States and several countries in Europe (7), including Italy, and is responsible for several disease outbreaks. In Italy, this genotype represents 61% (70/115) of all serogroup C strains collected during 2012–2015. Two outbreaks caused by this strain were reported in Italy in 2007 (8) and in 2012 (9). The N. meningitidis factor H binding protein and heparin binding protein alleles were 1.13 and 20, respectively. The N. meningitidis adhesin A variant had an insertion sequence that disrupted this gene, as described for ET-15 meningococci (10). On the basis of results of cgMLST, the strain was determined to be related to strains responsible for an outbreak in Italy in 2015.

In summary, we report a case of meningococcal pericarditis caused by a strain of N. meningitidis. This strain belongs to hyperinvasive clonal complex cc11 and was identified as C:P1.5–1,10–8:F3–6:ST-11(cc11), an emerging strain in Italy and worldwide. Timely diagnosis and complete molecular characterization of this strain, which causes a rare form of invasive disease (4), is needed for appropriate management of patients with this disease.

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References
1. Finkelstein Y, Adler Y, Nussinovitch M, Varsano I. A new classification for pericarditis associated with meningococcal infection. Eur J Pediatr. 1997;156:585–8. http://dx.doi.org/10.1007/s004310050669
2. Herrick WW. Meningococcic pericarditis, with report of 12 cases: medical service of the base hospital, Camp Jackson, S.C. Med Clin North Am. 1918;2:411–26.
3. Orgain ES, Postom MA. Pericarditis with effusion due to meningococcus. Am Heart J. 1939;18:368. http://dx.doi.org/10.1016/S0002-8703(39)90637-X
4. Blaser MJ, Reingold AI, Alsever RN, Hightower A. Primary meningococcal pericarditis: a disease of adults associated with serogroup C Neisseria meningitidis. Rev Infect Dis. 1984;6:625–32. http://dx.doi.org/10.1093/clinids/6.5.625
5. Nkosil Thakrar A, Kumar K, Ahmadie F, Fang T, Lytwyn M, et al. Meningococcal serotype Y myopericarditis. Diagn Microbiol Infect Dis. 2009;63:223–7. http://dx.doi.org/10.1016/j.diagmicrobio.2008.09.015
6. Zhu H, Wang Q, Wen L, Xu J, Shao Z, Chen M, et al. Development of a multiplex PCR assay for detection and genotyping of Neisseria meningitidis. J Clin Microbiol. 2012;50:46–51. http://dx.doi.org/10.1128/JCM.00918-11
7. Aubert L, Taha MK, Boo N, Le Strat Y, Deghmane AE, Sanna A, et al. Serogroup C invasive meningococcal disease among men who have sex with men and in gay-oriented social venues in the Paris region: July 2013 to December 2014. Euro Surveill. 2015;20:pii:21016.. http://dx.doi.org/10.2807/1560-7917.ES2015.20.3.21016
8. Fazio C, Neri A, Tonino S, Carannante A, Caporal MG, Salmaso S, et al. Characterization of Neisseria meningitidis C strains causing two clusters in the north of Italy in 2007 and 2008. Euro Surveill. 2009;14:pii:19179.
9. Stefanelli P, Fazio C, Neri A, Isola I, Sani S, Marelli P, et al. Cluster of invasive Neisseria meningitidis infections on a cruise ship, Italy, October 2012. Euro Surveill. 2012;17:pii:20336.
10. Elias J, Vogel U. IS1301 fingerprint analysis of Neisseria meningitidis strains belonging to the ET-15 clone. J Clin Microbiol. 2007;45:159–67. http://dx.doi.org/10.1128/JCM.01322-06

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Ecologic Study of Meningococcal B Vaccine and Neisseria gonorrhoeae Infection, Norway

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To the Editor: Gonorrhea is a sexually transmitted disease that can cause pelvic inflammatory disease, ectopic pregnancy, and salpingitis in women and infertility in men and women. Rates vary; incidence is 12.5 cases/100,000 population in Europe (1) and ≈6,000 cases/100,000 population in parts of sub-Saharan Africa (2). Recurrent infection is common, antimicrobial drug resistance is growing, and no licensed vaccine is available to protect against gonorrhea infection. Components of some meningococcal B (MenB) vaccines could provide protection against the causative bacterium, Neisseria gonorrhoeae (M. Pizza, pers. comm.), because the meningococcus bacterium is of the same Neisseria genus and the 2 bacteria share key protein antigens, such as the outer membrane vesicle (OMV). Ecologic evidence from Cuba supports a decline in gonococcus

1Now a GlaxoSmithKline company.
infection after a nationwide OMV vaccine campaign in the 1980s (3). In Norway, a trial of another OMV MenB vaccine (MenBvac, Norwegian Institute of Public Health, Oslo, Norway) was conducted among teenagers during 1988–1992. We retrospectively examined associations between MenB vaccine coverage during 1988–1992 and national gonorrhea rates for persons ≥16 years of age during 1993–2008 in Norway.

Age, year of birth, and year of diagnosis for anonymized, laboratory-confirmed gonorrhea patients were available from the Norwegian Surveillance System for Communicable Diseases from 1993 onwards. We collected aggregate vaccination coverage by MenB vaccine from the electronic trial register for 1988–1992. Only children enrolled in secondary schools and ≈13–15 years of age were offered the vaccine. This group accounted for 63% of the 148,589 children resident in Norway during the trial period and born during 1973–1976. Using annual population estimates, we derived gonorrhea notification rates and compared 3 cohorts: the vaccinated cohort (VC) born during 1973–1976; the unvaccinated cohort born during 1965–1972 (pre-VC); and the unvaccinated cohort born after 1976 (post-VC). Using Poisson log-linear regression adjusted for year of diagnosis, we calculated incidence rate ratios (IRRs; number of new diagnoses of gonorrhea per 100,000 population) for VC and post-VC and compared them with IRRs for pre-VC (the reference group).

During 1993–2008, a total of 2,601 cases of gonorrhea were reported. To avoid case ascertainment bias, cases reported from 2009 onwards were excluded because >50% were diagnosed by PCR. In men, notification rates fell from 27.3/100,000 population in 1993 to 12.3 in 1995 but increased to 22.9 in 2008 (IRR 1.02, 95% CI 1.00–1.03, p = 0.001). The recent increase is largely attributable to transmission between men who have sex with men, a group accounting for 12% of cases among men in 1993 but 40% in 2008. Among women, incidence rates dropped from 20.7/100,000 population in 1993 to 3.1 in 1999 and have remained stable since 2008, when the rate was 4.1.

We hypothesized that an immunogenic effect of MenB vaccination on gonorrhea notification rates might be most expected among persons in their early 20s in the mid-1990s, a group at risk for gonorrhea infection during years closest to the vaccination campaign. A small but significant reduction occurred in crude incidence rates for women 20–24 years of age in the VC (IRR 0.58, 95% CI 0.42–0.8); however, this reduction was not significant after incidence rates were adjusted for year of diagnosis (IRR 0.72, 95% CI 0.51–1.02). No change in incidence occurred for other age groups or birth cohorts among women (Figure). Reduced

Figure. Incidence rate ratios (IRRs) of cases of gonorrhea by age group and birth cohort, adjusted for year of diagnosis in A) women and B) men, Norway. IRRs were calculated by using Poisson log-linear regression and adjusted for year of diagnosis. IRRs indicate incidence of gonorrhea in the vaccinated cohort (VC), (i.e., persons born during 1973–1976, 63% of whom were vaccinated), represented by triangles, and in the post-VC (i.e., those born after 1976), represented by circles, relative to the reference birth cohort, the pre-VC (i.e., persons born during 1965–1972), represented by the horizontal dotted line. Also shown are 95% CIs. Differences in the IRR were considered statistically significant if the CI did not include 1. A) Among women, the adjusted IRR in the VC 20–24 years was of borderline significance: 0.72, 95% CI 0.51–1.02. No effect was seen in other age groups or birth cohorts. B) Among men, a reduced adjusted incidence was evident for those 20–24 years of age in the VC (IRR 0.68, 95% CI 0.51–0.93) and persisted among post-VC who were 20–24 years of age (IRR 0.51, 95% CI 0.33–0.78). IRR increased for men 25–29 years of age in the post-VC (IRR 1.6, 95% CI 1.06–2.43).
incidence also occurred for men 20–24 years of age in the VC (adjusted IRR 0.68, 95% CI 0.51–0.93) and persisted in the post-VC (adjusted IRR 0.51, 95% CI 0.33–0.78); the IRR increased for men 25–29 years of age in the post-VC (adjusted IRR 1.6, 95% CI 1.06–2.43). (Figure).

Overall, rates of gonorrhea dropped among men and women after the vaccination campaign; however, rates had already been in decline since the mid-1970s (4). A limited age-specific vaccine effect occurred among men and women 20–24 years of age. No vaccine effect was found among women in other age groups or birth cohorts. Among men 20–24 years of age, the persistent decline occurring among the post-VC could be explained by a herd effect, but rates subsequently increased for men 25–29 years of age. Different effects for men and women may be explained in part by changing transmission patterns and sexual behavior occurring among men who have sex with men and by inadequate adjustment of the underlying gonorrhea trend in men; however, differences by sex are difficult to interpret.

Among study limitations, our ecologic study design could not distinguish between long-term trends or behavioral factors and vaccine effects. For example, in the early 1990s, condom use increased in Norway, especially among persons in their early 20s (5), possibly in response to the evolving HIV epidemic (6,7). In addition, moderate population vaccine coverage in the VC (i.e., 63%) and a time lag between the vaccination program (1988–1992) and the start of surveillance (1993) may have diluted tangible vaccine effects. For further examination of cross-immunity of MenB vaccines with gonococci, vaccine effectiveness studies in regions where OMV vaccine was used (e.g. New Zealand) and evaluation of new protein-based vaccines are warranted.

References
1. European Centre for Disease Prevention and Control. Annual epidemiological report. Reporting on 2011 surveillance data and 2012 epidemic intelligence data. Sexually transmitted infections, including HIV and blood-borne viruses. Stockholm: The Centre; 2013.
2. World Health Organization. Global incidence and prevalence of selected curable sexually transmitted infections 2008. Geneva: The Organization; 2012.
3. Perea O, del Campo J, Cuello M, González E, Nuñez N, Cabrera O, et al. Mucosal approaches in Neisseria vaccination. Vaccimonitor. 2009;18:55-57. Epub ahead of print.
4. The Norwegian Institute of Public Health [Smittevernboeka]. Gonorrhoea—guide for healthcare personnel [in Norwegian]. 2010 Feb 25; modified 2015 [cited 2015 Feb 6]. http://www.fhi.no/artikler/?id=82735
5. Træen B, Stigum H, Magnus P. Report from sexual surveys in 1987, 1992, 1997 and 2002 [in Norwegian]. Oslo: Norwegian Institute of Public Health; 2003.
6. Piot P, Islam MQ. Sexually transmitted diseases in the 1990s. Global epidemiology and challenges for control. Sex Transm Dis. 1994;21(Suppl):S7–13.
7. Schwarze SK, Kellogg TA, Kohn RP, Katz MH, Lemp GF, Bolan GA. Temporal trends in human immunodeficiency virus seroprevalence and sexual behavior at the San Francisco municipal sexually transmitted disease clinic, 1989–1992. Am J Epidemiol. 1995;142:314–22.

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Neisseria gonorrhoeae Resistant to Ceftriaxone and Cefixime, Argentina

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To the Editor: Antimicrobial resistance in Neisseria gonorrhoeae is increasing globally. In recent years, gonococcal strains with resistance to the extended-spectrum cephalosporin (ESC) ceftriaxone have been reported from many countries (1). In South America, 7 ceftriaxone-resistant strains (MICs >0.25 μg/mL) were reported from Brazil in 2007; however, these isolates have not been characterized (2). Emergence of cephalosporin-resistant gonorrhea would substantially limit treatment options and represent a major public health concern. We report an N. gonorrhoeae isolate in Argentina that was resistant to ceftriaxone and cefixime.

In September 2014, a 19-year-old heterosexual man with no underlying disease was admitted to a hospital emergency department in Rio Negro, Argentina. Physical examination showed a purulent urethral discharge. He reported having had unprotected insertive vaginal sex with multiple partners in the past few months. The patient denied recent travel outside Argentina. He had a history of gonococcal urethritis (March 2014), which was treated with a single dose (500 mg) of ceftriaxone; a gonococcal isolate obtained at that time was not sent to a reference laboratory as part of the Argentinian Gonococcal Antimicrobial Susceptibility Surveillance Program for determination of the antimicrobial susceptibility profile.

At the time of admission, we obtained and cultured a urethral swab specimen and identified an isolate as N. gonorrhoeae by using conventional methods (3). The isolate was sent to a reference laboratory, and its identity was confirmed by using the Phadebact GC Monoclonal Test (MKL