



FIG. 3. Transformation of the full-length *penA* allele (*penA*_{H041}) from the high-level ceftriaxone-resistant *Neisseria gonorrhoeae* strain H041 (Donor) into *N. gonorrhoeae* strains (Recipients) with different ceftriaxone MICs and genetic resistance determinants affecting the susceptibility to ceftriaxone. The ceftriaxone MICs using the Etest method (shown as mean results of three repeated experiments) of the donor strain, recipient strains (R), and transformants (T) and the MIC ratio (T/R) are given. The breakpoint for ceftriaxone resistance is according to reference 8.

ceftriaxone resistance also and, although the biological fitness of ceftriaxone resistance in *N. gonorrhoeae* remains unknown, the gonococcus may soon become a true superbug that initiates a future era of untreatable gonorrhoea. To at least slow the spread of ESC (cefexime and ceftriaxone) resistance, a reduction in global gonorrhoea burden by enhanced disease prevention and control activities is crucial. As well, the implementation of much wider strategies for general AMR control, better understanding of the mechanisms and global monitoring of the emergence and spread of AMR, and global and national public health response plans (including sustainable clinical, microbiological, and epidemiological components) are needed. Any such plan alone will most probably not be able to prevent the emergence, establishment, and spread of ceftriaxone resistance; nevertheless, these plans will be valuable to delay and limit a global spread of ESC resistance (cefexime and ceftriaxone). Ultimately, a major focus important for public health globally is the timely development of effective new drugs (for single or combined use) for the treatment of gonorrhoea.

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