

Overcoming and containing bacterial resistance: appropriate antibiotic use in community-acquired RTIs

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The global spread of antibiotic resistance is driving the need for more thoughtful antibiotic prescribing. This paper reviews the principles of appropriate antibiotic therapy for community-acquired respiratory tract infections. Appropriate therapy options include: prescribing antibiotics only when they are beneficial to the patient; using agents that target the likely pathogens, taking into account local resistance patterns and risk factors for infection by resistant pathogens while not affecting the bowel flora and other nonrespiratory organisms; and using a dosing schedule and treatment duration that optimizes efficacy, tolerability and adherence to treatment.

The spread of antibiotic resistance among bacteria responsible for common infections has prompted calls for improved use of antibiotics in the USA and elsewhere [1,2]. Community-acquired respiratory tract infections (RTIs), such as acute bacterial sinusitis (ABS), acute exacerbations of chronic bronchitis (AECB), and community-acquired pneumonia (CAP), account for the majority of antibiotic prescriptions in outpatients [3].

The widespread and unnecessary use of antibiotics for viral RTIs contributes substantially to the selective pressure driving resistance [4]. Educational campaigns have had some success in reducing total antibiotic use in this setting [3]. However, in the absence of diagnostic tests the differentiation of viral and bacterial infections based on clinical signs and symptoms remains a major challenge [5]. A number of guidelines have sought to encourage better discrimination of viral from bacterial infections [6–8]. These generally recommend that acute RTIs among patients with no comorbidities or other predisposing factors are unlikely to be bacterial in origin for infections of short duration. Antibiotics are of no benefit for RTIs caused by viruses. They should thus be reserved for infections of more than 7 days' duration, which are more likely to be bacterial in origin, and for patients with specific risk factors for bacterial infection [5].

The spread of antibiotic resistance among bacteria responsible for common infections has prompted calls for improved use of antibiotics in the USA and elsewhere [1,2]. Frequently, the emphasis has been on reducing the amount of antibiotics prescribed, but there is a potential drawback to this approach, which is highlighted by the excess mortality that may occur [9]. This emphasizes the need to accurately identify patients with bacterial rather than viral RTIs.

Attention must now shift towards improving antibiotic prescribing for bacterial RTIs, with regard to the quality rather than merely the quantity of antibiotics used [10–12]. According to the CDC, appropriate antibiotic use aims to maximize therapeutic impact while minimizing toxicity and the development of resistance [13]. This involves the selection of the most appropriate antibiotic, dose and treatment duration, in addition to promoting adherence to therapy (e.g., by educating patients and using antibiotics with more convenient and simpler dosing regimens). In outlining these aspects of therapy, this paper reviews the concepts behind appropriate prescribing for community-acquired RTIs of bacterial origin and how these concepts can overcome and help to contain antibiotic resistance in this setting. In practice, different countries recommend different antibiotics for similar situations in their guidelines [14–16]. These guidelines differ in their use of evidence and expert consensus and must also take account of the local patterns of antibiotic resistance.

Spectrum of activity

A causative organism is rarely isolated before antibiotic treatment for community-acquired RTIs is begun. Hence, antibiotics for community-acquired RTIs are usually chosen empirically according to the likely causative pathogens and their antibiotic susceptibilities. The organisms most commonly responsible for community-acquired RTIs of bacterial origin are *Streptococcus pneumoniae* (the 'pneumococcus'), *Haemophilus influenzae* and *Moraxella catarrhalis* (Table 1). Atypical and intracellular organisms, such as *Chlamydia pneumoniae* (previously *Chlamydia pneumoniae*), *Legionella pneumophila* and *Mycoplasma*

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Table 1. Bacterial pathogens responsible for the most common community-acquired respiratory tract infections in the USA.

Pathogen	CAP (%)	AECB (%)	ABS (%)
Typical pathogens			
<i>Streptococcus pneumoniae</i>	20–60	15–25	34
<i>Haemophilus influenzae</i>	3–10	30–59	35
<i>Moraxella catarrhalis</i>	2	3–22	0–8
Atypical/intracellular pathogens			
<i>Legionella</i> spp.	2–8	NA	NA
<i>Chlamydomphila pneumoniae</i>	4–6	NA	NA
<i>Mycoplasma pneumoniae</i>	1–6	NA	NA

ABS: Acute bacterial sinusitis; AECB: Acute exacerbations of chronic bronchitis; CAP: Community-acquired pneumonia; NA: Not available.

Adapted from [44].

pneumoniae, are also potential causes of CAP and, according to recommendations by the Infectious Diseases Society of America, should be covered by empirical therapy [17]. These atypical organisms are also implicated in a minority (<10%) of AECB cases [18].

Antibiotics for the empirical treatment of community-acquired RTIs should have a targeted spectrum of activity covering implicated pathogens, without unnecessary broad-spectrum effects on bowel flora and other organisms. If a causative organism is identified, then it is possible to choose antibiotics with even greater specificity. Agents available for use in the treatment of RTIs and currently recommended in treatment guidelines [14–16] include β -lactams (e.g., amoxicillin, cefuroxime), β -lactam/ β -lactamase inhibitor combinations (e.g., amoxicillin–clavulanate), macrolides/azalides (e.g., erythromycin, azithromycin), doxycycline, trimethoprim–sulfamethoxazole, respiratory fluoroquinolones (e.g., levofloxacin, moxifloxacin) and ketolides (e.g., telithromycin). Typically, β -lactams, macrolides/azalides and doxycycline are recommended as first-line agents with newer drugs, such as ketolides and fluoroquinolones, reserved for patients with more severe disease and those who have experienced treatment failure with first-line agents. The new agent, linezolid, has a spectrum of activity limited to Gram-positive organisms and is not recommended for empirical treatment of ambulatory patients [19].

Resistance

Resistance patterns

S. pneumoniae – a primary bacterial cause of ABS [20], AECB [18] and CAP [17] – is commonly resistant *in vitro* to β -lactams, macrolides, trimethoprim–sulfamethoxazole and tetracyclines.

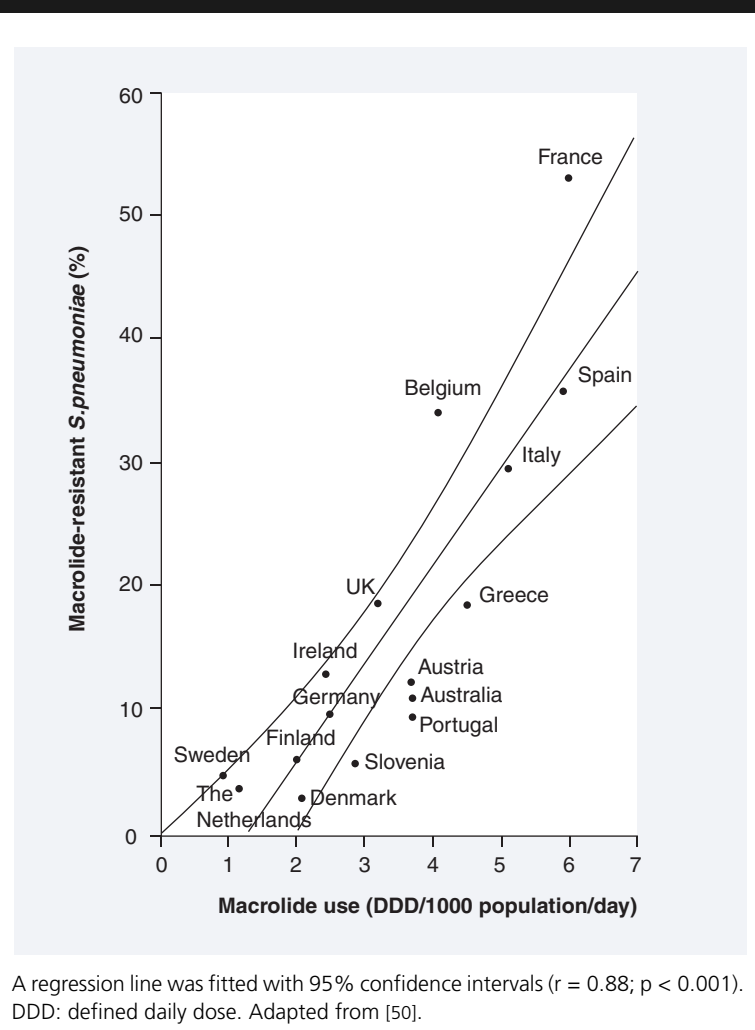
A number of studies (including the Alexander project [21], the SENTRY [22] and TRUST [23] antimicrobial surveillance programs) demonstrate that antibiotic resistance rates have tended to increase over time. The most recent data from the ongoing Prospective Resistant Organism Tracking and Epidemiology for the Ketolide Telithromycin (PROTEKT) US study show that, among 31,001 *S. pneumoniae* isolates collected between 2000 and 2003, 29.4% were resistant to erythromycin and 22.5% to penicillin [24]. The prevalence of macrolide resistance was also found to differ between US states, with the highest rate (48.2%) reported in Louisiana (LA, USA) and the lowest rate (15.2%) reported in Vermont (VT, Canada) in the period 2001–2002 [25]. This suggests that a switch away from macrolides as first-line agents may be justified in states with high rates of macrolide resistance and physicians should be aware of local antibiotic resistance surveillance data.

A particular concern is the rise in prevalence of multiple antibacterial drug-resistant *S. pneumoniae* strains. Approximately 31% of *S. pneumoniae* isolates collected in the PROTEKT US study between 2000 and 2003 were resistant to two or more antibiotic classes [24]. Macrolide resistance in the USA is most commonly associated with the *mef(A)* genotype, but isolates with the dual *erm(B)* + *mef(A)* genotype, which confers high-level macrolide resistance and multidrug resistance, are becoming more common [25].

The ‘respiratory’ fluoroquinolones and telithromycin remain reliably active against *S. pneumoniae*. Fluoroquinolone resistance in *S. pneumoniae* remains rare, with a prevalence of approximately 1% [24,26]. However, the trend in fluoroquinolone resistance in the USA has been towards increasing resistance over time [26] and this trend is expected to continue [27]. Infection with a fluoroquinolone-resistant isolate is more likely in nosocomial infections and infections acquired in residential care homes compared with isolates active in the general community [28]. *S. pneumoniae* isolates with low-level resistance to telithromycin have been reported [29] but US resistance surveillance data suggest that *in vitro* telithromycin resistance among *S. pneumoniae* remains very low (<0.8%) [30].

The other major pathogens implicated in community-acquired RTIs, *H. influenzae* and *M. catarrhalis*, commonly produce β -lactamase enzymes that inactivate many β -lactam agents, including penicillin, amoxicillin and some early

Figure 1. Relationship between macrolide use in the outpatient setting and prevalence of macrolide-resistant *Streptococcus pneumoniae* in 16 industrialized countries.



cephalosporins. In the USA, 28% of *H. influenzae* isolates and over 90% of *M. catarrhalis* isolates produce β -lactamases [30,31]. Agents with undiminished activity against β -lactamase-positive strains include amoxicillin–clavulanate, azithromycin, fluoroquinolones and telithromycin [30,31].

Impact of resistance

In the USA, macrolides/azalides are favored as the first-line agents in treating RTIs. Macrolides are effective against typical RTI pathogens and, in contrast to β -lactams, are also effective against atypical/intracellular pathogens [32] and β -lactamase-producing Gram-negative organisms [33]. The primary macrolide resistance mechanism observed in the USA (*mef*[A]) confers intermediate *in vitro* resistance compared with Europe where the *erm*(B) mechanism confers higher-

level resistance [25] and where the use of β -lactams as first-line agents is more common. However, the spread of higher macrolide resistance clones in the USA may require a re-evaluation [14] or limitation [34] of the use of macrolides.

Bacteriologic eradication is the main determinant of clinical outcome in the treatment of community-acquired RTIs [35]. Surprisingly, the effect on clinical outcomes of infection by *in vitro* antibiotic-resistant strains is poorly characterized. Small studies and case reports support a link between macrolide resistance and treatment failure in CAP and bacteremia [34–37] and have led some experts to suggest that the impact of macrolide-resistant *S. pneumoniae* may be underestimated [38]. The concern about increasing prevalence of macrolide resistance in *S. pneumoniae* may explain the shift in prescribing in the USA towards fluoroquinolones in the treatment of CAP [39]. This trend is worrying, since the guidelines reserving fluoroquinolone use for higher-risk patients do not appear to be being followed [39] and there are documented case reports of fluoroquinolone treatment failures among patients with CAP caused by fluoroquinolone-resistant *S. pneumoniae* [40–42].

Antibiotic resistance has the potential to increase the healthcare and socioeconomic costs associated with infections [43]. In hospitalized patients, antibiotic-resistant infections have been associated with increased healthcare utilization in terms of duration of hospitalization, antibiotic drug costs and nursing charges [44,45]. Further research is needed in order to better define and quantify the burden of resistance in the outpatient setting [46].

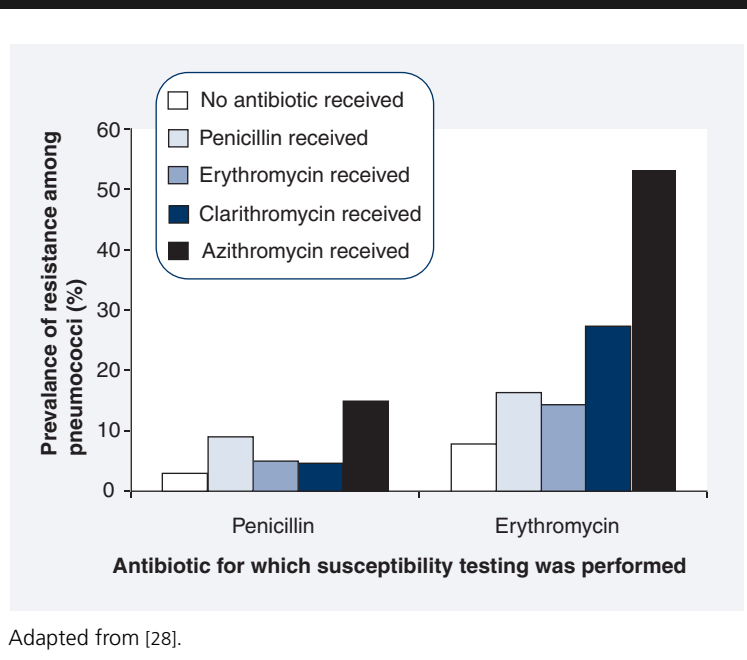
Selection & induction of resistance

Antibiotic consumption in the community is linked with bacterial-resistance patterns through complex relationships [47], thus the propensity for different antibiotic regimens to select for or induce resistance is a factor to be considered when choosing therapy.

Resistance at the population level

Ecological studies have demonstrated a correlation between β -lactam and macrolide consumption at the population level with high levels of pneumococcal resistance to these agents [48–52]. For example, Albrich and colleagues correlated macrolide use with resistance levels in 16 European countries (Figure 1) [50]. Data suggest that it is macrolides and not β -lactams that are

Figure 2. Association between the use of antibiotics during the 3-month period before invasive pneumococcal infection and the susceptibility of the infecting isolate to penicillin or erythromycin.



the main agents promoting both penicillin and macrolide resistance [49]. Recent studies have implicated the increased use of once-daily macrolides (e.g., azithromycin) as the principal driver of macrolide resistance [49,51,52]. The prolonged half-life of azithromycin is suggested to result in subinhibitory tissue concentrations that favor the selection of resistant strains [52].

Increased use of levofloxacin in the USA has also been correlated with so far modest increases in fluoroquinolone resistance in *S. pneumoniae* [53]. However, the rise of fluoroquinolone use in the community has been associated with increased levels of resistance in Gram-negative organisms, such as *Pseudomonas aeruginosa* and Enterobacteriaceae, which can cause serious infections in hospitalized patients [54,55]. In contrast, telithromycin appears to have a lower ecological impact on gastrointestinal flora than other agents [56].

According to the CDC, the widespread use of azithromycin, clarithromycin and fluoroquinolones in the USA warrants concern in light of resistance trends that threaten the utility of these agents in hospitalized patients [11].

Predicting resistance in individuals

At the level of the individual patient, previous antibiotic use is predictive of infection by a resistant organism, as demonstrated by a prospective

Canadian study involving 3339 patients with invasive pneumococcal infections [28]. Upon multivariate analysis, use of clarithromycin or azithromycin in the previous 3 months was significantly associated with infection by a macrolide-resistant strain, with odds ratios (ORs) of 3.93 (95% confidence interval [CI]: 2.16–7.16; $p < 0.001$) and 9.93 (95% CI: 4.85–20.3; $p < 0.001$) for these two agents, respectively (Figure 2). Previous azithromycin use was also significantly associated with resistance to penicillin and trimethoprim–sulfamethoxazole, while fluoroquinolone use predicted fluoroquinolone resistance with an OR = 12 ($p < 0.001$). Fluoroquinolone use within the previous month is also predictive for the emergence of resistant Gram-negative bacilli in the gastrointestinal flora [57].

Factors affecting the increase of resistance

Bactericidal agents (e.g., penicillins, fluoroquinolones and ketolides) may be less likely to select resistant strains than bacteriostatic agents (e.g., macrolides) owing to more rapid eradication of organisms by the former [58]. Resistance selection may be more likely when bacteria are exposed to prolonged, subtherapeutic antibiotic concentrations. This not only explains the aforementioned resistance selection by azithromycin, but also why β -lactams administered at low daily doses for long treatment durations (>5 days) are associated with an increased risk of nasopharyngeal carriage of penicillin-resistant *S. pneumoniae* as compared with high-dose, short-course regimens [59,60]. This highlights the need to prescribe antibiotics at appropriate doses based on pharmacokinetic/pharmacodynamic principles [14]. High-dose amoxicillin courses have thus been developed to combat the increase in penicillin resistance among *S. pneumoniae* strains [61].

Certain antibiotics may have inherent characteristics that reduce their propensity to induce or select for resistance. For example, ketolides, unlike macrolides, do not induce the macrolide–lincosamide–streptogramin_B resistance phenotype coded for by the *erm(B)* genotype [57,62] and the activity of telithromycin has not changed between 1999 and 2003, including in those countries where this antibiotic is in clinical use [63]. Similarly, respiratory fluoroquinolone resistance in *S. pneumoniae* has remained at approximately 1% in the USA [24,26]. This is because resistance selection mainly occurs via a stepwise mechanism, with mutations required in both the topoisomerase IV and gyrase genes. Relatively often strains have mutations in one gene but not both [64].

Highlights

- The spread of antibiotic resistance requires improved use of antibiotics in clinical practice.
- Inappropriate use of antibiotics in viral infections helps drive resistance, but differentiation of viral and bacterial respiratory tract illness remains a major challenge.
- Antibiotic-resistant infections are associated with increased healthcare utilization and increased costs.
- Empiric antibiotic treatment of community-acquired respiratory tract infections (RTIs) must be effective against all of the likely causes of the infection – *Streptococcus pneumoniae*, *Haemophilus influenzae*, *Moraxella catarrhalis*, and also 'atypical' organisms, such as *Chlamydophila pneumoniae*, *Legionella pneumophila* and *Mycoplasma pneumoniae*.
- Antibiotic resistance surveillance studies are providing vital local resistance data that can inform local antibiotic prescribing.
- Previous antibiotic use by a patient is predictive of infection by a resistant organism.
- Drug treatment regimens and course duration are important considerations in determining patient adherence to antibacterial treatment. Poor adherence to treatment can expose bacteria to subtherapeutic drug levels, leading to treatment failure and resistance selection.

Adherence & tolerability

According to a recent meta-analysis, mean adherence to antibiotic dosing regimens may be as low as 62% [65]. Poor adherence to antibiotic regimens can expose bacteria to subtherapeutic doses, and hence may lead to treatment failure and resistance selection. Patients are more likely to comply with dosing regimens that are short and conveniently administered, preferably once daily [66]. Short-course therapy is of direct benefit in reducing the selective pressure for resistance [60,61] and a growing body of evidence supports its use in community-acquired RTIs [67]. However, once-daily administration must be coupled with an appropriate pharmacokinetic/pharmacodynamic profile in order to avoid subtherapeutic concentrations and the associated risk of resistance selection.

Tolerability also has an important influence on compliance, as adverse events may prompt patients to discontinue therapy [66]. Thus,

physicians must consider the relative tolerability of different antibiotics in the relevant patient population when selecting therapy for community-acquired RTIs. The use of shorter treatment courses may also contribute to reducing the risk of adverse events.

Expert commentary

The association between antibiotic use and widespread bacterial resistance in pathogens responsible for community-acquired RTIs underscores the importance of selecting appropriate antibiotic therapy that maximizes clinical outcomes and cost-effectiveness, while limiting the selective pressure for resistance.

Agents with appropriate antibacterial spectra, documented efficacy, good tolerability, a low potential for resistance induction, and a convenient dosing regimen will best meet this need.

Outlook

Ongoing antibacterial resistance surveillance studies suggest that, despite an increasing awareness of the problems of antibiotic resistance, β -lactam and macrolide nonsusceptibility will remain significant problems for the treatment of community-acquired RTIs in the foreseeable future. At a time when there is increasing pressure on healthcare provision services to contain costs, the expense of increased morbidity or mortality resulting from treatment failure with older, established antibiotics will have to be balanced against the costs of newer, more effective antibiotic drugs. Some of the information required for these decisions will be the increased provision of accurate, localized antibacterial surveillance data. The introduction of new antimicrobials will have an impact, but newer agents are likely to be in the same classes as existing drugs, underlining the need for appropriate prescribing now.

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Highlights

- Many strategies have been used to reduce immunogenicity issues with monoclonal antibodies (mAbs), including chimerization, humanization and human mAbs. The development of antibody fragments, such as antigen-binding fragments (Fab's), is a further advance.
- Fab's are associated with several benefits compared with whole antibodies including amenability to microbial expression and lack of potential toxicities mediated by the Fc portion of the antibody.
- Certolizumab pegol is a PEGylated, humanized Fab' fragment of an anti-tumor necrosis factor (TNF)- α mAb.
- PEGylation of the Fab' fragment is compatible with subcutaneous administration of certolizumab pegol, with potential advantages in terms of convenience and safety.
- The bioavailability after subcutaneous administration is 80–100% and the half-life is approximately 2 weeks.
- Certolizumab pegol has a high affinity and potency for human TNF- α . Preclinical and clinical data have shown the efficacy and tolerability of certolizumab pegol in Crohn's disease and rheumatoid arthritis.

therapy ('responders') were randomized at week 6 to either certolizumab pegol or placebo. At Week 26, the rates of clinical response (decrease in baseline Crohn's Disease Activity Index [CDAI] score ≥ 100 points) and remission (CDAI score ≤ 150 points) irrespective of C-reactive protein levels were significantly higher following treatment with certolizumab pegol compared with placebo. Certolizumab pegol was generally well tolerated, with mild-to-moderate headache being the most commonly reported adverse event [56].

Positive results for a second Phase III trial (PRECiSE 1) have also been presented recently [57]. The primary outcomes were met and the adverse-event profile was in line with that observed in PRECiSE 2.

Collectively, these studies indicate that the Fab' fragment certolizumab pegol has the potential to be clinically effective, safe and well tolerated for the treatment of CD and RA.

Comparison of certolizumab pegol with other anti-TNF agents

The characteristics of certolizumab pegol, etanercept, adalimumab and infliximab are summarized in Table 1.

Expert commentary

Since the inception of targeted antibody technologies in the mid-1960s, immunotherapy has undergone a number of developments in order to improve disease treatment and reduce

immunogenicity. Chimeric and humanized mAbs have become important therapeutic and diagnostic tools for a variety of diseases, including CD; however, such antibodies are associated with the production of HACA and HAMA. Antibody Fab' fragments represent an advance in the field of immunotherapy owing to their small size, flexibility and amenability to rapid production on a large scale.

Certolizumab pegol is a PEGylated humanized Fab' fragment of an anti-TNF- α mAb. PEGylation ensures a half-life for certolizumab pegol comparable to those of other anti-TNF full IgGs and produces a compound compatible with subcutaneous administration, while humanization of the Fab' fragment may reduce the potential to cause an immune reaction. Unlike full-length IgG1 antibodies, Fab' fragments do not mediate ADCC and CDC owing to the absence of an Fc – this may have positive consequences from a safety perspective. Furthermore, Fab' fragments can also be produced via microbial fermentation – assisting a rapid and reproducible process and a reliable supply.

Certolizumab pegol exhibits high and specific affinity for TNF- α and high *in vitro* potency for TNF- α neutralization. It has also shown efficacy in an animal model of RA, as well as promising efficacy and tolerability results in Phase II and III trials. In comparative *in vitro* studies, the affinity and potency of certolizumab pegol were higher than that observed for the other anti-TNF agents infliximab and adalimumab.

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