

much reinfection within partnerships contributes to the persistent spread of chlamydia in heterosexual populations.

**Methods** We derive an expression for the basic reproduction number of an SIS infection in a population with long term monogamous partnerships. The basic reproduction number contains an explicit term quantifying the contribution of re-infection within partnerships to the overall reproduction number. The derivation is then extended to include partner notification and treatment. Finally, the model is extended to include two types of partnerships with differing average duration.

**Results** For small recovery rates and low transmission probabilities reinfection plays a minor role in sustaining chlamydia transmission. However, there is an optimal combination of duration of infection and transmission probability for which reinfection contributes substantially to keeping chlamydia endemic in a population. We discuss the functional dependency of the basic reproduction number on these parameters. Using a more complex model numerical scenarios were simulated showing that partner notification prevents a large proportion of re-infections.

**Conclusions** The effect of screening depends, in part, on whether or not it succeeds in moving the basic reproduction number away from the transmission optimum via reinfection. This can be achieved either by choice of the screening interval or by rescreening those individuals who tested positive in a first screening test. There is an optimal time interval for retesting that minimises the basic reproduction number. The precise numerical value depends on partnership durations and transmission probabilities.

#### P1-S5.42 PREVALENCE OF TRANSITIONAL AND EMBEDDED CONCURRENCY WITHIN A PROSPECTIVE COHORT OF URBAN ADOLESCENT FEMALES

doi:10.1136/sextrans-2011-050108.220

<sup>1</sup>P Matson, <sup>1</sup>S E Chung, <sup>2</sup>J D Fortenberry, <sup>1</sup>J Ellen. <sup>1</sup>Johns Hopkins School of Medicine, Baltimore, USA; <sup>2</sup>Indiana University-Purdue University Indianapolis, Indianapolis, USA

**Background** Studies have found that concurrent sexual partnerships, as opposed to cumulative number of sex partners or serially monogamous partnerships play a pivotal role in rates of STI transmission. Different patterns of concurrency have different consequences to STI transmission as these patterns result in diverse network structures from within which to disseminate disease. Our objective was to determine the prevalence and types of concurrent partnerships within a prospective cohort of urban adolescent females.

**Methods** A cohort of adolescent females (N=63), aged 16–19 at baseline, were recruited from health clinics and community venues in Baltimore, MD and completed daily diaries about each current sex partner on a Palm Smartphone continuously for up to 18 months. As data collection is ongoing, relationships that were at least 8 weeks long were selected for this analysis. Partner names were confirmed by participants at regular intervals to ensure the same partner was followed over time. Sex partners were considered concurrent if they were named by the participant in the same week. Concurrent partners were categorised as either embedded or transitional relative to each relationship. A side partner was categorised as embedded if the week(s) that partner was named were completely contained within the weeks of the relationship. A side partner was categorised as transitional if weeks that the partner was named either preceded or persisted beyond the relationship or if the partner was only named in a single week within the first or last 4 weeks of the relationship.

**Results** Fifty-five participants had a relationship of at least 8 weeks duration, of these 55 relationships 28 (51%) had a concurrent

partner within the relationship, with a mean (SD) of 2.7 (2.6) concurrent partners. Of the 28 relationships that contained concurrency, transitional concurrency was most prevalent (43%), followed by both transitional and embedded (39%), and embedded only (18%). Among relationships with concurrency, 25% had a gap of >4 weeks within the relationship, as a result partners initially classified as transitional subsequently became embedded. There was no difference in relationship duration for those with and without concurrency.

**Conclusions** Transitional concurrency may reflect adolescents' intentions of serial monogamy. Future data collection is needed to gain an improved understanding of adolescents' motivations for concurrent sexual partnerships.

### Epidemiology poster session 5: Transmission dynamic: Income/race disparities

#### P1-S5.43 DISPARITIES IN SEXUALLY TRANSMITTED DISEASES ACROSS RACE-INCOME "COUNTIES IN THE USA: A RACE-SPECIFIC MORBIDITY ANALYSIS"

doi:10.1136/sextrans-2011-050108.221

K Owusu-Edusei, H Chesson, J Leichter, C Kent, S Aral. Centers for Disease Control and Prevention, Atlanta, USA

**Background** We examined the association between race-specific incidence rates for three major bacterial STDs and race-specific household income at the county level for all counties in the 48 contiguous states in the USA.

**Methods** Race-specific county-level median household incomes were obtained from the 2000 decennial census. We used the overall US median household income (ie, \$41 994, in 2000 dollars) as the cut-point to categorise counties into six race-income county groups. "Race-income" county groups were defined by black household income and white household income at the county level (see Abstract P1-S5.43 table 1). County-level race-specific morbidity for chlamydia, gonorrhoea and primary and secondary (P&S) syphilis were obtained from the National Electronic Telecommunications System for Surveillance (NETSS) for 1999–2001. For each of the six county groups, we calculated race-specific STD rates (new cases per 100 000 residents) using the total number of cases and population size for all three years. In a supplemental analysis, we used a spatial regression technique to examine the association in more detail, controlling for county-level socio-demographic factors.

Abstract P1-S5.43 Table 1 Average race-specific STD rates (number per 100 000 residents), 1999–2001 for six "race-income" county groups

County group	Description of race-specific household income in county group	Chlamydia		Gonorrhoea		P & S syphilis	
		Black	White	Black	White	Black	White
1	Black and white incomes below national median	695.2	88.8	597.0	22.9	10.7	0.5
2	Black income below national median, white income above national median	691.1	67.5	625.3	23.5	14.0	0.7
3	Black income above national median, white income below national median	279.3	61.3	125.4	5.6	6.2	0.1
4	Black and white incomes above national median	220.6	29.2	176.1	6.3	4.2	0.3
5	No black households in county; white income below national median	—	55.3	—	3.0	—	0.2
6	No black households in county; white income above national median	—	55.1	—	3.8	—	0.0

**Results** STD rates for Blacks in each category were at least 4 times higher than for whites. For chlamydia, county group 1 had the