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Current stage in inflammatory bowel disease: What is next?

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important economic impact on the healthcare system. Advances in recent years in pharmacogenetics and clinical pharmacology have allowed for the development of treatment strategies adjusted to the patient profile. Concurrently, new drugs aimed at inflammatory targets have been developed that may expand future treatment options. This review examines advances in the optimization of existing drug treatments and the development of novel treatment options for IBD.

Key words: Inflammatory bowel disease; Future directions; Pharmacogenetic; Pharmacokinetics; New drugs

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Core tip: The incidence and prevalence of inflammatory bowel disease (IBD) has been increasing worldwide. In recent years, the treatment objectives, the monitoring of IBD, and the drug treatments for controlling the disorder have been evolving. This review summarizes recent developments in pharmacogenetics, clinical pharmacology, and the use of new drug molecules that may expand IBD treatment options in the future.

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Abstract

In recent years, the incidence of inflammatory bowel disease (IBD) has been on the rise, extending to countries where it was infrequent in the past. As a result, the gap between high and low incidence countries is decreasing. The disease, therefore, has an

INTRODUCTION

Inflammatory bowel disease (IBD), Crohn's disease (CD), and ulcerative colitis (UC) are important public health problems. According to recent studies, the annual incidence of UC varies between 19.2-24.3 cases per 100000 inhabitants in Europe and 6.3 cases

per 100000 inhabitants in Asia and the Middle East^[1]. For CD, the estimated incidence is 12.7-20.2 cases per 100000 inhabitants in Europe and the United States vs five cases per 100000 inhabitants in Asia and the Middle East. The incidence of IBD is currently growing in areas where the disease was previously infrequent. As a result, the gap between high- and low-incidence countries is closing^[2]. This rise runs parallel to technological development, improvements in living standards, and a greater interest in this disease among physicians^[3]. The underlying pathogenesis remains uncertain, although the most widely accepted theory revolves around changes in the host immune response in genetically susceptible individuals to the intestinal microbiota that is triggered by environmental stimuli. None of these alterations alone can cause the disease, and the interactions among these four factors in the pathogenesis are very complex. In recent decades there have been important advances regarding each of these factors. Progress in the field of genetics has resulted from the performance of genome-wide association studies (GWAS), although they only account for 20%-25% of the cases of IBD^[4]. Knowledge of epigenetic mechanisms could explain the influence of environmental factors and the microbiota upon IBD and the low correlation to concrete genes^[5,6]. These developments have opened the door to personalized medicine^[7].

Knowledge of the immunological mechanisms involved in the manifestation of IBD has led to the development of new biological drugs. The first major advance is represented by the anti-tumor necrosis factor (TNF)- α drugs, which have revolutionized the treatment of IBD, since they are able to induce and maintain mucosal healing of the disease^[8], a key factor for modifying the natural course of the disorder^[9,10]. Nevertheless, despite these advances, one-third of all patients with CD fail to respond to anti-TNF- α therapy (primary non-responders), and 10% do not tolerate or do not respond to any of the drugs used to treat CD^[11,12]. In the case of UC, the reported colectomy rate reaches up to 21% after an initial response to anti-TNF- α drugs^[13]. This has led to the search for new therapeutic targets and further optimization of existing treatment options. Clinical pharmacology allows us to determine therapeutic drug concentrations (thiopurine agents and anti-TNF- α drugs) and, if needed, to explain their loss of responsiveness and their adverse effects. In the coming years, personalized medicine, where treatments will be prescribed according to the risk factors in each individual patient and the probability of achieving response to a given drug substance, will be initiated. There have been developments in the way IBD is monitored, with the adoption of reliable and scanty aggressive techniques, such as noninvasive imaging tests, stool markers, breath tests, *etc.*^[14], which fall beyond the scope of this review. We provide below a description of the current advances in pharmacogenetics and possible new drug

substances.

PHARMACOGENETICS

Personalized medicine seeks to find the ideal drug for each individual patient at the appropriate dose and administered *via* the best possible route. This approach allows for increased effectiveness, with the least risk of side effects, and at the lowest possible cost. Physicians try to identify patients with more serious disease, with a view to introducing early and more effective treatment in order to prevent long-term complications, distinguishing them from those individuals with less severe disease and a more favorable prognosis in which aggressive treatment poses a higher risk of undesired effects. Patient response to drug treatment is dependent upon many factors, including the severity of the disease and genetic and environmental factors.

Pharmacogenetics is the study of the association between the different polymorphisms of a gene and the variability of response to treatment or its toxicity with a given drug. It has been estimated that polymorphisms can account for 20%-95% of the variability of a response to a drug^[15].

A number of drugs are currently available for the treatment of IBD: 5-aminosalicylates, corticosteroids, immunosuppressors (thiopurine drugs, calcineurinic agents, methotrexate), and biological agents (anti-TNF- α drugs).

Aminosalicylates

The aminosalicylates are among the main agents used to treat patients with UC, and their colon cancer chemoprophylactic effect allows them to be used in UC with pancolonial disease involvement. The metabolism of both sulfasalazine and mesalazine is mediated by the enzyme N-acetyltransferase (NAT). For almost six decades, the population has been divided into fast and slow acetylators. There are two NAT isoenzymes (NAT1 and NAT2), and different polymorphisms have been described in different ethnic groups^[16]. NAT1 metabolizes mesalazine, and it has no demonstrable associations with clinical effects. NAT2 metabolizes salazopyrin derived from sulfasalazine breakdown. In 1983, a link between NAT2 slow acetylators, who accumulate higher drug levels in blood, and an increased number of side effects was shown. Twenty-five years later, and thanks to our knowledge of single nucleotide polymorphisms (SNPs), it has been possible to confirm the association between NAT2 with a slow acetylator phenotype and dose-dependent side effects^[17]. There are fewer studies on 5-acetylsalicylic acid (5-ASA) than with immunosuppressors and biological drugs, since 5-ASA is only used to reduce side effects that are usually not serious. However, since more prolonged treatment with 5-ASA was proposed due to its chemoprotective effect against colon cancer, the pharmacogenetic studies have become more important.

Glucocorticoids

Glucocorticoids (GLCs) are used in moderate and severe flare-ups of IBD, and although they are very effective, 16%-20% of all patients are refractory to GLCs in the Caucasian population, and 28%-36% are corticoid-dependent^[18-20]. The GLCs exert their anti-inflammatory effect by inhibiting T cell activation and cytokine secretion, following binding of the drug to the intracellular glucocorticoid receptors (GR- α), which modify their structural conformation as a result. Three potential mechanisms can cause GLCS treatment to be ineffective: inadequate receptor function; an excess of proinflammatory cytokines, which would reduce affinity between the drug and its intracellular receptor; and a decrease in intracellular corticosteroids secondary to expulsion from the cell^[21]. This latter mechanism is dependent upon glycoprotein P-170 (P-gp), which is found in lymphocytes and in the apical membrane of the enterocytes, among other locations. An increase in P-gp at cell surface level causes drug release into the bloodstream. This protein is encoded by the ABCB1/MDR1 gene of chromosome 7. The expression of this gene is reportedly increased in IBD presenting a greater need for surgery because of a poor response to drug treatment^[22]. Different allelic variants (the most widely studied being C3435T and G2677T) are associated with an increased risk of developing extensive UC, although no association to CD has been observed^[23]. Studies with larger patient series and stable corticosteroid doses are needed to determine the precise relationship between P-gp and the lack of response to such drugs.

The studies that have explored the different cytokines implicated in corticosteroid response offer contradictory results, and the underlying polymorphisms have not been established^[24].

Genetic studies related to the gene encoding for the intracellular glucocorticoid receptor (*hGR*) have also been performed. Polymorphism N363S is associated with a good response^[25], while polymorphism ER22/23EK is associated with corticosteroid resistance^[26]. Knowing the genetic susceptibility of corticosteroid resistant patients is an important step forward, since it would help avoid important morbidity among patients who stand to derive no benefit from such treatment. None of these pharmacogenetic markers are of use in routine clinical practice.

Thiopurine drugs

Thiopurine drugs are used to maintain remission in patients with moderate to severe IBD. The effects are only observed after 3 mo of treatment. Purine metabolism is complex and involves different enzymes; this results in important genetic variability in the efficacy and toxicity of these drug substances (Figure 1).

The thiopurine drugs (TPs) are able to control the disease in 66% of patients, although an important

proportion (10%-25%) must suspend the medication because of serious (leukopenia, pancreatitis, infections, or malignancies) or mild side effects (rash, nausea, vomiting, flu syndrome, or joint pain). A clear association has been demonstrated between thiopurine methyltransferase (TMPT) deficiency and bone marrow suppression, although this explains only one-third of adverse effects. No clear explanation has been found for the remaining effects. Knowing in advance whether these drugs will be tolerated and the risk of side effects in a given patient may prove useful in daily practice.

TPMT: TPMT is the most widely studied enzyme in IBD. The identification of genetic mutations before onset of treatment with TPs is currently the only pharmacogenetic test performed in IBD.

In 1980, Weinshilboum and Sladeck were the first to describe the trimodal distribution of TMPT in Caucasian patients: 90% of the subjects have normal TMPT activity, almost 10% have intermediate activity, and 0.3% have almost zero activity. Posteriorly, over 30 allelic variants have been described, with different distributions depending on the ethnic group considered. The correlation between genotype and phenotype (expressed enzyme activity) is very good in 77%-99% of the cases. The differences can be explained by genetic and epigenetic factors (such as the use of concomitant drugs that inhibit TMPT), the age of the patient, and the existence of recent transfusions - since two different enzyme populations (donor and recipient) may be measured^[27,28]. The genetic study of this enzyme allows us to distinguish among homozygous individuals (without enzyme activity) at a high risk of suffering bone marrow suppression; ultra-fast methylators (high enzyme activity) with high liver toxicity and a low response to treatment; and patients with normal and intermediate enzyme activity, which are the individuals that stand to benefit most from this medication and are at lesser risk of adverse effects. Furthermore, it can help determine which dose a treatment should be started (Table 1).

Allopurinol reduces the levels of 6-methylmercaptopurine (6-MeMp) and increases those of 6-thioguanine (6TG). Although its mechanism of action is not clear, it has been suggested that the drug inhibits the enzyme xanthine oxidase through competitive inhibition or reduces the availability of its substrate^[29,30]. In daily practice, the TP dose should be reduced by 25% in those patients who require allopurinol and present with a normal TPMT genotype (Table 1). Taking into account that only about one-third of all cases of bone marrow suppression in patients receiving TPs are explained by genetic disposition and that the origin is, therefore, multifactorial and will require constant laboratory test monitoring, many authors have questioned whether this strategy is cost-effective. Nevertheless, most current clinical

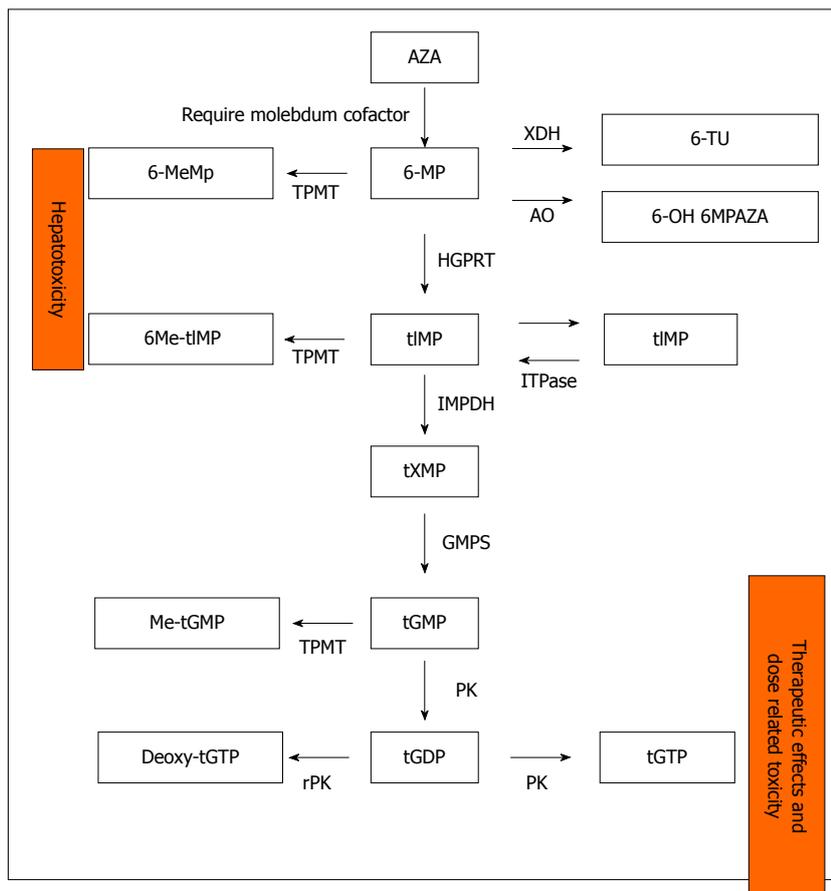


Figure 1 Metabolism of azathioprine and 6-mercaptopurine. AZA: Azathioprine; 6-MP: 6-mercaptopurine; 6-MeMP: 6-methylmercaptopurine; 6-TU: 6-thiouric acid; 8-OH 6MP: 8-hydroxymercaptopurine; 6Me-tIMP: 6-methyl thioinosine monophosphate; 6-tIMP: 6-thioinosine monophosphate; 6-TIDP: 6-thioinosine diphosphate; 6-TITP: 6-thioinosine triphosphate; 6-MeTITP: 6-methylthioinosine triphosphate; tXMP: Thioxanthine monophosphate tGMP: Thioguanine monophosphate; tGDP: Thioguanine diphosphate; tGTP: Thioguanine triphosphate; Me-tGMP: Methylthioguanine monophosphate; Deoxy-tGTP: Deoxythioguanine triphosphate; GST: Glutathione S-transferase; TPMT: Thiopurine methyltransferase; XDH: Xanthine dehydrogenase; AO: Aldehyde oxidase; HGPRT: Hypoxanthine guanine phosphoribosyltransferase; IMPDH: Inopine monophosphate dehydrogenase; GMPS: Guanosine monophosphate synthetase; PK: Phosphokinase; rPK: Reductase phosphokinase.

Table 1 Azathioprine treatment basis of individual thiopurine methyltransferase status

TMPT genotype	TPMT phenotype (pmol/mgHb)	Treatment dosis (mg/kg)
Homozygous	< 10	Avoid or consider 0.1-0.2
Heterozygous	10-24	1-1.5
Wild type (normal)	25-35	2-2.5
Wild type (high)	> 35	0.5 + 100 mg allopurinol

TMPT: Thiopurine methyltransferase.

guides do not recommend genetic study before starting treatment with TPs. On the other hand, such studies are not available in many hospitals; laboratory test monitoring is, therefore, the safest alternative.

Less scientific evidence is available regarding the other enzymes.

The enzyme xanthine oxidase (XO) is the second most important enzyme in the metabolism of TPs. It is found in many tissues, although its main activity is located in the small bowel and liver. Until a few

years ago, it was only known that XO activity varied from one person to another. Discordant variations depending on patient gender or ethnicity had been described^[31,32], together with changes induced by environmental factors, such as smoking and the diet. In 2008, a Japanese group described three polymorphisms of the gene encoding for XO (G514A, A3326C and A3662G) that are linked with enzyme activity, with the population being divided into low, normal, or high activity^[33]. These polymorphisms have not been studied in Caucasians or in patients with IBD. At present, we are only able to extrapolate that individuals with low enzyme activity would be at greater risk of suffering adverse effects, while those with high activity would experience treatment failure.

The enzyme inosine triphosphate pyrophosphatase (ITPase) controls the intracellular levels of inosine triphosphate (ITP), transforming it into inosine monophosphate, which acts as a substrate for other enzymes. When the enzyme is deficient, ITP accumulates in enterocytes. Deficiencies of this enzyme have been known for almost half a century and

Table 2 Genetic association studies of adverse events secondary to azathioprine in inflammatory bowel disease patients

Study (year)	No. of patients	Conclusion
Marinaki <i>et al</i> ^[34] (2004)	130	Significant association with flu-illness, rash, and pancreatitis No association with mielotoxicity
Allorge <i>et al</i> ^[35] (2005)	72	No association with flu-illness, rash, pancreatitis, or mielotoxicity
Gearry <i>et al</i> ^[36] (2004)	147	No association with flu-illness, rash, and pancreatitis
De Ridder <i>et al</i> ^[37] (2006)	72	No association with side effects
Hindorf <i>et al</i> ^[38] (2006)	60	No association with side effects
von Ahsen <i>et al</i> ^[39] (2005)	71	Early withdrawal of therapy but no association with specific adverse events
Ansari <i>et al</i> ^[40] (2008)	202	Association with flu-like symptoms but not withdrawal of therapy
van Dieren <i>et al</i> ^[41] (2005)	109	Not associated with an increased risk for the development of leucopenia and other side effects
Zelinkova <i>et al</i> ^[42] (2006)	262	Increased risk of leucopenia
Uchiyama <i>et al</i> ^[43] (2009)	16	Increase risk of mielotoxicity (leucopenia)
Shipkova <i>et al</i> ^[44] (2011)	160	Increase risk of mielotoxicity
Kim <i>et al</i> ^[45] (2010)	248	No association with leucopenia
Zabala-Fernández <i>et al</i> ^[46] (2011)	232	Significant association with artralgia

have been evaluated in different ethnic groups (with incidences of 5%-7% among Caucasians and African populations and up to 15% in Asian populations). Five polymorphisms have been described, of which only two are associated with enzyme inactivity: C94A, with an activity between 0% and less than 25% of normal and IVS2 + 21AC, with an activity of 60% of normal^[24].

The results from IBD studies are conflicting regarding side effects (pseudoinfluenza syndrome, rash, and pancreatitis) and bone marrow toxicity of azathioprine (Table 2). These studies, in general, have few patients and an even smaller number of patients with the relevant polymorphisms and have been carried out in different ethnic groups. Reliable conclusions, therefore, cannot be drawn. Pseudoinfluenza syndrome causes a large number of patients to abandon the medication. Currently, the importance of the genotype of ITPase in relation to side effects and early treatment suspension is not known.

Prior to the year 2006, it was believed that the conversion of azathioprine to 6-mercaptopurine (6-MP) was not mediated by enzyme action. That year, Eklund demonstrated that it was catalyzed by the glutathione-S-transferases (GSTs)^[47]. This author analyzed 14 variants, where GST-A1, GST-A2, and GST-M1 were the three with the highest enzyme activity. All of them are polymorphic. The studies in patients with IBD only allow us to affirm that individuals with low enzyme activity will have low 6-MP levels and, therefore, will not respond to the medication, while patients with ultra-fast activity are at an increased risk for adverse effects due to high 6-MP levels. If this is confirmed, the clinical application would be evident, since the problem could be overcome by directly prescribing 6-MP at the correct dose.

Other enzymes, such as inosine monophosphate dehydrogenase, hypoxanthine phosphoribosyl-transferase, *etc.*, have been studied, although their application at the present time is unclear.

Likewise, studies have attempted to determine the relationship between the levels of metabolites

(6-TG and 6MeMP) and the response to treatment or development of adverse effects. Once again, the results have been contradictory, with some studies describing a relationship between activity and the metabolite levels, while others indicating the opposite. A meta-analysis suggested that 6-TG levels above 260 pmol/8 × 10⁸ imply that the patient has a greater probability of disease remission^[48]. Blood 6-TG levels above 400 pmol/8 × 10⁸ red blood cells increase the risk of bone marrow toxicity^[49], and 6-MeMP levels above 5700 pmol/8 × 10⁸ red blood cells increase the risk of liver toxicity^[50,51]. At present, metabolite determination is not available on a generalized basis in clinical practice, and its use is controversial. However, in patients lacking a clinical response, it may help in deciding medication changes (Figure 2).

Methotrexate

Methotrexate (MTX) is usually used as an alternative to treatment with TPs in CD both for flare-ups and as a maintenance therapy. Its usefulness in UC is more controversial.

The mechanism of action of MTX in IBD has not been clearly established. The drug is a folic acid antagonist and blocks purine and pyrimidine synthesis. Those tissues characterized by greater cellular regeneration (turnover) show more toxic effects. Consequently, the main adverse effects of MTX are bone marrow suppression, mucositis, and gastrointestinal and hepatic alterations. Folic acid supplementation reduces these side effects, although up to 30% of all patients have to suspend the medication.

Most pharmacogenetic studies of MTX have focused on patients with hematological tumors and rheumatoid arthritis (RA). In both of these disease conditions, the dosage and administration route are very different from those used in IBD; the extrapolation of results is therefore not possible.

Herrlinger has carried out the only study to date in patients with IBD. Patients with the 1298C allele of the enzyme MTHFR are more susceptible to adverse

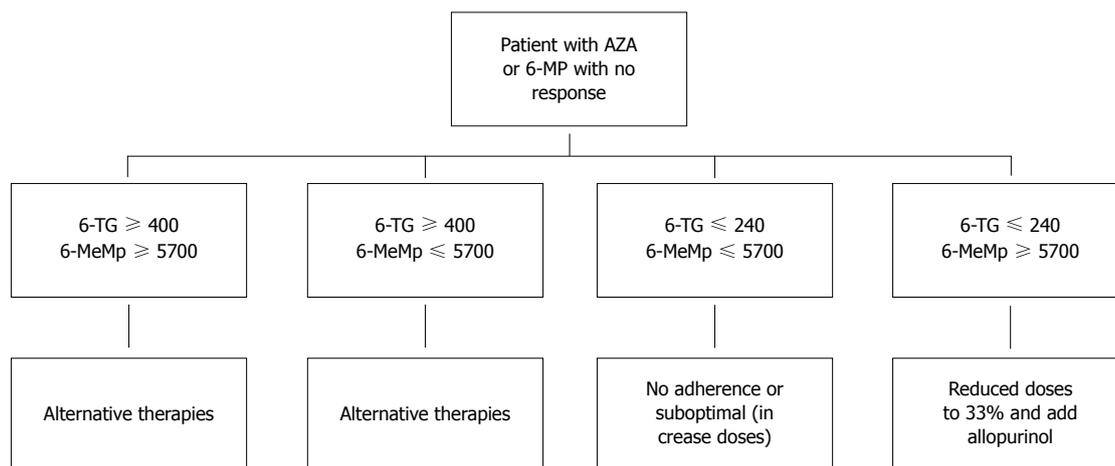


Figure 2 Algorithm of treatment for non responders to thiopurine drugs. AZA: Azathioprine; 6-MP: 6-mercaptopurine; 6-TG: 6-thioguanine; 6-MeMp: 6-methyl mercaptopurine.

effects, although these data are in contrast with those recorded for other diseases^[52].

Anti TNF- α drugs

Infliximab (IFX) is a chimeric IgG1 monoclonal antibody (Ab) targeted to TNF- α . It is used for induction and maintenance in patients with moderate to severe flare-ups of IBD. IFX is very useful and has been shown to produce mucosal healing and to reduce the number of flare-ups, hospital admissions, and surgeries. A number of clinical factors indicative of a good response have been described, such as the start of patient treatment at a young age, colonic location of the disease, and associated immunosuppressor therapy. It seems that a shorter duration of the disease, non-smoking, and elevated C-reactive protein levels at the start of therapy also favor a good response^[53,54]. Even so, 25% of all patients are primary non-responders, and 20%-30% lose responsiveness over time (secondary non-responders). Most of these latter cases are a consequence of the production of antibodies against IFX^[11-13]. Furthermore, the drug is expensive and has potentially serious side effects.

Table 3 summarizes the genetic studies on treatment of IBD with IFX. Studies have been carried out on genes that encode TNF- α , the TNF- α receptor, genes regulating the expression of TNF- α (NOD2/CARD2), apoptotic mechanisms, and other proinflammatory cytokines. To date, it has not been possible to demonstrate an association between drug response and a specific gene. There are two main issues: on the one hand, many studies failed to reach statistically significant results because of the small number of patients involved, and on the other hand, the studies are not always comparable, since different response criteria were used (C-reactive protein, CDAI score, Harvey-Bradshaw index), patients with different degrees of disease activity were included, and different doses were administered.

Perhaps, in the near future, information regarding the genotype of TNF- α and its receptor may help us to identify non-responders to anti-TNF- α therapy.

PHARMACOKINETICS

Understanding the pharmacokinetics of a drug is very important when adjusting the dose required to guarantee therapeutic concentrations, since many factors influence drug concentration in blood. In clinical practice, the determination of blood drug levels has been used to monitor treatments with drug substances that have a narrow therapeutic margin or window. In IBD, such monitoring has been applied to cyclosporine A and tacrolimus; and in recent years, it has become particularly important in the management of anti-TNF- α drugs.

Differences in the administration route, degradation, and clearance of anti-TNF- α determine its concentration in blood and, in turn, its treatment response. In this regard, achieving adequate blood drug levels is correlated with clinical and endoscopic remission of the disease^[71,72]. When the anti-TNF- α drug is administered intravenously (*e.g.*, IFX), the maximum blood drug concentration is reached immediately after infusion, with little variability among patients. However, when the anti-TNF- α drug is administered subcutaneously (*e.g.*, antidrug antibodies (ADAs), golimumab, and certolizumab), the maximum concentration is reached after approximately 10 days, and the bioavailability ranges between 50%-100%^[73,74].

The clearance of anti-TNF- α drugs from blood is complex and multifactorial. Variables that increase drug clearance include those that depend upon or reflect the severity of the disease (hypoalbuminemia, decreased hemoglobin levels, C-reactive protein elevation, TNF- α , leukocytosis, and increased IFX intestinal losses), demographic parameters (increased body mass, male gender, and age under 40 years) and immunogenicity

Table 3 Genetic association studies of infliximab response in inflammatory bowel disease patients

Study (year)	Patients recruited	Response criteria	Genes investigated	Conclusion
Taylor <i>et al</i> ^[55] (2001)	75	CDAI	Polymorphisms TNF/LTA region	Homozygosity for the LTA 1-1-1 haplotype may identify subgroups with poorer response
Louis <i>et al</i> ^[56] (2002)	226	CDAI	TNFA	No association with treatment outcome
Mascheretti <i>et al</i> ^[57] (2002)	90	CDAI	TNF and TNFR polymorphism	196Arg homozygotes had poorer clinical response than 196Met heterozygotes and homozygotes ($P = 0.036$)
	444			No predictive treatment outcome
Mascheretti <i>et al</i> ^[58] (2002)	534	CDAI	CARD15/NOD2	A strong relation to susceptibility to CD but not association with treatment outcome
Vermeire <i>et al</i> ^[59] (2002)	245	CDAI	CARD15/NOD2	Not predictive of treatment outcome
Pierik <i>et al</i> ^[60] (2004)	166	CDAI	TNF/TNFR	Biological response to infliximab was lower in patients carrying TNFR1-36G
Matsukura <i>et al</i> ^[61] (2008)	80	HBI	TNFRSF1A TNFRSF1B	28% of G allele heterozygotes and homozygotes responded compared to 73% of A allele homozygotes ($P = 0.04$) 5% of patients with AT haplotype responded compared to 20%-40% of patients with other haplotypes ($P = 0.01$)
Louis <i>et al</i> ^[62] (2004)	200	CDAI	FcγRIIIa	Positive (V/V genotype) association with good treatment outcome
Urcelay <i>et al</i> ^[63] (2005)	40	CDAI	IBD5(5q31)	Polymorphisms TT is associated with negative response
Hlavaty <i>et al</i> ^[64] (2005)	287	CDAI	FASL/CASP9	Positive association
Hlavaty <i>et al</i> ^[65] (2007)	287	CDAI	FASL	Negative association (statistical model)
Willot <i>et al</i> ^[66] (2006)	189	CRP	CRP	Polymorphisms evaluated are not associated with treatment outcome
Dideberg <i>et al</i> ^[67] (2006)	214	CDAI	TNF/LTA region	No association
Dideberg <i>et al</i> ^[68] (2006)	186	CDAI and CRP	ADAM17	Minor allele homozygotes for each SNP associated with clinical response ($P < 0.002$)
Jürgens <i>et al</i> ^[69] (2010)	90	CAI	IL-23R IL-2/IL-21	Homozygous carriers of IBD risk-increasing IL-23R variants more likely to respond to infliximab than homozygous carriers of IBD risk-decreasing IL-23R variants ($P = 0.001$)
Dubinsky <i>et al</i> ^[70] (2010)	94	HBI and Partial Mayo score	rs2241880 2q37/ ATG16L1 rs2188962 5q31 rs6908425 6p22/ CDKAL1 rs762421 21q22/ ICOSLG rs2395185 6p21/ HLA-DAQ1 rs2836878 21q22/ BRWD1	Six known susceptibility loci associated with primary nonresponse ($P < 0.05$). Only the 21q22.2/BRWD1 loci remained significant in the predictive model

Available from: URL: <http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4419078/>.

(the development of antibodies against the drug)^[75]. The concomitant use of immunosuppressors (TP drugs and MTX) is associated with decreased immunogenicity and increased anti-TNF- α drug levels.

Up to 40% of all patients that respond to anti-TNF- α drug treatment will require one or more dose adjustments in order to maintain treatment efficacy. After 1 year of treatment, efficacy is maintained in only one-third of all responders^[76].

The need for dose adjustment of these drugs may occur at two time points during treatment: at the start (primary failure) or during the maintenance phase (secondary failure).

Primary failure, or a primary lack of response, refers to the absence of improvement in signs and symptoms of the disease that leads to treatment suspension during the induction phase^[77]. The time point at which this primary lack of response is assessed varies among different studies. In patients treated with IFX, as in the ACCENT I trial^[78], assessment was made 2 wk after the

first IFX infusion. In contrast, assessment in the ACCENT II trial was made 10-14 wk after induction^[79]. In studies including patients treated with ADA, the response was assessed in week 4 in the CLASSIC I trial^[80] and in week 6 in the CLASIC II trial^[81]. Among the different published studies, the lack of primary response to treatment rate with anti-TNF- α drugs varied from 10%-40% and was found to be higher in UC than in CD. Lack of response was noted particularly in severe presentations of UC. This observation could be explained by increased drug clearance as a result of a greater inflammatory intestinal surface^[77].

Secondary failure, or secondary lack of response, is defined as failure occurring in the course of treatment in those patients that have responded to induction therapy and is observed in approximately 13% of the patients treated with IFX^[82] and in 10%-24% of those treated with ADA^[83]. Failure is more frequent in the first year of therapy.

Table 4 and Table 5 describe the risk factors asso-

Table 4 Risk factors associated to primary failure

Crohn's disease	Ulcerative colitis
Duration of the disease > 2 yr	Old age
Smoking	Anti-neutrophil cytoplasmic (ANC) antibodies
Extensive small bowel involvement	Negative antibodies against <i>Saccharomyces cerevisiae</i>
Low C-reactive protein levels	Previous exposure to anti-TNF- α drugs
Genetic mutations or polymorphisms of the apoptosis and caspase 6 genes and locus IBD 5	

IBD: Inflammatory bowel disease.

Table 5 Risk factors associated to secondary failure or loss of response

Individual differences in bioavailability and pharmacokinetics
Symptoms not due to inflammatory bowel disease
Lack of adherence to therapy
Drug loss in stools
Intermittent treatments
Non-inflammatory symptoms
Structuring disease pattern
Smoking
Development of antibodies

ciated with primary and secondary failure^[11,84].

In secondary failure, the most widely investigated factor is the development of ADAs. When a loss of treatment response occurs and once the possible existence of intercurrent processes and lack of adherence to therapy have been discarded, the patient requires dose adjustment or a switch to some other molecule. In this context, it is very useful to know the blood drug levels and whether or not ADAs have developed.

The development of ADAs is conditioned by the patient immune condition and is much more common in intermittent treatments (over 60% of the cases) than in maintenance therapy (10%-20%). Furthermore, those patients who have developed ADAs and remain without treatment for extended periods exhibit slow clearance of these antibodies^[85].

Many studies have been carried out to determine when and how antibody titers should be measured in order to perform the necessary adjustments and to define the interval during which therapeutic blood drug levels are present. In routine clinical practice, if this is not possible, dose adjustment is made empirically, shortening the interval between doses or switching to another molecule when loss of response occurs or an immune-mediated reaction is observed. Three different methods can be used to measure drug levels.

The most widely used option is the enzyme linked immunosorbent assay (ELISA), which can also be used to determine ADA titers. This technique requires the anti-TNF- α drug to be undetectable in blood, since it

binds to the antibodies and forms immune complexes that are not detected.

Radioimmunoassay (RIA) is similar to ELISA, although its use is limited since it involves the use of a radioactive reagent. Furthermore, as for ELISAs, it cannot detect the presence of ADAs if there are detectable drug levels present in blood.

Another method for determining the drug and antibody is variable mobility testing, which allows for the detection of ADAs (against IFX and ADA) in the presence of drug in blood.

The determination of anti-TNF- α drug levels is performed immediately before administration of the next dose. ADAs are only detected when the drug levels are undetectable, since the most widely used technique in clinical practice is the ELISA.

Different studies have correlated the development of ADAs with an increased risk of infusion reactions and increased drug clearance from blood^[86]. However, not all studies have established correlations to loss of response. In this regard, a systematic review published by Chaparro *et al.*^[87] found no differences between maintenance or loss of response according to whether ADAs develop or not, while the meta-analysis conducted by Nanda *et al.*^[88] found that patients who develop ADAs have a 3-fold greater risk of loss of treatment response than patients who do not develop such antibodies.

Although there is great variability among studies in defining the minimum effective concentration of anti-TNF- α drugs, the determination of drug levels has been correlated with improved disease control and to clinical and endoscopic remission. This is important in designing treatment algorithms^[71]. To date, the management approach in clinical practice depends on the drug values and on the presence or absence of ADAs (Table 6).

Current studies are evaluating the role of routine anti-TNF- α drug level measurements in blood during therapy, as is done with other drug substances, in order to facilitate better dose adjustment^[86]. However, the different studies propose different cutoff values when defining the therapeutic range. Therefore, studies are needed to determine the adequate therapeutic interval or window.

FUTURE TREATMENTS

The biological drugs currently authorized for the treatment of IBD are monoclonal antibodies targeted to TNF- α (IFX, ADA, certolizumab, and golimumab) and monoclonal antibodies targeted to the leukocyte integrins (natalizumab, approved by the United States Food and Drug Administration (FDA) for refractory CD; and vedolizumab, approved by the European Medicines Agency (EMA) and the FDA).

A number of lines of research have been developed with the aim of blocking the inflammatory process at

Table 6 Treatment algorithm according to antidrug antibodies and drug levels

Anti-TNF- α drug levels	Antibodies	Action
Low	Negative	Increase dose
Low	Positive	Switch drug
High	Not determined	Switch to a drug with a different mechanism of action

different levels. In the coming years, new drugs will be introduced that will contribute to the expansion of therapeutic options.

The drug options that are presently in the most advanced stages of development are described below.

Interleukins

Interleukins (ILs) are soluble inflammatory response messenger (signaling) molecules. Their role in IBD has been clearly established, and research in this field has been particularly wide-ranging and advanced^[89].

Anti-IL-12/23 drugs: Ustekinumab is the drug with the most advanced results available to date. Ustekinumab is a fully humanized IgG1k anti-IL-12/23 monoclonal antibody. It specifically binds to the p40 protein subunit shared by both of the mentioned ILs. Binding prevents the mentioned subunit from interacting with the IL-12R β 1 receptor protein, which is expressed on the surface of immune cells - thereby inhibiting innate and adaptive immune response stimulation. In an inflammatory environment, naïve CD4+ T cells are induced to interferon (IFN)- γ producing Th1 cells by the action of IL-12 and to Th17 cells by the action of IL-23. The Th17 cells, in turn, are responsible for the production of proinflammatory cytokines, such as IL-17, IL-17F, IL-6, and TNF- α ^[90,91]. Blocking this pathway has been successfully employed in animal models^[92,93], and both ILs play a key role in the inflammatory processes of CD^[91,94,95].

The results of a first double-blind and placebo (PB)-controlled phase II clinical trial were published in 2008^[96]. This study had a complicated design that included two patient populations. In population 1, the results at the primary endpoint were discouraging, with a clinical response rate in week 8 of 49% in the group of patients treated with ustekinumab vs 40% in the PB group ($P = 0.34$). However, in a subgroup of 49 patients previously treated with IFX, statistical significance vs PB was reached, with a response rate of 59% and 26%, respectively ($P = 0.05$), in week 8. In population 2, the clinical response rate with ustekinumab in week 8 was 43% in the subcutaneous treatment group and 54% in the intravenous treatment group. Failure of the study to confirm the primary endpoint was attributed to the high percentage response observed in the PB group. No serious adverse events were detected in week 8, and the recorded problems were similar to those

seen in the PB group. Overall, the final conclusion was favorable regarding the capacity of the active drug to elicit a clinical response in the induction phase.

The CERTIFI trial was published in 2012^[97]. This was a randomized, double-blind, PB-controlled phase II a study on the efficacy of ustekinumab in patients with moderate to severe CD refractory to IFX. A total of 526 patients resistant to treatment (50% of the subjects having received at least two anti-TNF- α drugs) were randomized to three intravenous induction treatment arms (1, 3, or 6 mg/kg of ustekinumab) and a PB arm. The 145 patients responding to ustekinumab in week 6 were randomized to subcutaneous maintenance therapy in weeks 8 and 16 with PB vs ustekinumab 90 mg. The primary endpoint was the clinical response rate in week 6, where the recorded percentages were 36.6%, 34.1%, and 39.7% (1, 3, or 6 mg/kg of ustekinumab) vs 23.5% in the PB arm. Only the 6 mg dose reached statistical significance vs PB. In the maintenance phase, 41.7% of the patients treated with ustekinumab showed clinical remission vs 27.4% of the patients in the PB arm ($P = 0.03$), and the clinical response rate was 69.5% vs 42.5%, respectively ($P < 0.001$). The safety profile was found to be similar to that of other biological drugs.

The publication of the results of three phase III clinical trials is currently pending. The first of these studies (UNITI-1)^[98] is a randomized, double-blind trial designed to evaluate the efficacy and safety of induction with ustekinumab in patients with moderate to severe CD who have failed or are intolerant to anti-TNF- α therapy. The primary endpoint is clinical response in week 6, while the secondary endpoints are remission and clinical response in week 8. A total of 769 patients have been randomized to three arms: (1) intravenous PB; (2) intravenous ustekinumab 130; and (3) intravenous ustekinumab 6 mg/kg as a single dose. The study ended in July 2013. The second study (UNITI-2)^[99] has the same design as the first, although the included patients are naïve to biological drugs and show failure or intolerance to immunosuppressors or corticosteroids. This study ended in October 2014 and includes a total of 642 patients. Lastly, the IM-UNITI^[100] trial is also a randomized, double-blind, PB-controlled, parallel group multicenter study. This trial was designed to determine efficacy in the maintenance phase of CD and is currently in the recruitment stage. The study plans to include 1310 patients from the two previously mentioned trials, with conclusion in November 2018.

There are two other anti-IL-12/23 molecules: briakinumab (ABT-874), which targets the p40 subunit, and apilimod mesylate, which is a small molecule administered *via* the oral route that inhibits the transcription of IL-12 and IL-23. The initial results with both molecules have not been significant^[101,102].

Anti-IL-6 drugs: Interleukin-6 is a proinflammatory cytokine produced by different types of cells. It

participates in a series of processes including T lymphocyte activation and immunoglobulin secretion through the differentiation of B cells into plasma cells^[103,104]. Interleukin-6 exerts its action *via* membrane or soluble receptors^[105]. In healthy individuals, the IL-6 levels are low and increase in the context of immune processes^[103]. This cytokine is increased in CD in the same way as its soluble receptor, and its levels are correlated with the C-reactive protein concentrations^[106,107].

Tocilizumab is an IgG1 monoclonal antibody indicated in RA. It binds specifically to the soluble and membrane receptors of IL-6. In one study, 36 patients with active CD were randomly assigned to two treatment arms (intravenous 8 mg/kg every 2 wk or every 4 wk) or PB^[108]. The clinical response rate in the group administered tocilizumab every 2 wk was 80% vs 31% in the PB arm, although only 20% achieved clinical remission. The drug was well tolerated, although studies in RA have shown neutropenia, altered liver biochemical parameters, and hyperlipidemia. In this regard, dyslipidemia might prove to be a safety problem of the drug long term^[109].

Two phase II studies involving two monoclonal antibodies (BMS-945429, formerly ALD518, and PF-04236921) targeted to IL-6 are currently ongoing. No results are yet available, since patient recruitment has ended only recently^[110,111].

Other anti-interleukin drugs: Interleukin-2 is another molecule that plays a key role in T cell activation and proliferation.

Basiliximab and daclizumab are monoclonal antibodies targeted to CD25, which is the alpha-chain of the IL-2 receptor. Both drugs are used for the prevention of renal graft rejection. Initial studies with basiliximab documented clinical remission in eight out of 10 patients with UC refractory to corticosteroid therapy^[112], and a later study recorded a remission rate of up to 65%, without a control group though^[113]. In the case of daclizumab, a comparative study vs PB failed to demonstrate positive efficacy results^[114].

IL-13 is produced by naive T cells and activates natural killer (NK) cells, which in turn synthesize IL-13. IL-13 has been shown to play a key role in the pathogenesis of UC^[115,116].

Two phase II trials (IMA-648 [arunkinzumab] and CAT-345 [tralokinumab]) published in 2014 randomized the patients to the antibody^[117,118] at different doses (in the case of the first study) or PB. Neither study recorded differences in terms of treatment response or clinical remission. Both studies documented a tendency towards lower activity index scores, and the safety profile was favorable. A third phase II study in patients with perianal CD has been designed to evaluate the efficacy and safety of another antibody targeted to IL-13 (QAX576) vs IFX for comparison^[119]. Ten patients have been included, and

results from the trial are pending after the end of the recruitment phase.

Vidofludimus is an immunosuppressor that inhibits the release of IL-17 and IFN- γ [by interfering with the janus kinase/signal transducer and activator of transcription (JAK/STAT) pathway and nuclear factor kappa B (NF- κ B)], which blocks the enzyme dihydroorotate dehydrogenase (DHODH). This is a small molecule administered *via* the oral route. The results of the ENTRANCE study, a non-controlled multicenter trial, were published in 2013^[120]. This study evaluated 34 patients with corticosteroid-dependent IBD (CD and UC) treated with vidofludimus 35 mg/d orally administered (*per os*, *po*) over 12 wk. The primary endpoint was clinical remission without corticosteroids in week 12, and this was reached by 56% of the patients with CD and 50% of those with UC. The drug was well tolerated, and no serious adverse events were reported - although additional studies with larger patient series are needed to confirm this finding.

Other targets currently under investigation are antibodies against IL-17 (AMG 827), although the study that has been closed because of worsening of patient symptoms^[121], IL-18 (GSK1070806)^[122], and IL-21 (PF-05230900)^[123]. These interleukins had been implicated previously in the pathogenesis of IBD^[124-126].

Anti-inflammatory interleukins: Lastly, studies have been performed to evaluate the efficacy of the anti-inflammatory interleukins IL-10, IL-11, and IFN- β . The results to date have not been encouraging^[127-131].

Chemokine antagonists

Chemokines are small cytokines that induce chemotaxis by interacting with the chemokine transmembrane receptors bound to protein G^[132,133]. Their ligand is CCL25, which in the intestine is fundamentally expressed by the luminal epithelial cells^[134]. In IBD, and especially in CD, high levels of CCL25 and of T lymphocytes expressing CCR9 have been detected^[135].

Vercirnon: CCX-282B (vercirnon) is a molecule administered *via* the oral route that acts as a CCR9 receptor antagonist and has been suggested to reduce lymphocyte trafficking towards the intestine^[136]. The data derived from the PROTECT-1 multicenter trial were published in 2013^[137]. This was a double-blind multicenter study randomizing 436 patients with CD to three vercirnon treatment groups (250 mg/d *po*, 250 mg twice daily *po*, 500 mg/d *po*) vs PB over an induction phase lasting 12 wk. Subsequently, in the case of clinical response, the patients were again randomized to vercirnon (250 mg twice daily *po*) vs PB until week 36. The results were not statistically significant, although in week 8 a tendency towards greater efficacy, in terms of clinical response, was observed in the group administered 500 mg/d vs PB

(49% vs 60% with odds ratio (OR) = 1.53, $P = 0.111$). In week 52, significance was reached in terms of the percentage of patients in remission (47% vs 31% with OR = 2.01, $P = 0.12$). The safety profile was found to be favorable, with similar serious and non-serious adverse events rates in both groups.

BMS-936557: Regarding UC, a parallel line of research is based on IFN- γ -induced protein 10 (IP-10, also known as CXCL10). This protein is secreted by monocytes, endothelial cells, and fibroblasts in response to stimulation by IFN- γ ^[138]. This protein is implicated in chemotaxis and interaction phenomena with T cells through the CXCR3 receptor^[139]. Inhibition of this pathway could incline the Th1 response towards a Th2 response^[140]. In patients with UC, CXCL10 is over-expressed in plasma and colon tissue^[141], and in animal studies it has shown a reasonable efficacy profile warranting the start of evaluations in humans^[142]. BMS-936557 (MDX-1100) is a monoclonal antibody targeted to protein IP-10. Its safety profile has been evaluated in phase I studies^[143], and based on the evidence obtained in patients with RA^[144], a specific double-blind multicenter trial on active UC has been started^[145]. This phase II trial randomized 109 patients to PB or four doses of intravenous 10 mg/kg BMS-936557 every 2 wk. The primary endpoint (clinical response on day 57) was better among the patients that had received the antibody, although statistical significance was not reached (52.7% vs 35.2%, $P = 0.083$). Nevertheless, the results were statistically significant in the group of subjects with the highest titers in blood (87.5% vs 37%, $P < 0.001$). The number of infections was greater in the active treatment group (12.7%) compared with PB (5.8%), and drug suspension because of adverse events was necessary in 3.6% of cases. At present, we are awaiting the publication of the results of two other phase II trials randomizing patients to different doses of the molecule vs PB in application to moderate or severe CD^[146] or UC^[147].

Other chemokine antagonists: There are other lines of research involving chemokines in autoimmune diseases, such as the binding of CXCL10 to its CXCR3 receptor^[148], which has been shown to be reduced in patients with CD, in parallel to reduction in the C-reactive protein levels^[149]. On the other hand, it has been reported that patients with UC show serum and tissues elevations of exotaxin-1, a chemokine that acts by recruiting eosinophils at the intestinal level.

Bertilizumab is a humanized IgG4 monoclonal antibody that blocks exotaxin-1 activity^[150]. A phase II study involving 42 patients with moderate or severe UC is planned, with the aim of assessing the efficacy and safety of the drug^[151].

Endogenous anti-TNF: TNF-kinoid

Another line of investigation involves the generation

of anti-TNF- α polyclonal antibodies through active immunotherapy. This strategy is based on the use of a TNF- α derivative as a vaccine. The compound is known as TNK-kinoid (TNF-K), and while biologically inactive, it can interrupt B cell tolerance of their own cytokines, resulting in the production of high titers of antibodies^[152]. A first phase I/II study was presented at Digestive Disease Week in 2011^[153], involving 21 patients with moderate to severe CD assigned to different doses on an open-label basis. The safety profile was found to be favorable, with no serious adverse events, and antibodies were generated in 81% of the cases. Clinical remission in week 12 was achieved by 50% of the patients. The results of a second randomized phase II study were published in 2012, involving patients with moderate to severe CD and loss of response or with intolerance to conventional anti-TNF- α drug therapy. The trial included 68 patients randomized vs PB into two cross-over treatment arms with intramuscular doses of 180 μ g on days 0, 7, 28, 84, 91, and 112, with switching of the active drug to PB in the fifth dose and of PB to the active drug in the third dose in the control group. The safety data were again favorable in this case, with only one serious adverse event related to worsening of CD, although the efficacy results were not reported^[154].

Janus kinase antagonists

The Janus kinases (JAKs) are a group of proteins corresponding to enzymes associated to cytokine receptors. They form part of a complex system of signal transmission from outside the cell towards the nucleus, activating transcription of the genes that intervene in important cell processes, such as growth, differentiation, proliferation, or migration. The process begins when the membrane receptor is stimulated by a chemical messenger, *e.g.*, a cytokine. This receptor activates JAK, which undergoes auto-phosphorylation and, in turn, phosphorylates the STAT protein. The latter protein then binds to another phosphorylated STAT protein (*i.e.*, it undergoes dimerization) and is translocated to the cell nucleus where DNA transcription factors are activated. This system, known as the JAK/STAT system^[155], has been implicated in the pathogenesis of different diseases^[156,157], including specifically IBD^[158].

Tofacitinib: This is a small molecule administered *via* the oral route that selectively inhibits JAK1 and JAK3, affecting the signaling pathways of cytokines such as IL-2, 4, 7, 9, 15, and 21^[159,160]. Blocking these pathways could suppress the activation and proliferation of lymphocytes while maintaining T cell regulatory function^[161,162]. In 2002, the United States FDA approved the drug for the treatment of RA. In the case of IBD, the initial data are not encouraging in reference to CD regarding disease response or remission - although the drug is associated to a significant decrease in biomarkers (C-reactive protein

and calprotectin)^[162]. In reference to UC, the results of a randomized, double-blind phase II trial involving four treatment doses (0.5, 3, 10, or 15 mg/d po) or PB in 194 patients with moderate to severe UC were published^[163]. The primary endpoint was clinical response in week 8. In the high dose groups (10 and 15 mg, respectively), the clinical response was greater than in the PB series - statistical significance being reached with the 15 mg dose (61% and 78% vs 42%, $P = 0.10$ and $P < 0.001$). Clinical remission reached statistical significance vs PB with both doses (48% and 41% vs 10%, $P < 0.001$). A subsequent sub-analysis demonstrated improvement in patient quality of life^[164]. Further studies involving larger patient samples are needed to confirm the efficacy and safety of the drug.

Inhibition of IL-13: In this line of research, studies have been performed on the inhibition of the mentioned pathway with monoclonal antibodies targeted to IL-13, which is responsible for activating the JAK/STAT pathway^[165]. The studies are cited in the section on interleukin antagonist drugs.

Anti-adhesion molecules

Adhesion molecules constitute one of the most advanced lines of research in IBD. Adhesion molecules are transmembrane receptors with three domains (intracellular, transmembrane, and extracellular) that induce cellular changes following stimulation by external molecules. These molecules include the integrins and lymphocyte homing receptors.

Natalizumab: The first anti-adhesion drug investigated was natalizumab, and it consists of an IgG4 monoclonal antibody targeted to integrin subunit $\alpha 4$. Natalizumab initially showed favorable results in CD^[166,167], although the risk of progressive multifocal leukoencephalopathy (PML) secondary to reactivation of the JC virus has limited its use^[168,169]. Natalizumab acts by blocking the interaction of $\alpha 4\beta 7$ with mucosal vascular addressin cell adhesion molecule-1 (MAdCAM-1) and the interaction of $\alpha 4\beta 1$ with vascular cell adhesion molecule-1 (VCAM-1), which is critical to lymphocyte trafficking towards the central nervous system - thereby giving rise to the risk of JC virus reactivation^[170].

Vedolizumab: Vedolizumab is another IgG1 monoclonal antibody that binds to integrin $\alpha 4\beta 7$, preventing it from binding to its specific intestinal ligand, MAdCAM-1. As a result, T lymphocyte migration towards the inflamed intestinal areas is inhibited. In contrast to natalizumab, it does not bind to integrins $\alpha 4\beta 1$ and $\alpha E\beta 7$ and does not antagonize the interaction of integrin $\alpha 4$ with VCAM-1. At present, and based on the results of clinical trials^[171,172], vedolizumab has been approved by both the FDA and the EMA for the treatment of patients with moderate to severe CD or UC who fail to respond to conventional treatment or therapy with

anti-TNF- α drugs.

AMG 181: AMG81 is another humanized IgG2 monoclonal antibody likewise targeted to integrin $\alpha 4\beta 7$ ^[173]. Early-stage studies warrant the safety, pharmacological, and tolerability profile of the drug. AMG 181 is currently being investigated in the context of two randomized, PB-controlled trials in application to both CD^[174] and UC^[175]. More evidence will be obtained in the coming years.

AJM300: AJM300 is a small molecule administered *via* the oral route that inhibits the $\alpha 4$ receptor. It is known to inhibit the binding of integrin $\alpha 4\beta 1/\alpha 4\beta 7$ - expressing cells to VCAM-1/MAdCAM-1 and has efficacy in the prevention of colitis in animal studies^[176]. The results of a double-blind multicenter phase IIa trial were published in 2012^[177]. The study involved the randomization of 102 patients with active-moderate UC to receive AJM 300 at a dose of 960 mg or PB three times daily for 8 wk. The primary endpoint was the clinical response rate in week 8, which was found to be 62.7% vs 25.5% in the AJM300 group and PB group, respectively (OR = 5.35; $P = 0.0002$). The secondary endpoints (clinical remission and mucosal healing in week 8) were also favorable to the study molecule (23.5% vs 3.9% with OR = 7.81; $P = 0.0099$ and 58.8% vs 29.4% with OR = 4.65; $P = 0.0014$). No serious adverse events (progressive multifocal leukoencephalopathy) were documented over the short term.

Etrolizumab: Etrolizumab is a humanized monoclonal antibody targeted to the B7 subunit present in integrins $\alpha 4\beta 7$ and $\alpha E\beta 7$. The results of a double-blind multicenter phase II trial were published in 2013^[178]. The study involved the randomization of 124 patients with refractory moderate to severe UC to etrolizumab in two treatment arms (subcutaneous 100 mg monthly or subcutaneous 300 mg monthly plus a loading dose of subcutaneous 420 mg between weeks 0 and 2) or PB. The primary endpoint was the clinical remission rate in week 10, with significant results in both active treatment arms vs PB (20.5% and 10.3% vs 0%, $P = 0.004$ and $P = 0.049$). In the subgroup of patients naïve to anti-TNF- α drug treatment, the differences obtained with the 100 mg dose were even greater (43.8% vs 0%, $P = 0.007$). The adverse event rates were similar in all three groups.

MECA-367 and PF-00547,659: The pharmacological properties of two monoclonal antibodies (MECA-367 and PF-00547,659) targeted to MAdCAM, with inhibition of binding of the latter to integrin $\alpha 4\beta 7$, were presented in 2009^[179], but subsequent data are only available for PF-00547,659. In 2011, a double-blind study randomized 80 patients with active UC to different single or multiple subcu-

taneous or intravenous doses of PF-00547,659 vs PB. Good efficacy vs PB was recorded, with endoscopic improvement of the lesions, a good safety and tolerability profile, and no immunogenicity. Results of the TURANDOT trial were published this year^[180]. This is a double-blind, PB-controlled multicenter efficacy and safety study in 357 patients with moderate to severe UC randomized to PB or to doses of the antibody (7.5, 22.5, 75, or 225 mg every 4 wk for three doses). The primary endpoint was the remission rate in week 12, and the secondary endpoints were the response rate and mucosal healing rate in week 12. Remission and mucosal healing were significantly greater in the 22.5 mg and 75 mg dose groups vs PB, while response was significantly greater for the 22.5 mg and 225 mg groups vs PB. The safety profile remained favorable.

Other lines of research

Laquinimod: Laquinimod is a molecule administered *via* the oral route, with great bioavailability and with purported regulatory activity upon antigen-presenting cells (APCs) and T lymphocytes^[181,182]. A phase II trial randomizing 180 patients to different doses of active treatment vs PB reported that increasing dose was inversely proportional to the percentage response or remission^[183]. Specifically, the lowest dose (0.5 mg) elicited a clinical response in 55.2% of the patients treated with the active drug vs in 31.7% of the patients administered PB, with respective remission rates of 48.3% vs 15.9% in week 8.

Masitinib: Gastrointestinal mast cells are usually found beneath the epithelial surfaces and are able to release cytokines, chemokines, prostaglandins, histamine, and heparin. The proliferation of these cells increases at intestinal mucosal and submucosal levels in CD^[184,185]. Masitinib is a selective tyrosine kinase inhibitor that targets the c-kit receptor (expressed by mast cells), platelet-derived growth factor receptor- α/β , lymphocyte-specific kinase, Lck/Yes-related protein, fibroblast growth factor receptor 3, and the focal adhesion kinase activation pathway^[186]. A phase II b/III phase trial with 450 CD patients is currently underway with this molecule^[187].

Visilizumab, rituximab, and abatacept: Attempts have been made to invert the natural inflammation process in which T cell proliferation vs apoptosis is observed. Visilizumab (a humanized monoclonal antibody against T cell receptor CD3) and rituximab (a chimeric monoclonal antibody targeted to B cell receptor CD20) were evaluated in IBD, where they were shown to have an unfavorable safety profile^[188-193]. Abatacept is a recombinant protein that blocks T cell co-stimulation by the antigen-presenting cells (APCs). Its use has been approved for RA, although the results for IBD have been discouraging^[194,195].

Morgensen: Immunosuppressive cytokine transforming growth factor (TGF)- β 1 is a secreted cytokine with known functions in growth, proliferation, differentiation, and apoptosis. It has been linked to immune regulating functions, depending on the cell upon which it acts and the environment in which it is found. Diminished TGF- β 1 activity has been reported in CD. This is due to the binding of an intracellular protein called SMAD 7 to the TGF- β 1 receptor^[196]. A molecule known as Morgensen (GED301) was first described in 2001. This is an antisense oligonucleotide that hybridizes to the human *SMAD7* messenger RNA (mRNA) and facilitates RNase H-mediated RNA degradation through a classic antisense mechanism. Its release is pH-dependent; accordingly, it is released in the ileum and right colon^[196]. Favorable safety results from a phase I study in 15 patients with CD were published in 2012^[197]. Recently, a phase II trial has evaluated 166 patients with active CD assigned to three active drug treatment groups vs PB^[198]. The clinical remission rates associated with the two highest drug doses were 55% and 65% vs 10% for PB ($P = 0.001$), while the clinical response rates were 58% and 72% vs 18% ($P = 0.04$). The total and serious adverse event rates were similar, with no reported neoplasms.

RPC1063 and GLPG0974: Lastly, the results of two randomized, double-blind, PB controlled trials were presented at the European Crohn's and Colitis Organisation (ECCO) meeting in 2015 on two new molecules: RPC1063 and GLPG0974.

RPC1063 is a molecule administered *via* the oral route with selectivity for sphingosine 1-phosphate (S1P) 1 and 5 receptor modulator. The study included 197 patients with moderate or severe UC that were administered 0.5 mg or 1 mg of the active drug vs PB, once daily^[199]. The primary endpoint (the proportion of subjects in remission in week 8) reached statistical significance in the highest dose group (16.4% vs 6.2%, $P = 0.048$). The adverse events profiles were comparable between groups, with approximately 31% of patients experiencing a treatment emergent adverse event.

GLPG0974 is a selective free fatty acid receptor antagonist. Binding of the fatty acids to their receptor induces neutrophil activation and migration. The results were assessed in 45 patients with mild to moderate UC treated with GLPG0974 during 4 wk (200 mg/12 h *po* vs PB)^[200]. A decrease in calprotectin levels and myeloperoxidase-positive cells was recorded, although there was no difference in terms of response, clinical remission, or mucosal healing. The safety and tolerability profile was favorable.

Stem cells: Different stem cell therapies have been used in CD and UC. Stem cell therapy involves the use of autologous hematopoietic stem cell transplantation

Table 7 Treatment with herbal remedies

Author	n	Producto	Comparador	Indicación	Remisión/response herbal vs PB or drug (%)
Langmead	44	Aloe vera	Placebo	Induction remission CU	30 vs 7
Ben-Arye	23	Triticum aestivum	Placebo	Induction remission CU	91 vs 42
Khan	14	Bovine colostrum enema	Placebo	Induction remission CU	
Sandborn	224	HMPL-004	Placebo	Induction remission CU	38/60 vs 25/40
Fukunaga	30	Xilei-san suppository	Placebo	Induction remission CU	46 vs 0
Zhang	35	Xilei-san enema	Enema dexametasona	Induction remission CU	
Tang	120	HMPL-004	Mesalazina	Induction remission CU	21 vs 16
Gupta	30	Boswellia serrata	Sulfasalazina	Induction remission CU	70 vs 40
Cheng	153	Jian Pi Ling tablet	Sulfasalazina	Induction remission CU	53 vs 28/19
Wang	106	Kui Jie Quing enemas	Sulfasalazina Prednisolona	Induction remission CU	72 vs 9
Cheng	118	Yukui tang ablets	Prednisolona Neomicina Vitamina B	Induction remission CU	33 vs 17
Fernández Bañares	105	Plantago ovata sedes	Mesalazine	Maintenance remission CU	60 vs 65
Hanai	89	Curcumin	Placebo	Maintenance remission CU	95 vs 79
Greenfield	43	Oenothera biennis	Evening primrose oil and olive oil	Maintenance remission CU	
Omer	40	Artemisia absinthium	Placebo	Treatment and prevention recurrence EC	65 vs 0
Krebs	20	Artemisia absinthium	Placebo	Treatment and prevention recurrence EC	80 vs 20
Gerhardt	102	Boswellia serrata extract	Mesalazine	Treatment and prevention recurrence EC	36 vs 31
Ren	20	Tripterygium wilfordii	Placebo	Treatment and prevention recurrence EC	
Holtmeier	108	Boswellia serrata extract	Placebo	Treatment and prevention recurrence EC	60 vs 55
Tao	45	Tripterygium wilfordii	Mesalazine	Treatment and prevention recurrence EC	68 vs 61
Liao	39	Tripterygium wilfordii	Sulphasalazine	Treatment and prevention recurrence EC	94 vs 75

Adapted from Ng *et al*^[203].

in CD, mesenchymal stem cells administered systemically or locally in perianal fistulas, and other cell treatments where experience is more limited, such as regulatory T cells and dendritic cells. We refer readers to two recent reviews carried out by our group in this field^[201,202].

Herbal remedies: Studies have been made that specifically compare treatment with herbal remedies vs PB or even conventional therapy, although there are discrepancies in the results due to the lack of homogeneity among the different studies. The findings in terms of safety are favorable, and the predictable costs are lower than in the case of conventional treatment. We recommend a recent systematic review, which affords a more detailed analysis of this subject^[203] (Table 7). Among the different herbal remedies employed, special mention must be made of *Andrographis paniculata* extract, known as HMPL-004, which has been found to reduce TNF, IL-1 β , IFN- γ , and IL-22 in the development of experimental colitis^[204].

Fecal transplant: Fecal transplant is a therapeutic alternative in gastrointestinal processes, such as *Clostridium difficile*-infection, metabolic syndrome, constipation, pouchitis, irritable bowel syndrome, and IBD^[205]. In IBD, effectiveness appears to be related to the stability of the colonization of donated bacteria^[206]. The experience in CD is limited to six patients in total^[207,208]. In UC, there are data available for up to

106 patients^[206,208-212]. The study with more patients included 62 cases with UC, finding clinical improvement in 92% and clinical remission in 68%. In the remaining studies, the results have not been as favorable as the aforementioned studies, with clinical remission ranging from 0% to 30% and clinical response from 0% to 70%. The relatively small number of patients evaluated so far does not allow for the establishment of firm conclusions, but it stresses the importance of the microbiota in the pathogenesis of IBD.

CONCLUSION

At present there are a large number of ongoing studies in various stages of research on new molecules for the treatment of IBD. An analysis of mucosal healing is needed in order to evaluate fully the impact of these therapies. In this way, it is expected to change the course of treating IBD. Among the different alternatives, anti-adhesion molecules and interleukin drugs are promising anti-TNF- α treatments.

With developments in the near future in pharmacogenetics, clinical pharmacology, the use of indices that try to classify patients by defining profiles of severity, and new drug molecules, personalized tailoring of treatment strategies will be possible for IBD.

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