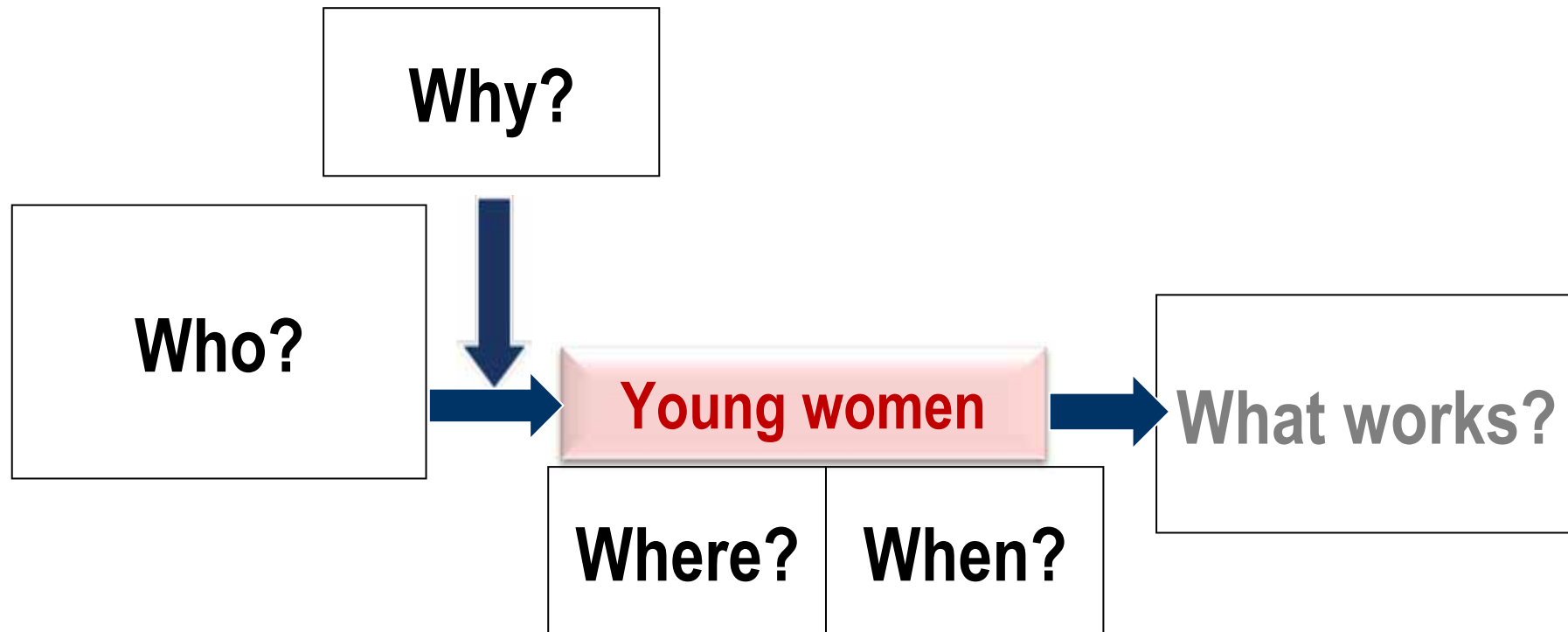
- Learners were sexually active  
1 in 4
- HIV prevalence  
High across all ages  
Higher in girls vs boys
- Girls with older sex partners  
3 to 4 times more likely to be HIV +
- Clues on HIV transmission  
Limited in and across schools
- Possible introductions  
Community

# Challenges to understanding HIV acquisition in adolescent girls and young women

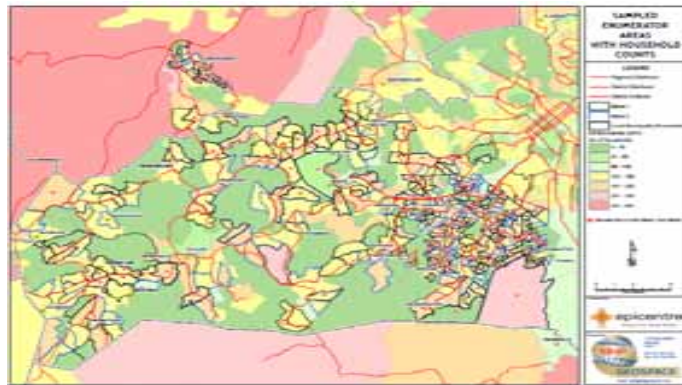
- Biological and structural factors
- Risk behaviors of young women
  - Low sexual frequency but high risk sex
  - Few have multiple concurrent partners
  - Why?
- Age-disparate relationships as a key contributor to HIV acquisition
  - Evidence
    - Consistent in CAPRISA studies
    - Conflicting with literature
      - Bias with study design
      - Analysis bias
      - Reliability of self reported data
      - Definitions



# Young women at high HIV risk



# Community Based Surveillance

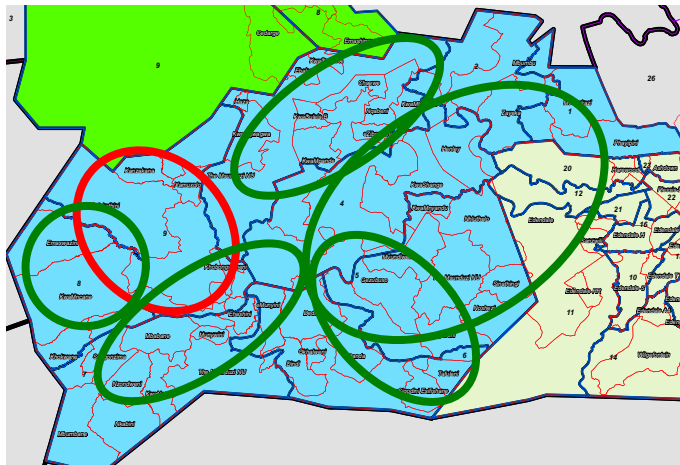
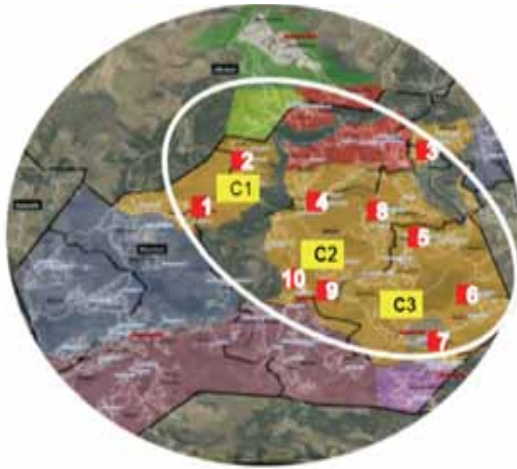


- **Purpose:** Evaluate population impact of programmatic scale-up of combination prevention efforts in a “real world”, non-trial setting in rural KZN
- **Study objectives:** Measure HIV prevalence and incidence in relation to ART and plasma viral load
- **Study Location:** 2 Sub-districts of uMgungundlovu district, KZN, SA
  - Population of ~370 000
  - Males ~176 418 / Females ~191 515
  - High levels Poverty / Unemployment
  - Highest HIV burden district in SA
  - (SA DOH\_ANC prevalence-40.7% in 2012)

Source: Kharsany AB et al. BMC Public Health 2015,15:1149.

# School Based Surveillance

## CANS Adolescent Network studies



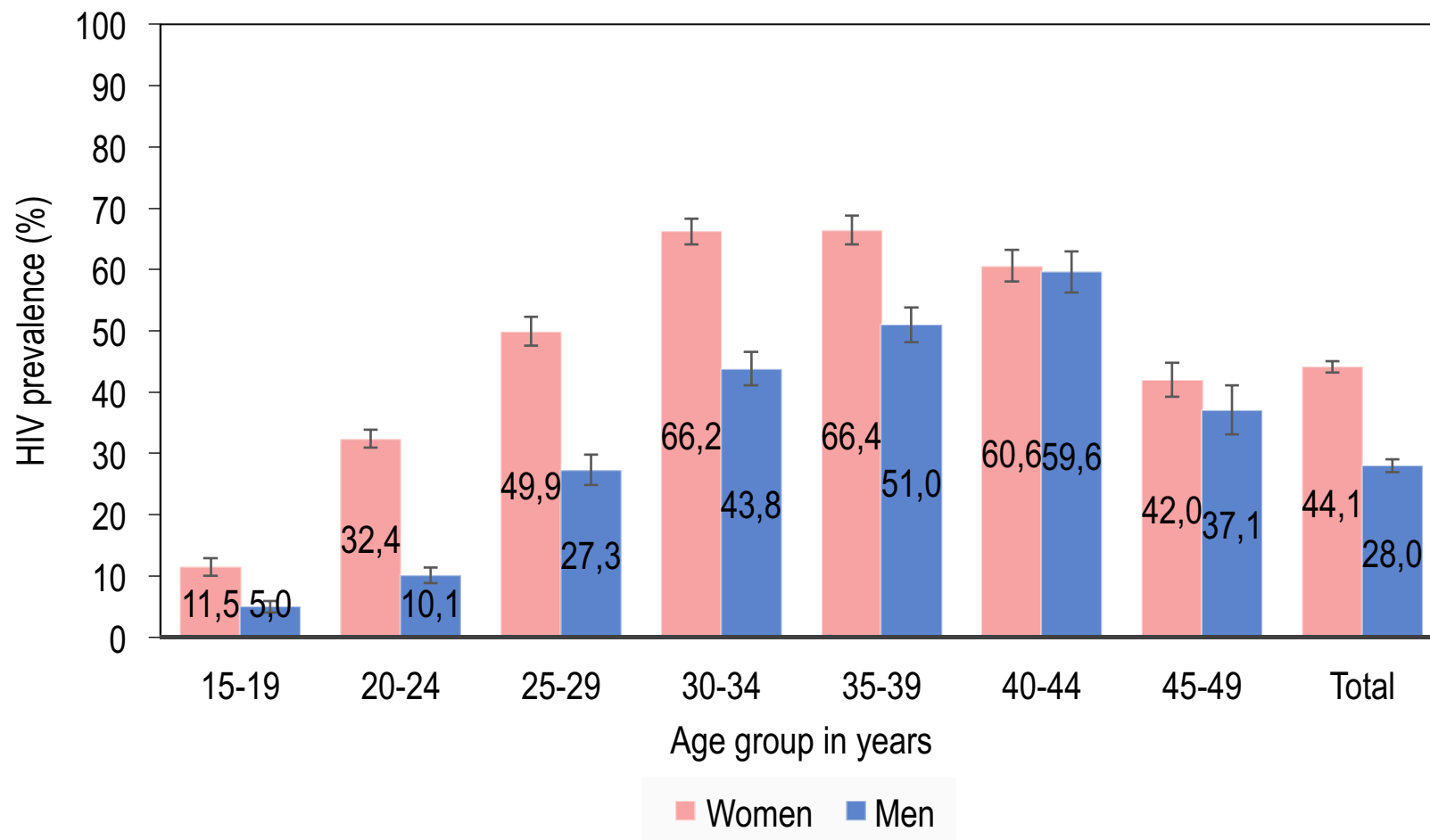
- **Purpose:** To understand the HIV transmission dynamics and sexual networks that adolescent may belong to, potentially increasing their risk for HIV infection.
- **Study objectives:** To identify HIV transmission linkages that contribute to the source of infection in adolescent girls in rural KZN
- **Study Sites :** Specified geographical location to maximise density and coverage
  - All high schools  $n=+/-8000$
  - Facilities – out of school pregnant girls  $n=+/-1000$
  - Out of school high risk adolescent boys (RDS)  $n=+/- 1000$ .

Source: Kharsany et al :Identifying sources of HIV infection in adolescent girls in rural South Africa National Institutes of Health, RO1, R01HD083343 (Multi-PI: Kharsany and Kohler).

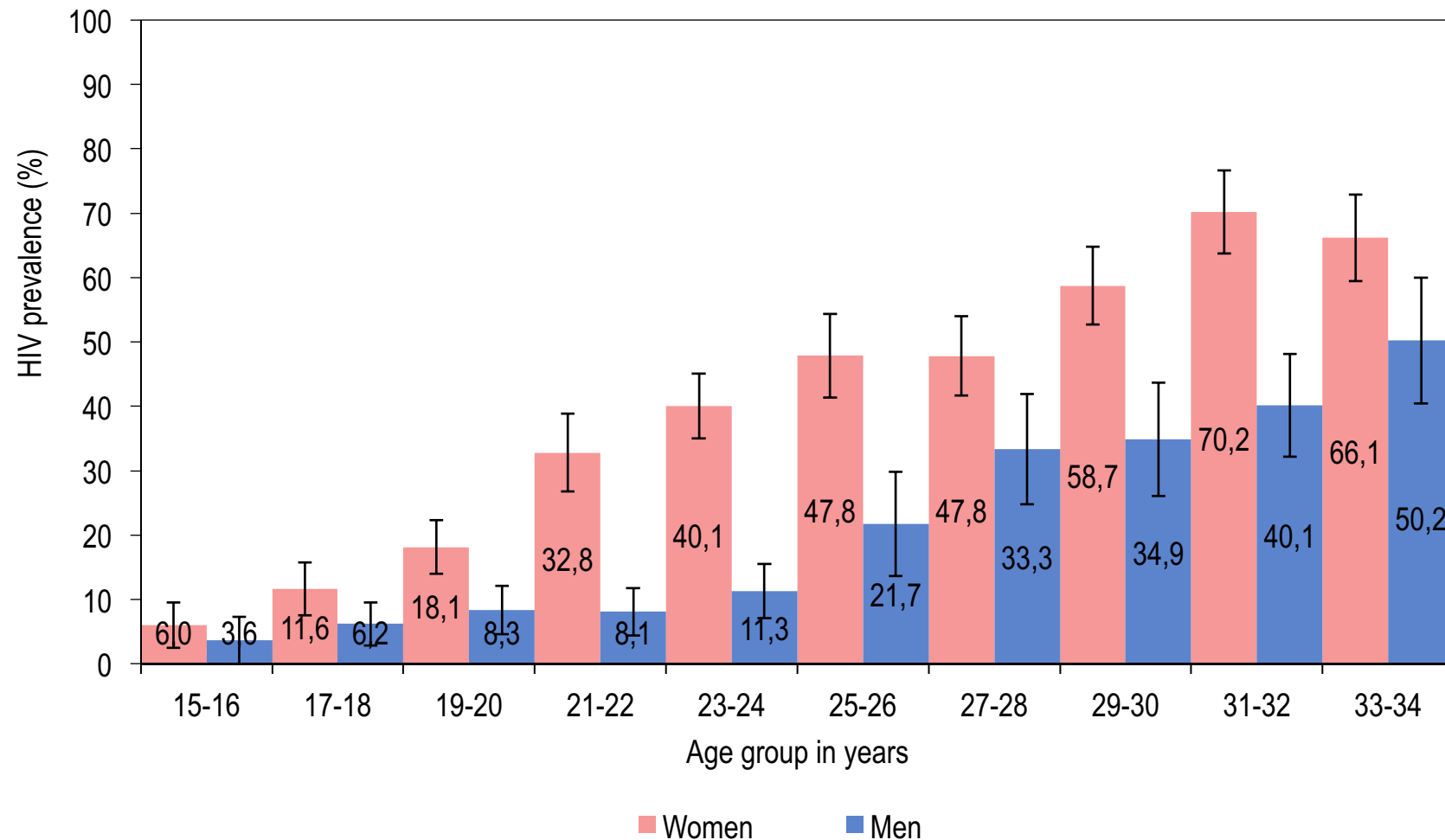
# Community-wide phylogenetic study

- Cross-sectional multi-stage random sampling
- Duration: 2014 - 2016
- 86% consent rate
- People tested for HIV: 9 812
- HIV positive: 36.3% (CI: 35-38) (n=3,969)
- Knew HIV+ status: 59.8% (n=2,337)
- On ARVs: 42.3% (n= 1,590)
- Viral load >1000: 47.1% (n= 1,847)

# HIV sero-prevalence by age and gender in a rural and peri-urban community in KZN, South Africa, 2014-2015

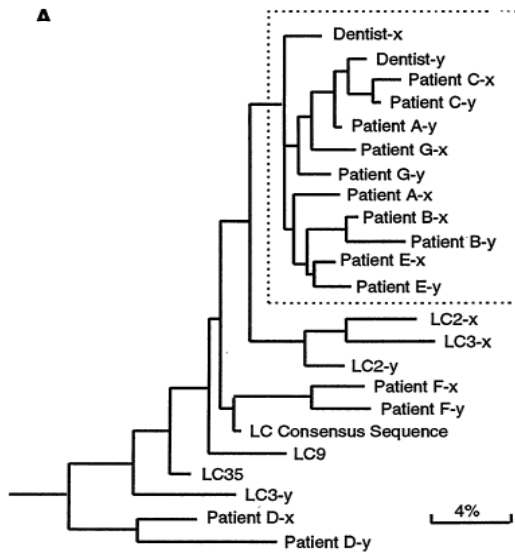


# HIV sero-prevalence in young men and women in a rural and peri-urban community in KZN, South Africa, 2014-2015



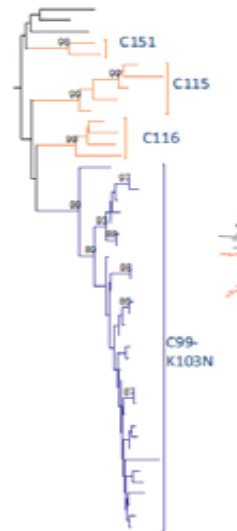


# Phylogenetic analyses to assess HIV genetic diversity for identifying linked viruses



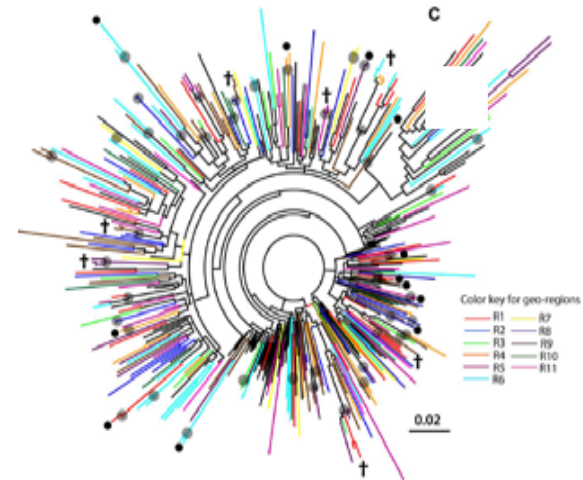
**Individual linkage analysis**  
Transmission linkages in 5 patients who became infected with HIV while receiving care from a dentist with AIDS

Source: Ou CY et al : Science 1992, 256:1165-1171.



**MSM epidemic in Montreal**  
Unique transmission clusters three small clusters and one large cluster with K103N resistance mutation

Source: Brenner B et al AIDS. 2013;27(7): 1045-57.

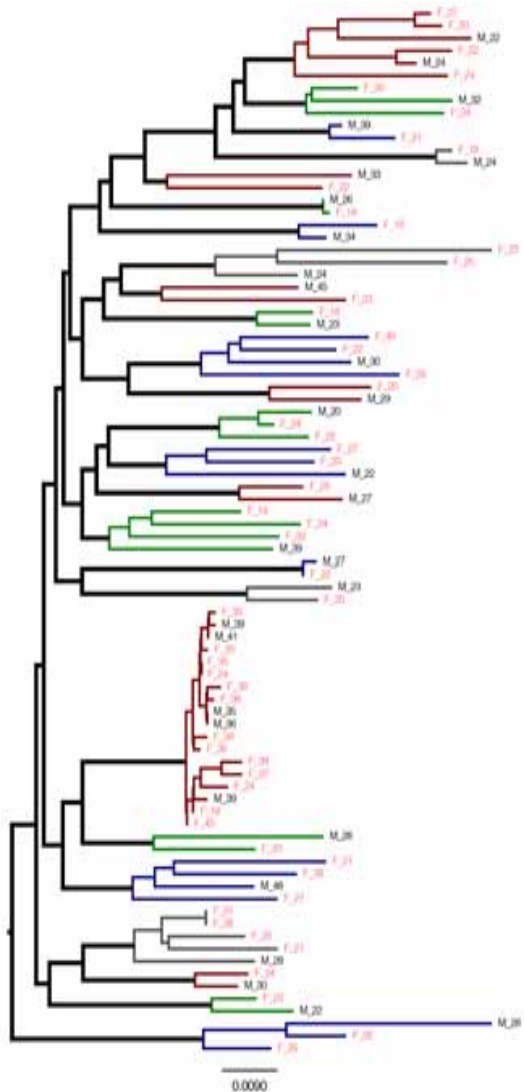


**Rakai, Uganda**

Significant viral introductions  
39% within households  
40% from extra household contacts / outside communities.  
Localised key population drive the epidemic

Source: Grabowski MK, et al. PLoS Med. 2014;11(3):e1001610.

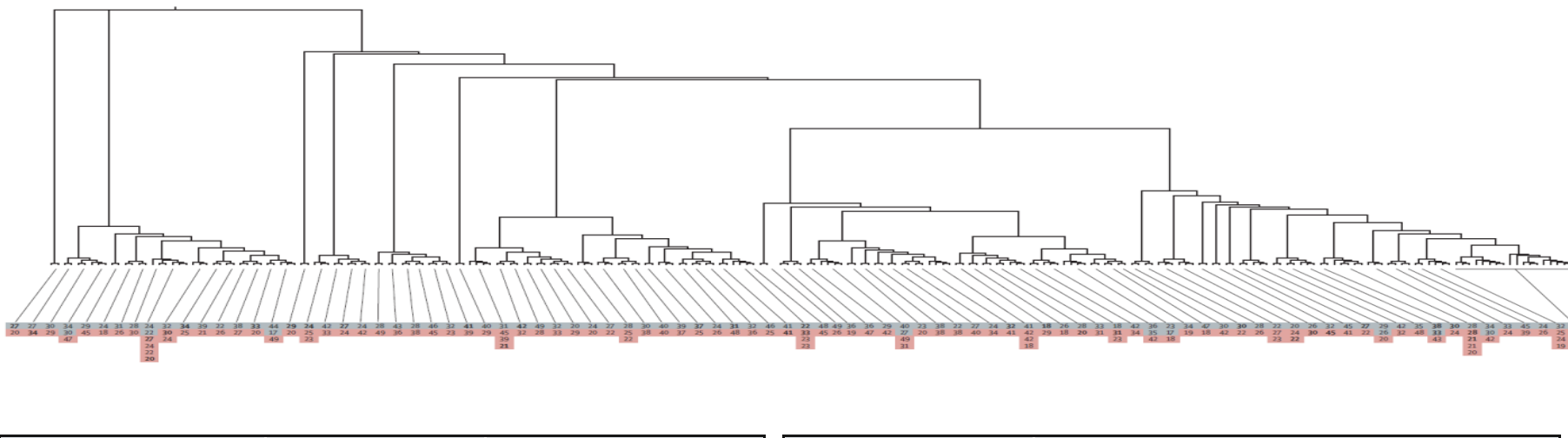
# Phylogenetics to map HIV transmission



- **1,589 of 1847 viruses sequenced**
- ***pol* region**
- **Maximum likelihood tree**  
**Branch support > 90%**  
**Genetic diversity < 4,5%**
- **Phylogenetic linkage = probable recent transmission as those with more established infection were on ART**
- **202 phylogenetically linked clusters of 469 sequences**
  - 168 with 2 individuals
  - 22 with 3 individuals
  - 8 with 4 individuals
  - 1 with 5 individuals
  - 2 with 6 individuals
  - 1 with 18 individuals

# Phylogenetic analysis: cluster linkages

90 clusters (Male-Female linkages) –123 females linked to 103 males (188 possible pairings)



	Women	Men
Median VL	16,629c/ml	31,000 c/ml
on ARVs	4.1%	4.9%
Known HIV+	38.9%	26.2%

Women age group	Age difference with male partners
16-20	11.5 yrs
21-25	7.0 yrs
26-30	1.5 yrs
31-35	1.7 yrs
36-40	0.7 yrs

< 25: 8.7 yrs

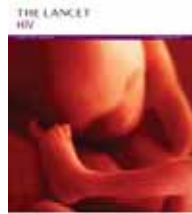
≥ 25: 1.1 yrs

# Phylogenetic pairing and HIV-1 prevalence

Females Community HIV prevalence	Males Community HIV prevalence			Total
	<25yrs (7.6%)	25-40yrs (40.3%)	40-49yrs (47.2%)	
<25 yrs (22.3%)	18	37	5	60
25-40 yrs (59.8%)	13	58	21	92
40-49 yrs (50.1%)	2	24	10	36
<b>Total</b>	33	<b>119</b>	36	188

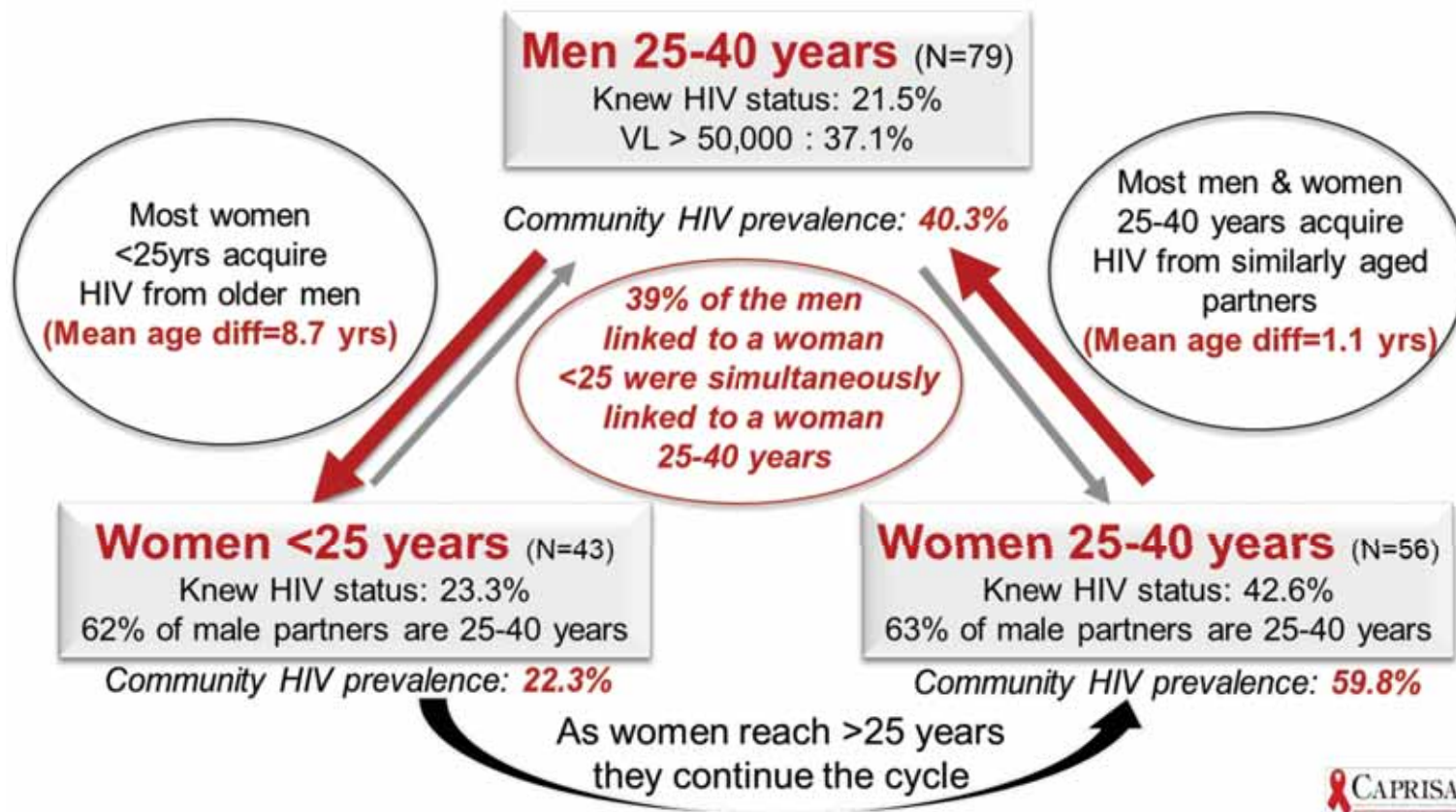
- Pairing across all age groups, but the highest linkage is of men 25-40 (group that has the highest prevalence of viraemia including PVL >50 000 c/ml)
- HIV-1 transmission occurs from high-prevalence to low prevalence, unless there are super-spreaders

# Why: HIV transmission networks in rural South Africa



## Transmission networks and risk of HIV infection in KwaZulu-Natal, South Africa: a community-wide phylogenetic study

Tulio de Oliveira\*, Ayesha B M Kharsany\*, Tiago Gräf, Cherie Cawood, David Khanyile, Anneke Grobler, Adrian Puren, Savathree Madurai, Cheryl Baxter, Quarraisha Abdool Karim, Salim S Abdool Karim

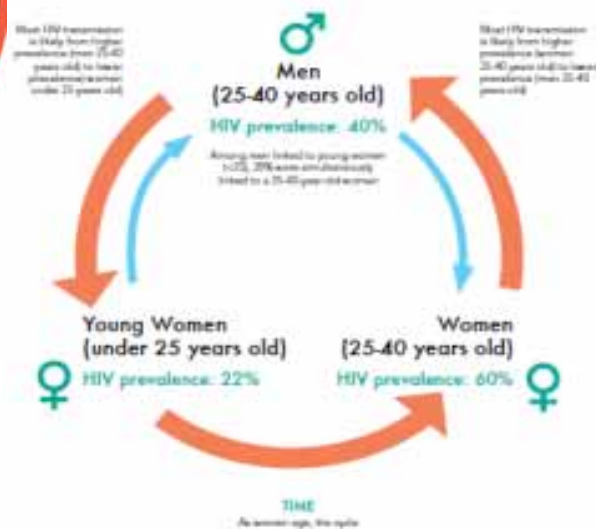




# UNAIDS AIDS Update 2016

## Get on the Fast-Track

### The Life-cycle approach to HIV



# LET OUR ACTIONS COUNT

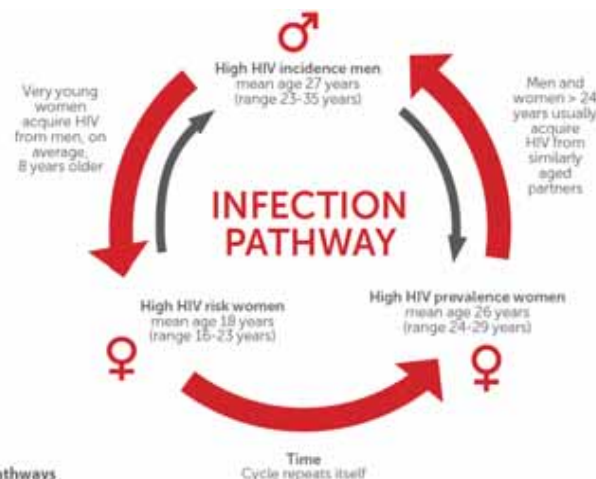
## SOUTH AFRICA'S NATIONAL STRATEGIC PLAN ON HIV, TB and STIs 2017-2022



**Goal 1:**  
Accelerate prevention in order to reduce new HIV and TB infections and new STIs.  
- Breaking the cycle of transmission



The NSP sets out intensified prevention programmes that combine biomedical prevention methods, such as medical male circumcision (MMC) and the preventive use of antiretroviral drugs (ARVs) and TB medication, with communication designed to educate and encourage safer sexual behaviour in the case of HIV and STIs, and environmental interventions to control TB infection.



Transmission pathways

Time Cycle repeats itself

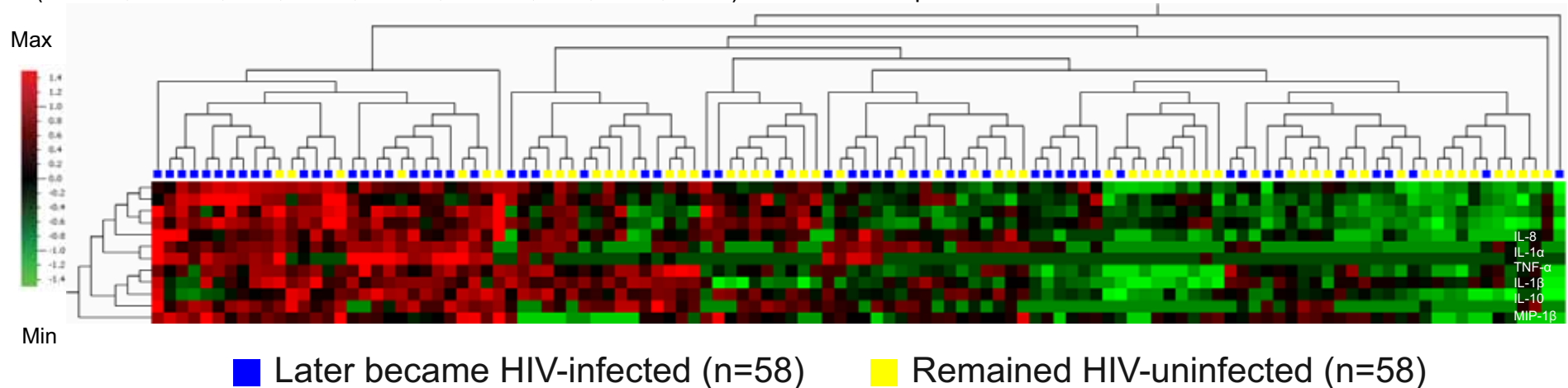
## Important public health benefit



# Why: Vaginal Microbiome : genital inflammation

## Cytokine and chemokine responses

(MIP-1a, MIP-1b, IL-8, IP-10, TNF-a, MCP-1, IL-6, IL-1a, IL-1b) above the 75<sup>th</sup> percentile



**Women who later became HIV-infected had pre-infection genital inflammation – what is the cause?**

Only **20%** of HIV infections could be attributed to an STI  
*T. vaginalis* was the most strongly predictive of genital inflammation

Source: Masson L, Passmore AJ, et al. *J Infect Dis* 2015; **212**:200-208

# Association between genital inflammation and HIV acquisition

	HIV+	HIV-	Total
Genital inflammation present*	19	6	25
Genital inflammation absent	39	52	91
Total	58	58	116

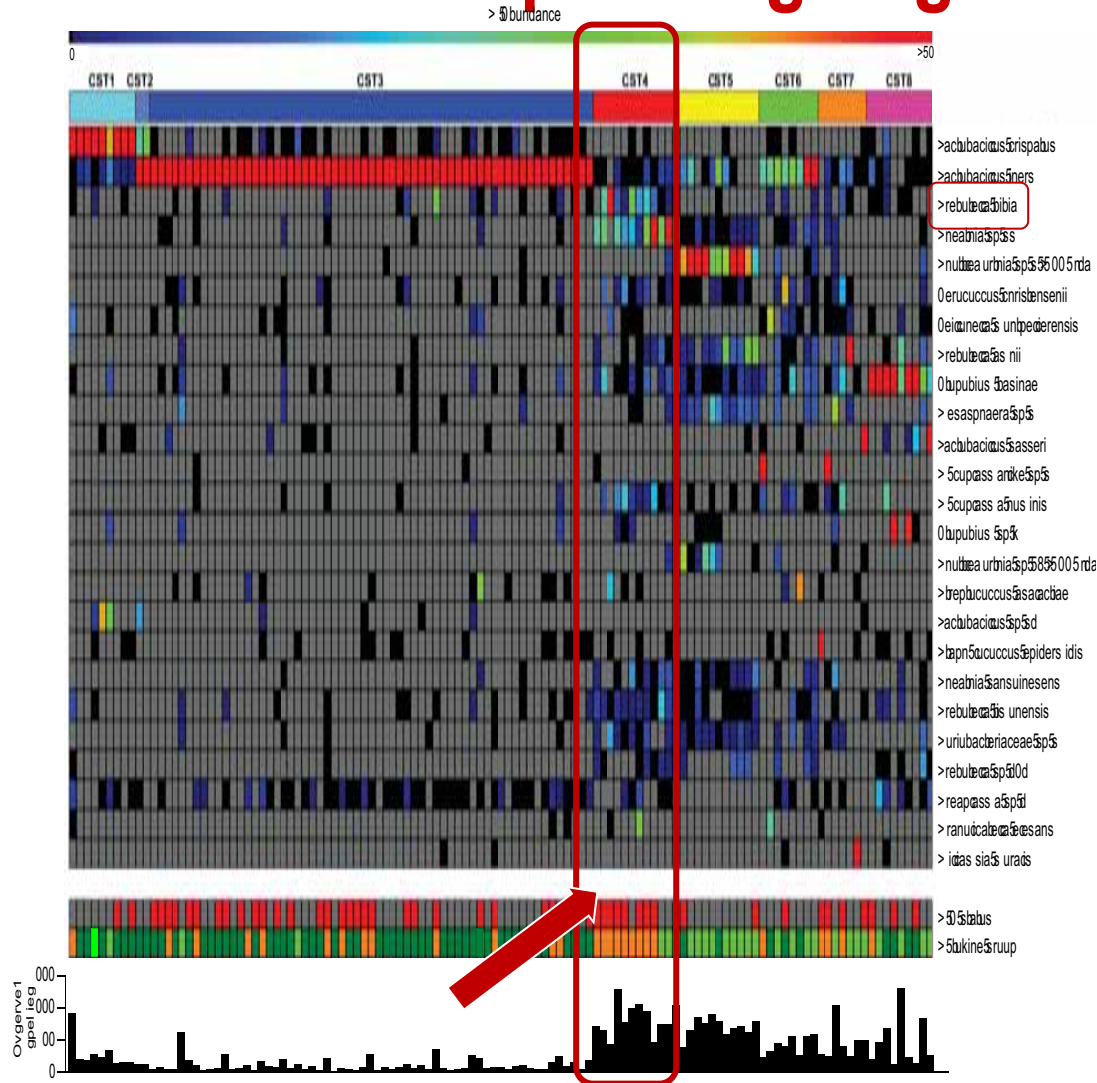
**Odds Ratio 3.2 (95% CI: 1.3 – 7.9), p-value 0.014**

\*Women with 5 or more **pro-inflammatory cytokines or chemokines** (MIP-1a, MIP-1b, IL-8, IP-10, TNF-a, MCP-1, IL-6, IL-1a, IL-1b) above the 75<sup>th</sup> percentile

Significant after adjusting for age, urban/rural, condom use, hormonal contraceptives, number of sex acts, number of returned used applicators, HSV-2 status

# Vaginal Microbiome

## 16s DNA sequencing of genital swab samples



Over 3 million sequences

1368 species identified

- Predominance of Cluster Community State Type (CST) 4 related organisms
- *Prevotella bivia*
- Linked with
  - genital inflammation
  - HIV acquisition

## ***Prevotella bivia* is strongly associated with genital inflammation and HIV acquisition**

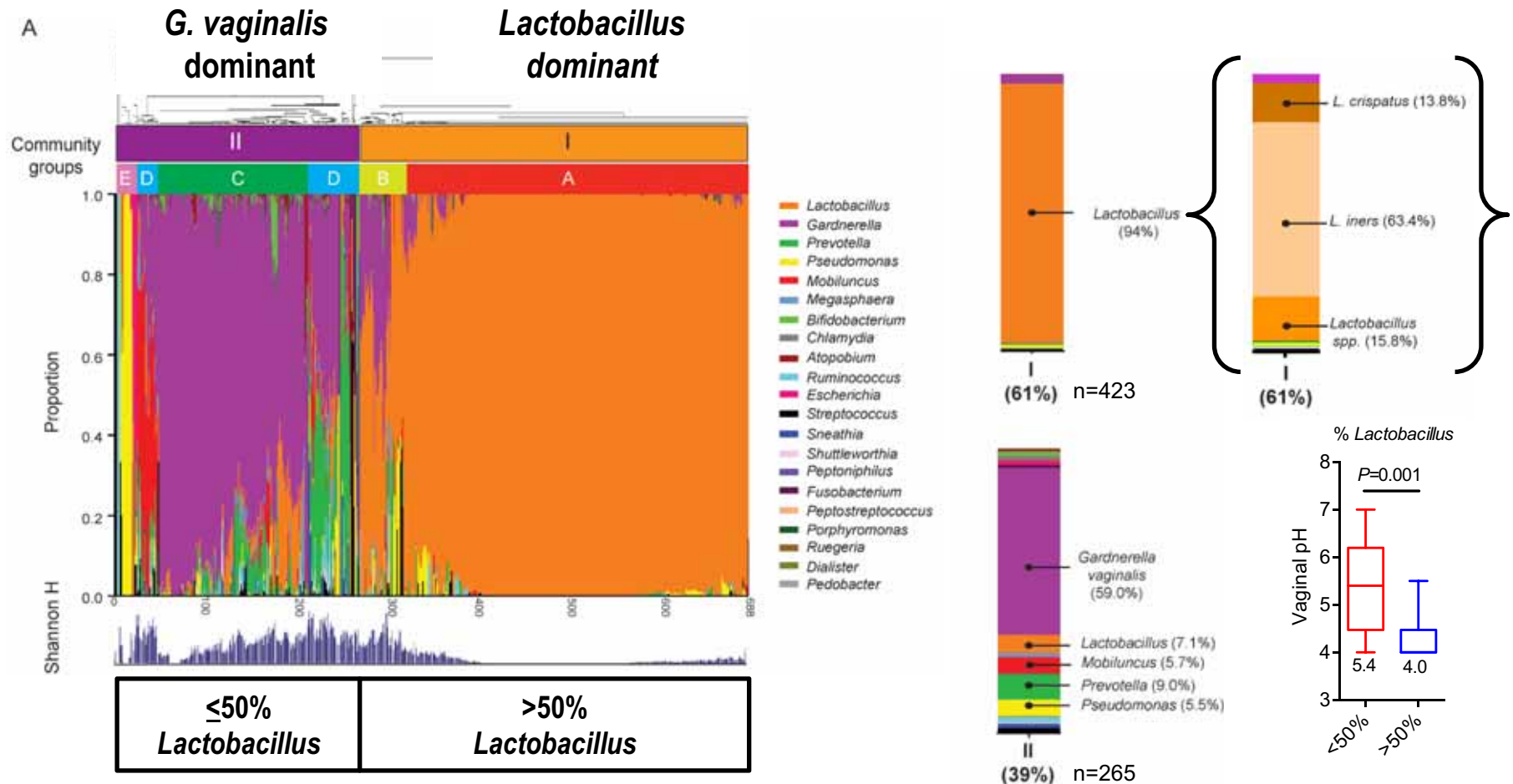
	<b><i>P. bivia</i>+ OR*</b>	<b>P value</b>
GI	19.2 (95% CI: 4.0-92.4)	p<0.001
HIV+	12.7 (95% CI: 2.1-77.8)	p=0.006

*\*adjusted odds ratio*

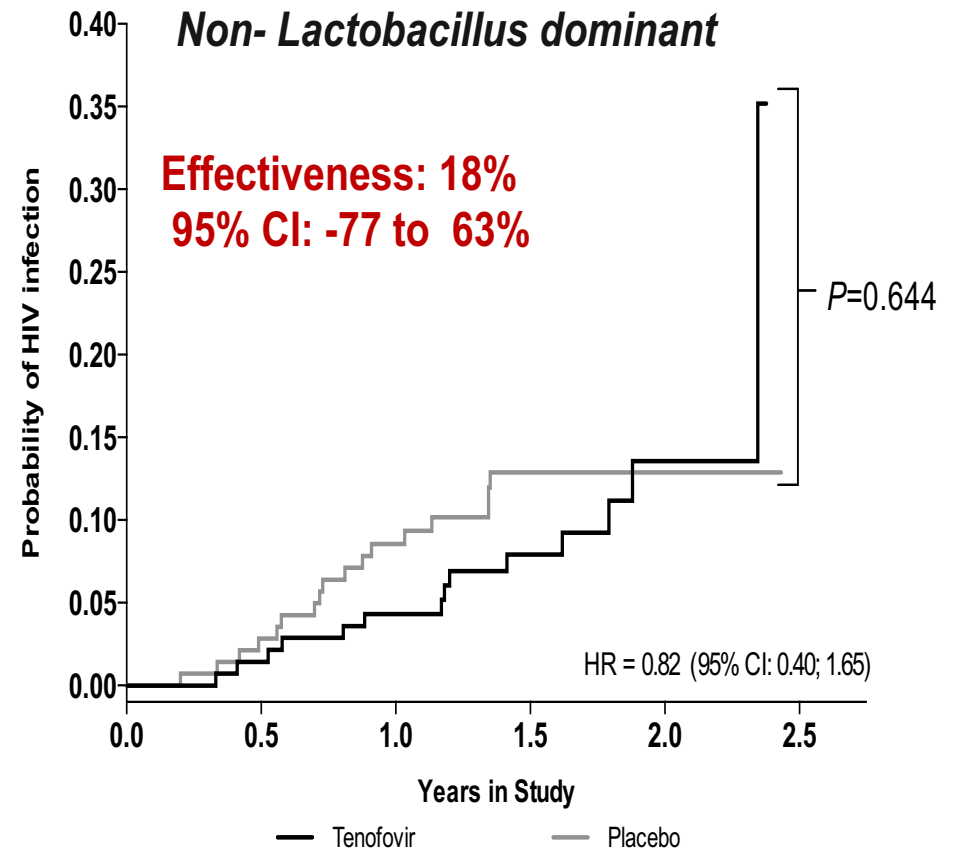
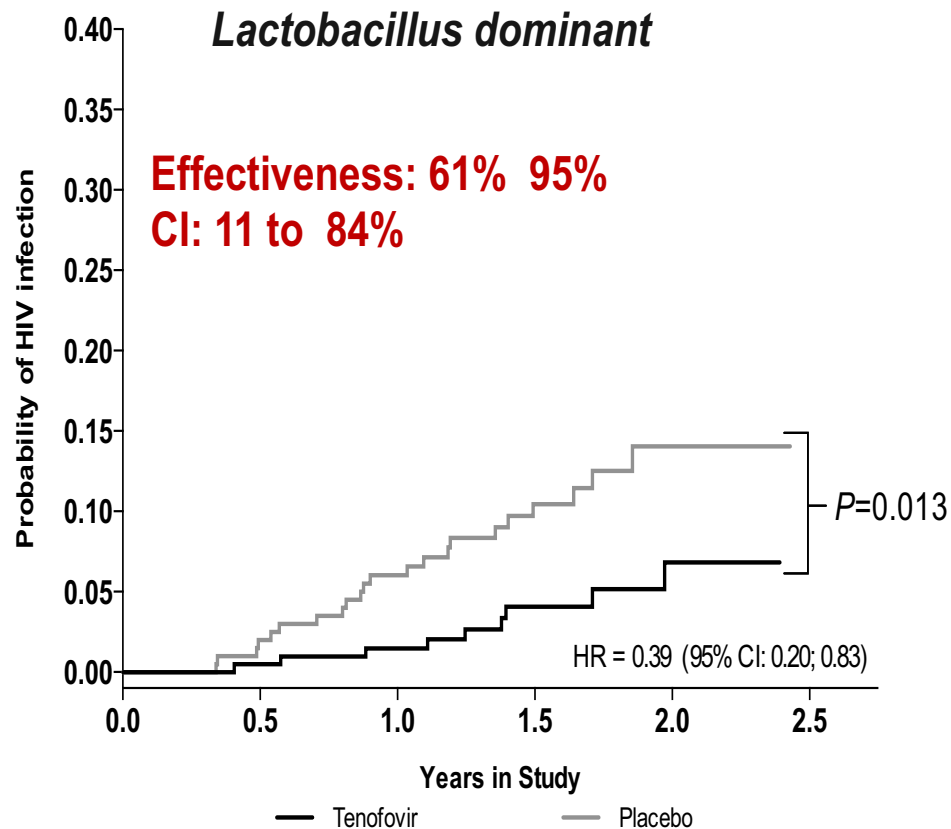
**22 women were HIV positive & had inflammation  
– 9/22 (41%) had *P. bivia***

Women with *P. bivia* were **19 times** more likely to have genital inflammation and  
**13 times** more likely to acquire HIV

# Proteomics: Identification of Vaginal microbial groups

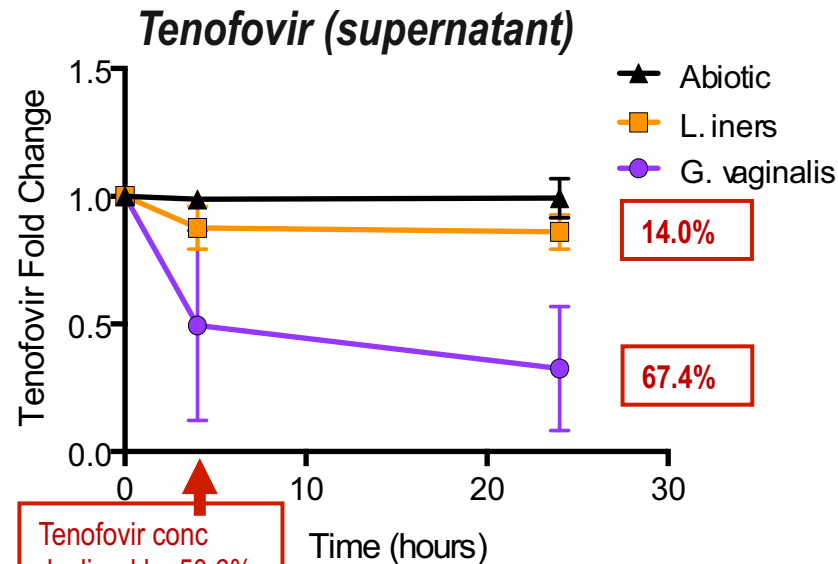


# Effectiveness of Tenofovir gel against HIV amongst women with *Lactobacillus* dominance





# Tenofovir rapidly depleted by *Gardnerella* but not by *Lactobacillus*



Tenofovir conc declined by 50.6% within 4 hrs

**4 hours:**

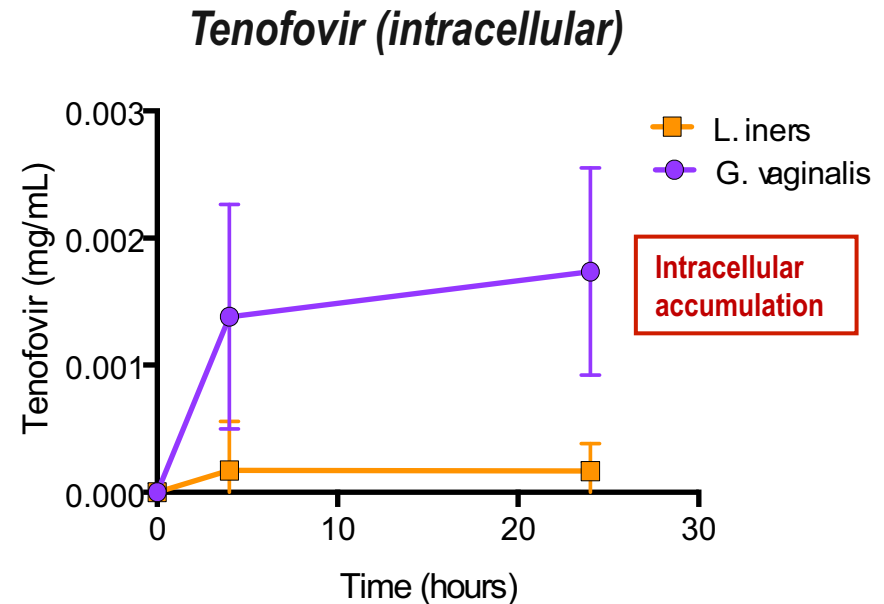
G. vag vs. L. iners:  $P=0.002$

G. vag vs Abiotic:  $P=0.005$

**24 hours:**

G. vag vs. L. iners:  $P<0.001$

G. vag vs Abiotic:  $P<0.001$



**4 hours:**

G. vag vs. L. iners:  $P<0.001$

**24 hours:**

G. vag vs. L. iners:  $P<0.001$

## Bio-degradation of tenofovir by *Gardnerella*

# Conclusion

- Unprecedented high burden of HIV in young women
- Greater understanding of WHY young women are at high risk
- Strong empiric evidence on
  - Age disparate partnerships
  - Genital inflammation
  - *Prevotella bivia*
  - Absence of *Lactobacilli*
  - *Gardnerella vaginalis* compromises the activity of tenofovir
- Implications for the design and testing of biomedical interventions
- Focussed effort to include young women who would benefit the most
- Ethical and programmatic challenges

**But need to pursue**

# Acknowledgements

## Investigators in the CAPRISA 251 (HIPSS) team

- Epicentre AIDS Risk Management
- Centre for the Programme of AIDS Research in South Africa (CAPRISA)
- Health Economics and HIV/AIDSs Research Division (HEARD)
- National Institute for Communicable Diseases, National Health Laboratory Service (NICD/ NHLS)

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- uMgungundlovu District office DOH
- HAST Unit, KZN Department of Health
- KZN Department of Health
- KRISP-UKZN
- University of Cape Town

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## CAPRISA

- Salim S Abdool Karim; Quarraisha Abdool Karim; Cheryl Baxter

