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# Novel Targets for Antiretroviral Therapy Clinical Progress to Date

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#### **Abstract**

The advent of HIV-1 resistance to antiretroviral medications, the need for lifelong antiretroviral therapy (ART) for HIV-infected individuals, and the goal of minimizing ART-related adverse effects and toxicity all drive the need for new antiretroviral drugs. Two new classes of antiretroviral medications for HIV treatment, the CCR5 and integrase inhibitors, have recently been approved for use in patients in whom previous HIV treatment regimens have failed. These new agent classes are a welcome addition to other antiretroviral classes, which include nucleoside reverse transcriptase inhibitors, non-nucleoside reverse transcriptase inhibitors.

Maraviroc is a CCR5 co-receptor antagonist that blocks HIV binding to the CCR5 receptor, which is a CD4 co-receptor necessary for cell entry. It is approved for use in ART-experienced patients with CCR5-tropic HIV, and was found to significantly reduce HIV viral load and increase CD4+ cell count when combined with an optimized background ART regimen (OBR). Treatment failure with maraviroc has been described and is primarily associated with the presence of CXCR4-tropic virus. Vicriviroc is another CCR5 co-receptor antagonist that is in late clinical trials.

Raltegravir is the first US FDA-approved HIV-1 integrase inhibitor. It is approved for use in ART-experienced patients and was found to significantly reduce HIV viral load and increase CD4+ cell counts compared with placebo in combination with an OBR. Raltegravir has also been studied in treatment-naive patients and was found to be non-inferior to an efavirenz-based regimen. Elvitegravir is another HIV-1 integrase inhibitor in clinical development.

Other new antiretroviral agents in clinical development include PRO140, a monoclonal antibody against CCR5, and bevirimat, a maturation inhibitor that prevents late-stage gag polyprotein processing. A number of other drug targets, such as CCR5 co-receptor agonists, CXCR4 co-receptor antagonists, novel fusion inhibitors, and alternative antiretroviral strategies, such as immune stimulation and gene therapy, are under investigation.

Antiretroviral therapy (ART) has improved life expectancy for HIV-infected patients. [1,2] For most patients, HIV has become a chronic illness,

requiring lifelong, uninterrupted ART, and attempts to suspend or stop treatment have resulted in inferior clinical outcomes.<sup>[3-5]</sup> Although ART

has resulted in virological suppression, improved immune function and improved quality of life, issues such as tolerability, drug-drug interactions and resistance persist.<sup>[6-9]</sup> Thus, there is a need for antiretroviral agents with new mechanisms of action, and different resistance and toxicity profiles.

The HIV replication cycle is well understood, providing many opportunities for drug targets. Major steps in the HIV life cycle that are targeted by antiretrovirals include attachment, entry, reverse transcription, integration, protein processing and maturation. Viral entry to the host cell begins with attachment of the HIV envelope glycoprotein (gp120) to the CD4 T-cell receptor.[10] This interaction produces a conformational change in gp120, which allows it to bind to the chemokine co-receptor, either CCR5 or CXCR4. CCR5 and CXCR4 bind the V3 region of HIV gp120, which is the site of resistance mutations to CCR5 receptor antagonists (or inhibitors).[11] Further conformational changes in gp120 occur after CCR5 or CXCR4 binding, allowing the HIV envelop gp41 fusion peptide to insert into the cellular membrane, causing the virion to fuse with the viral membrane.<sup>[12]</sup> Once the HIV virion enters the cell, it is uncoated and the viral RNA genome undergoes reverse transcription into proviral DNA. The proviral DNA becomes part of the pre-integration complex, which enters the nucleus.<sup>[13]</sup> The proviral DNA is then incorporated into the host-cell genome using the HIV integrase enzyme.<sup>[14]</sup> HIV viral polyproteins are then expressed using hostcell machinery and the large Gag-Pol polyprotein is cleaved by the HIV protease enzyme. Finally, after the virion buds from the host cell, processing of the HIV viral capsid protein to allow mature viral particles to be produced is accomplished in a process called maturation.[15] Study of HIV cell entry and integration has recently led to the approval of maraviroc, a CCR5 co-receptor antagonist, and raltegravir, an integrase inhibitor. This review discusses the clinical trial results that led to their approval, their role in clinical practice, and drugs for novel targets in clinical and preclinical development.

#### 1. CCR5 Co-Receptor Antagonists

HIV-1 enters and infects cells by interacting with the CD4 receptor and a co-receptor, either CCR5 or CXCR4 (sometimes also simply referred to as R5 and X4). The HIV-1 gp120 envelope protein binds either CCR5 or CXCR4, thereby infecting CCR5- or CXCR4-expressing cells. HIV-1 may also be dual-tropic (i.e. is able to use either receptor). Some patients have mixed HIV populations of CCR5- and CXCR4-tropic viruses. Because current tropism assays cannot distinguish between dual and mixed HIV populations, they are referred to as dual/mixed (D/M) tropic. CCR5-tropic virus predominates in less advanced disease, whereas CXCR4 virus emerges later and is associated with more rapid disease progression (table I).[16,17]

Phenotypic  $(Trofile^{TM})$ and genotypic (SensiTrop<sup>TM</sup>) tropism assays that detect CXCR4and/or CCR5-tropic virus are available to identify patients who are likely to benefit from CCR5 co-receptor antagonists. The original Trofile™ assay detected CXCR4 virus with 100% accuracy when CXCR4 comprised ≥10% of the viral population. Successful detection dropped when viral load was <1000 copies/mL. SensiTrop<sup>™</sup> was less sensitive than Trofile<sup>TM</sup> in detecting D/M or CXCR4 virus. Of 100 samples classified by Trofile<sup>TM</sup>, SensiTrop<sup>TM</sup> identified a higher proportion as CCR5 (69 vs 52) and a lower proportion as D/M (20 vs 31).[25] An enhanced Trofile<sup>TM</sup> assay has increased sensitivity for CXCR4 by 10- to 100fold and identified more patients with D/M virus compared with the original Trofile<sup>TM</sup> assay. [26,27] This assay is able to detect CXCR4 tropic virus 100% of the time when present at 0.3% of the total viral population.<sup>[28]</sup> An enhanced genotypic assay (SensiTrop<sup>TM</sup> II) has been developed, which demonstrated 87% concordance with the original Trofile™ assay and was also able to identify D/M populations in the clade C HIV subtype with significantly more sensitivity than the standard Trofile™ assay.<sup>[29]</sup> A comparison between the enhanced Trofile™ assay and SensiTrop™ II has not been performed. Finally, genotypic clonal analysis of HIV proviral DNA may be useful in determining tropism in patients with undetectable plasma

Table I. Prevalence of co-receptor tropism<sup>[18]</sup>

Study cohort	Co-receptor us	sage (%)		Reference	
	CCR5	CCR5/CXCR4	CXCR4		
Treatment-experienced patients					
Chelsea and Westminster cohort (n = 161)	78	22	<1	19	
Demarest et al. (n = 117)	67	28	5	20	
MOTIVATE 1 and 2 (n = 2560)	56	41.4	2.6	21	
TORO 1 and 2 (n=612)	50	48	2	22	
ACTG 5211 (n=391)	50	46	4	23	
Treatment-naive patients					
MERIT cohort (n = 1428)	85	14.7	0.3	21	
Homer cohort (n=979)	82	18	<1	24	
Chelsea and Westminster cohort (n = 402)	81	19	<1	19	
Demarest et al. (n=299)	88	12	0	20	

**ACTG** = AIDS Clinical Trials Group; **MERIT** = Maraviroc vs Efavirenz Regimens as Initial Therapy; **MOTIVATE** = Maraviroc Plus Optimized Background Therapy in Viremic, ART-Experienced Patients; **TORO** = T-20 (enfuvirtide) versus Optimised Regimen Only.

viral loads.<sup>[30]</sup> In CCR5 co-receptor antagonist-naive patients, 7% had baseline HIV envelope gene V3 loop mutations associated with resistance to marayiroc.<sup>[31]</sup>

Maraviroc is the first approved CCR5 coreceptor antagonist. The trials known as MOTI-VATE (Maraviroc Plus Optimized Background Therapy in Viremic, ART-Experienced Patients) 1 and 2 (table II) compared different doses of maraviroc with placebo in the presence of an optimized background regimen (OBR) in patients with CCR5-tropic virus, previous ART experience and antiretroviral resistance.[46,47] Mean CD4+ cell count increases and number of patients who achieved undetectable viral loads (<50 copies/mL) at 48 weeks were significantly greater with maraviroc compared with placebo (table II).[32,48] Virological response was better in those with a baseline viral load <100 000 copies/ mL. The most common maraviroc adverse events were diarrhoea, nausea, fatigue and headache. There were similar frequencies of serious adverse events, toxicity-driven discontinuations, laboratory abnormalities, AIDS-defining infections, and AIDS- or non-AIDS-defining malignancies among maraviroc and placebo arms; however, there were higher rates of grade 3 or 4 AST and ALT elevations in hepatitis C virus (HCV) co-infected patients in the maraviroc arm. [49] Virological failure was significantly associated

with a switch from CCR5 to D/M or CXCR4 virus (table III). Discontinuation of maraviroc resulted in reversion of D/M or CXCR4 to CCR5 virus in the majority of those who experienced treatment failure.[50] Additional virological failures were explained by changes in tropism (using the original Trofile™ assay) seen between screening and study entry (table III). V3 loop envelope gene mutations that confer genotypic resistance to maraviroc were found in all patients experiencing treatment failure with phenotypic resistance to maraviroc, whether with CCR5- or CXCR4-tropic virus.[11] Another recent analysis using the MOTIVATE 1 and 2 data found that the activity of the OBR was a stronger predictor of virological success in the maraviroc arm than baseline CCR5 tropism, reinforcing the concept that it is important to maximize the number of active drugs in ART regimens.<sup>[52]</sup>

The MERIT (Maraviroc vs Efavirenz Regimens as Initial Therapy) study (table II) compared maraviroc and efavirenz, in combination with zidovudine/lamivudine, in ART-naive patients. [33] It was a randomized, double-blind, phase III trial of patients with CCR5-tropic HIV-1 and a plasma HIV-1 RNA level ≥2000 copies/mL. The study failed to show maraviroc noninferiority using a noninferiority margin of −10% of maraviroc to efavirenz for the endpoint of viral load <50 copies/mL; however,

Table II. Summary of virological and immunological responses to new antiretroviral agents in representative studies

Study name	Dosage	No. of patients	Mean baseline VL (log <sub>10</sub> copies/mL)	Mean baseline CD4+cell count (cells/μL)	Mean VL response (log <sub>10</sub> copies/mL)	VL <50 copies/ mL at study endpoint (%)	Mean CD4+ cell count response (cells/μL)	Timepoint of results (wk)	References
MVC (maraviro	c)								
MOTIVATE 1	MVC 150, 300 mg bid + OBR	235	4.86	150	-1.82	46.0	+122	48	32
	MVC 150, 300 mg od + OBR	232	4.85	168	-1.66	42.0	+113		
	PL+OBR	118	4.84	160	-0.80	16.0	+54		
MOTIVATE 2	MVC 150, 300 mg bid + OBR	191	4.84	182	-1.87	45.0	+128	48	32
	MVC 150, 300 mg od+OBR	182	4.87	173	-1.72	45.0	+122		
	PL+OBR	91	4.89	174	-0.76	18.0	+69		
MERIT	MVC 300 mg bid + AZT/3TC	360	4.86	241 (median)	NR	65.3	+170	48	33
	EFV 600 mg od + AZT/3TC	361	4.88	254 (median)	NR	69.3	+144		
MERIT-ES	MVC 300 mg bid + AZT/3TC	361	4.88	236 (median)	NR	68.5	+174	48	34
	EFV 600 mg od + AZT/3TC	303	4.85	254 (median)	NR	68.3	+144		
VCV (viewiyiya a	A								
VCV (vicrivirod ACTG 5211	VCV 5 mg od + OBR	30	4.56 <sup>a</sup>	146ª	b	NR	NR	48	35,36
	VCV 10 mg od + OBR	30	4.56 <sup>a</sup>	146ª	-1.92	37	+130		35,36
	VCV 15 mg od + OBR	30	4.56 <sup>a</sup>	146ª	-1.44	27	+96		
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Table II. Contd

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Study name	Dosage	No. of patients	Mean baseline VL (log <sub>10</sub> copies/mL)	Mean baseline CD4+cell count (cells/μL)	Mean VL response (log <sub>10</sub> copies/mL)	VL <50 copies/ mL at study endpoint (%)	Mean CD4+ cell count response (cells/μL)	Timepoint of results (wk)	References
	PL+OBT	28	4.56 <sup>a</sup>	146ª	NR	11	NR		
VICTOR-E1	VCV 20 mg od + OBR with boosted PI	116°	4.5 <sup>d</sup>	200 <sup>d</sup>	-2.04	58	+100	48	37
	VCV 30 mg od + OBR with PI	116 <sup>c</sup>	4.5 <sup>d</sup>	200 <sup>d</sup>	-2.04	64	+94		
	OBR	116 <sup>c</sup>	4.5 <sup>d</sup>	200 <sup>d</sup>	-0.96	26	+56		
RAL (raltegray	rir)								
Protocol 004	RAL 100 mg bid+TDF/FTC	39	4.6–4.8 <sup>e</sup>	271–314 <sup>e</sup>	At least –2.2	85	+144-221 <sup>9</sup>	48	38,39
	RAL 200 mg bid+TDF/FTC	40	4.6–4.8 <sup>e</sup>	271–314 <sup>e</sup>	At least –2.2	83	+144–221 <sup>9</sup>		
	RAL 300 mg bid+TDF/FTC	41	4.6–4.8 <sup>e</sup>	271–314 <sup>e</sup>	At least –2.2	88	+144–221 <sup>9</sup>		
	RAL 400 mg bid + TDF/FTC	40	4.6–4.8 <sup>e</sup>	271–314 <sup>e</sup>	At least -2.2	88	+144–221 <sup>9</sup>		
	EFV 600 mg od + TDF/FTC	38	4.6–4.8 <sup>e</sup>	271 to 314 <sup>e</sup>	At least -2.2	87	+144-221 <sup>g</sup>		
STARTMRK	RAL 100 mg bid + TDF/FTC	281	5.01	219	NR	86	+189	48	40
	EFV 600 mg qhs + TDF/FTC	282	5.02	217	NR	82	+163		
Protocol 005	RAL 100 mg bid + OBR	43	4.6	186	-1.80	65.1	+62.9	24	41
	RAL 400 mg bid+OBR	45	4.8	115	-1.87	55.6	+112.8		
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Table II. Contd

Study name	Dosage	No. of patients	Mean baseline VL (log <sub>10</sub> copies/mL)	Mean baseline CD4+cell count (cells/μL)	Mean VL response (log <sub>10</sub> copies/mL)	VL <50 copies/ mL at study endpoint (%)	Mean CD4+ cell count response (cells/μL)	Timepoint of results (wk)	References
	RAL 600 mg bid + OBR	45	4.7	161	-1.84	66.7	+94.1		
	PL+OBR	45	4.7	188	-0.35	13.3	+5.4		
BENCHMRK 1	RAL 400 mg bid + OBR	232	4.6	140	-1.7 <sup>f</sup>	65	+120	48	42,43
	PL+OBR	118	4.5	105	-0.8 <sup>f</sup>	31	+49		
BENCHMRK 2	RAL 400 mg bid+OBR	75	4.7	102	-1.7 <sup>f</sup>	60	+98	48	42,44
	PL+OBR	41	4.7	132	-0.8 <sup>f</sup>	35	+40		
EVG (elvitegra	vir)								
GS183-0105	EVG 50 mg od	71	4.59 <sup>g</sup>	185 <sup>9</sup>	-1.5	38	+52	24	45
	EVG 125 mg od	73	4.59 <sup>g</sup>	185 <sup>9</sup>	-1.7	40	+61		
	PI	63	4.59 <sup>g</sup>	185 <sup>g</sup>	-1.2	30	+28		

a For all 118 subjects.

ACTG = AIDS Clinical Trials Group; AZT/3TC = zidovudine/lamivudine; BENCHMRK = Blocking Integrase in Treatment-Experienced Patients With a Novel Compound Against HIV; bid = twice daily; EFV = efavirenz; FTC = emtricitabine; MERIT = Maraviroc vs Efavirenz Regimens as Initial Therapy; MOTIVATE = Maraviroc Plus Optimized Background Therapy in Viremic, ART-Experienced Patients; NR = not reported; OBR = optimized background regimen; od = once daily; PI = ritonavir-boosted protease inhibitor; PL = placebo; qhs = at bedtime; STARTMRK = Safety and Efficacy of Raltegravir-Based versus Efavirenz-Based Combination Therapy in Treatment-Naive HIV-1 Infected Patients; TDF = tenofovir; VICTOR-E1 = Vicriviroc in Combination Treatment with an Optimized ART Regimen in HIV-Infected, Treatment-Experienced Subjects; VL = viral load; + indicates increase; - indicates decrease.

b This arm ended early because of virological failure.

c For all three groups.

d For all 116 subjects.

e For all 198 subjects.

f BENCHMRK 1 and 2 combined.

g For all patients.

noninferiority was demonstrated for suppression to <400 copies/mL. When analysed by baseline viral load of less than or greater than 100000 copies/mL, 71.6% and 69.6%, and 66.6% and 59.6% in the efavirenz and maraviroc arms, respectively, reached a viral load of <50 copies/mL. CD4+ cell count increases were greater in the maraviroc arm (table II). As in the MOTIVATE studies, 25 patients (3.4%) demonstrated a switch from CCR5-tropic virus at screening to D/M virus at study entry and treatment failure was seen in patients with CCR5-, D/M or CXCR4tropic virus<sup>[51]</sup> (table III). In the group with D/M virus at study entry, only 54.6% and 7.1% of patients in the efavirenz and maraviroc arms, respectively, achieved a viral load of <50 copies/mL. Of those in whom treatment failed, nucleoside reverse transcriptase inhibitor (NRTI) mutations were more likely to develop with D/M or CXCR4 virus than with CCR5.<sup>[51]</sup> In the MERIT-ES (enhanced sensitivity) reanalysis of 48-week efficacy and safety data, MERIT study samples were re-tested with the enhanced Trofile<sup>TM</sup> assay, which was able to detect D/M virus in 55% of the patients who had CCR5-tropic virus at screening but who developed D/M virus during the study. [34] Virological responses with maraviroc and efavirenz were similar: 68.5% and 68.3% of patients, respectively, achieved an HIV-1 RNA <50 copies/mL at week 48. More maraviroc treatment discontinuations resulted from lack of efficacy (9.3% vs 4.0%), whereas more efavirenz discontinuations were for adverse events including CNS adverse effects, rash and aminotransferase elevations (14.2% vs 4.2%).

Vicriviroc is an oral CCR5 co-receptor antagonist with the following characteristics: (i) plasma concentrations are markedly increased by cytochrome P450 (CYP) 3A4 inhibitors (e.g. ritonavir); (ii) a plasma half-life >24 hours; (iii) no food effects; and (iv) durable antiretroviral activity in CCR5-tropic ART-experienced subjects. [35,53,54] The ACTG (AIDS Clinical Trials Group) 5211 study evaluated vicriviroc in ART-experienced adults (table II). [35,36] Following optimization of background ART, vicriviroc (10 or 15 mg once daily) with ritonavir demonstrated sustained antiretroviral activity over

48 weeks in only a minority of patients, although median CD4+ cell count increased by 130 and 96 cells/mm³ for the 10 and 15 mg groups, respectively. Findings similar to maraviroc of tropism switching and tropism type with virological failure were seen with vicriviroc, and crossresistance can be expected if treatment failure is due to a change in tropism to CXCR4 (table III). The time to virological failure was quicker in those with D/M than CCR5 virus. [36] ACTG 5211 results were re-analysed with the enhanced Trofile™ assay, and demonstrated greater reductions in HIV-1 viral load at day 14 and week 24 among patients with CCR5- versus D/M-tropic virus at study entry. [55]

The VICTOR-E1 (Vicriviroc in Combination with an Optimized ART Regimen in HIV-Infected Treatment-Experienced Subjects) study compared two higher doses of vicriviroc (20 or 30 mg once daily) with placebo all plus OBR (at least three antiretroviral agents, including a ritonavir-boosted protease inhibitor [PI]) in ARTexperienced patients for 48 weeks (table II).[37] HIV-1 viral load was significantly reduced and CD4+ cell counts significantly increased in both vicriviroc arms compared with placebo. Viral tropism changes are described in table III.[56] Once-daily vicriviroc 30 mg was selected for ongoing trials in ART-experienced subjects. Adverse event data from 205 patients in the ACTG 5211 and VICTOR-E1 studies who were followed-up for 96 weeks found few treatment-emergent adverse events including AIDS-associated opportunistic infections and malignancies. A few patients (17 and 14) developed elevated AST and ALT, respectively, but these elevations were judged not to be related to vicriviroc.<sup>[57]</sup>

#### 1.1 Other CCR5 Co-Receptor Antagonists

Aplaviroc (GSK-873,140) was another CCR5 co-receptor antagonist that showed good antiretroviral activity but clinical development was discontinued because of hepatotoxicity. [58-60] Other small molecule CCR5 co-receptor antagonists in development have included INCB-9471, [61] SCH532706 in combination with low-dose ritonavir, [62] and PF-232798, which has potent

Table III. Tropism and outcome in clinical trials of R5 inhibitors

Trial name (drud) No. of pts with	No. of pts with	No. of pts with tropism	No. of pts with tropism Virological outcome of D/M at	Tropism at CCR5	Time to treatment failure	Reference
ò	pretreatment tropism shift (%)		study entry	inhibitor failure		
MOTIVATE 1 and 83 of 1042 (8) 2 (MVC)	83 of 1042 (8)	No data	Inferior virological outcome	2 of 3 pts had detectable CXCR4 virus at failure	30 days earlier if D/M or CXCR4	20
MERIT (MVC)	25 of 740 (3.4)	No data	Undetectable VL in 6 of 11 EFV pts and 1 of 14 MVC pts	32 R5 at baseline, 10 of 32 had D/M or CXCR4 at failure.	No data	51
ACTG 5211 (VCV)	12 of 118 (10)	4 of 30 (VCV 10 mg) 3 of 30 (VCV 15 mg) 3 of 28 (placebo) [2 after switch to VCV]	No data	9 of 26 (35%) had CXCR4 virus at failure	Shorter in D/M compared with CCR5	35
VICTOR-E1 (VCV)	6 of 114 (5)	9 of 39 [23% CXCR4] (VCV 30 mg) 7 of 40 [10% CXCR4] (VCV 20 mg)	Not necessarily associated with 13 of 108 (12%) virological failure	13 of 108 (12%)	No data	37

ACTG=AIDS Clinical Trial Group, D/M=dual/mixed tropism; EFV=efavirenz; MERIT=Maraviroc vs Efavirenz Regimens as Initial Therapy; MOTIVATE=Maraviroc Plus Optimized Background Therapy in Viremic, ART-Experienced Patients; MVC=maraviroc; pts=patients; VCV=vicriviroc; VICTOR-E1=Vicriviroc in Combination Treatment with Optimized Regimen in Experienced Subjects; VL = viral load. activity against CCR5-tropic, maravirocresistant HIV-1.[63] Monoclonal antibodies that block the binding of HIV to CCR5 in clinical HGS004<sup>[58,64]</sup> trials included have PRO140.[41,64-66] PRO140 was synergistic with maraviroc and vicriviroc, active against maraviroc- and vicriviroc-resistant strains in vitro, [67,68] was well tolerated<sup>[69]</sup> and, in a small number of HIV-infected patients with CCR5-tropic virus, showed a dose-dependent decrease in viral load.[41] One of 30 patients who received PRO140 had a CCR5 to D/M change, but analysis showed that the patient was likely to have had a pre-existing D/M virus rather than developing a mutation following treatment.<sup>[70]</sup> A randomized, placebo-controlled phase IIA trial compared PRO140 at 5 and 10 mg/kg single infusions, and found a mean 2 log<sub>10</sub>/mL decrease in viral load in the 10 mg/kg group that was durable for 4 weeks post-injection.<sup>[71]</sup> Dose administration every 3 weeks may be possible based on these results. Finally, aprepitant, a neurokinin-1 receptor antagonist and approved antiemetic. which downregulates CCR5 expression on macrophages in vitro, [72] is currently under investigation in HIV infection.[73]

## 1.2 Clinical Use of CCR5 Co-Receptor Antagonists

Maraviroc is approved for use in antiretroviral treatment-experienced patients with CCR5-tropic HIV-1, detectable viral loads, and resistance to multiple antiretroviral drugs based on results from the MOTIVATE 1 and 2 studies. Maraviroc is useful in the subset of patients who have some ART experience. However, 50% of patients with low CD4+ cell counts are likely to have D/M- or, less commonly, CXCR4-tropic virus; therefore, tropism testing is required to determine the utility of maraviroc in this population.<sup>[23]</sup> Furthermore, 5–10% of patients who were found to have CCR5 virus on the original TrofileTM assay will have minority CXCR4 populations capable of causing maraviroc failure. However, with the enhanced sensitivity Trofile™ assay, maraviroc will be used with more confidence because CXCR4 populations will be detected with greater certainty. Maraviroc should not be used in treatment-naive patients based on MERIT study results, where maraviroc did not meet criteria for noninferiority to efavirenz-based therapy. The MERIT-ES re-analysis of the MERIT data with the enhanced Trofile™ assay did find that maraviroc was noninferior to efavirenz when combined with zidovudine/lamivudine; however, maraviroc is not US FDA approved for use in ART-naive patients, and more information on pairing of maraviroc with other antiretrovirals and longer-term outcomes would be helpful.

If a patient develops virological failure on a maraviroc-containing ART regimen, then standard resistance testing should be performed, and a tropison assay may be useful in determining the cause of treatment failure. Some virological failures on maraviroc regimens are secondary to CCR5 to D/M or CXCR4 tropism changes (57% and 31% of patients with virological failure in the MOTIVATE and MERIT trials).[51,74] However, a small study has shown that it is possible for maravoric treatment to fail in patients with CCR5-tropic but maraviroc-resistant virus secondary to V3 loop mutations.[11] Additional virological failure results from other antiretroviral resistance in the regimen and standard antiretroviral resistance testing should be performed. Tropism testing can be considered at the time of maraviroc-containing regimen failure to prevent inappropriate maraviroc discontinuation; however, no commercially available testing can rule out the presence of CCR5-tropic but phenotypic maraviroc-resistant HIV.

The recommended dose of maraviroc is as follows: (i) 150 mg orally twice daily when used in combination with CYP3A inhibitors, including PIs except the combination of tipranavir plus ritonavir; azoles such as itraconazole and ketoconazole, and clarithromycin; (ii) 300 mg orally twice daily when given in combination with ritonavir boosted tipranavir, NRTIs, enfuvirtide, nevirapine and other drugs without strong CYP3A effects; and (iii) 600 mg orally twice daily with CYP3A inducers, including efavirenz, etravirine and rifampin. Hypericum (St John's wort) should be avoided, whereas hormonal contraceptives can be safely used.<sup>[75]</sup> Because

maraviroc is primarily metabolized by the liver, the FDA recommends caution when giving maraviroc to patients with hepatic impairment, as maraviroc levels may increase.<sup>[76]</sup> Further caution is indicated in patients with ischaemic heart disease as patients receiving maraviroc had a cardiovascular event rate of 1.34 compared with none in patients receiving placebo.<sup>[76]</sup>

### 2. Integrase Inhibitors

Integrase inhibitors are a new class of antiretrovirals that inhibit HIV proviral DNA integration into the host genome. Integration is a multi-step process and currently available integrase inhibitors inhibit the strand transfer step.<sup>[77]</sup> Raltegravir is a strand transfer inhibitor and was recently approved for use in HIV treatment. It is active against HIV strains with reverse transcriptase (RT) and protease (PR) resistance mutations, different HIV-1 subtypes, and CCR5 or CXCR4 strains. It is metabolized primarily by glucuronidation (UGT1A1) and is not a potent inhibitor or inducer of CYP3A4. Elvitegravir is another potent integrase inhibitor that is also active against antiretroviral (NRTI, non-nucleoside reverse transcriptase inhibitors [NNRTI], PI) resistant strains, different HIV-1 subtypes and HIV-2. Elvitegravir also has additive activity in vitro with other antiretrovirals and is a moderate inducer of CYP3A4.<sup>[78]</sup>

The phase IIa Protocol 004 study evaluated multiple doses of raltegravir compared with efavirenz in treatment-naive patients, all with tenofovir and lamivudine (table II). There was no significant difference in viral load or CD4+ cell count results between the raltegravir and efavirenz arms. Combining all raltegravir doses, there were no dose-related toxicities, and total cholesterol, low-density lipoprotein cholesterol (LDL-C)and triglycerides were not increased by raltegravir. [38,39] The STARTMRK (Safety and Efficacy of Raltegravir-Based versus Efavirenz-Based Combination Therapy in Treatment-Naive HIV-1 Infected Patients) study evaluated raltegravir 400 mg twice a day versus efavirenz 600 mg at bedtime with tenofovir/emtricitabine in treatment-naive patients.[40] There were no significant

differences between raltegravir and efavirenz in the percentage of patients achieving an undetectable HIV viral load, although the increase in absolute CD4+ cell count with raltegravir was greater than with efavirenz. Of 12 patients in the raltegravir arm who did not respond to treatment and had >400 copies/mL of HIV, four had integrase resistance mutations and three did not have the integrase gene amplified. Three of eight patients who did not respond to treatment with efavirenz had efavirenz resistance mutations. Lipid profiles were less affected with raltegravir than with efavirenz; LDL-C increased 6 and 16 mg/dL (0.15 and 0.4 mmol/L) on average and triglycerides decreased 3 and increased 37 mg/dL (0.033 and 0.4 mmol/L) on average, respectively.

The Protocol 005 study compared multiple doses of raltegravir plus OBR versus placebo plus OBR in ART-experienced patients and found that raltegravir resulted in a significantly greater reduction in viral load than placebo after 24 weeks (table II).<sup>[79]</sup> The BENCHMRK (Blocking Integrase in Treatment-Experienced Patients with a Novel Compound Against HIV) 1 and 2 studies compared raltegravir plus OBR with placebo plus OBR in ART-experienced patients who had antiretroviral resistance (table II). After 48 weeks, patients in the raltegravir arms were significantly more likely to achieve an undetectable viral load compared with those in the placebo arm. The best virological result occurred in those patients who received raltegravir with both darunavir and enfuvirtide in their OBR (89%). Remarkably, 50% of those receiving raltegravir with no other active drugs in the OBR achieved an undetectable viral load compared with only 2% of patients receiving placebo plus OBR. Adverse events were similar between raltegravir and placebo, and no increased risk of malignancy with raltegravir was seen.[42] There was a trend towards decreased AIDS-defining events or death at 48 weeks in the raltegravir arm versus placebo arm relative risk (relative risk [RR] 0.50; 95% CI 0.25, 1.03).<sup>[80]</sup>

Elvitegravir was evaluated in patients who were ART naive or not receiving their current regimen, at doses of 50, 400 and 800 mg, all with ritonavir, and was shown to reduce viral load by

 $2.0 \log_{10}/\text{mL}$ . [81] The phase II study, GS-US-183-0105, evaluated elvitegravir plus ritonavir plus OBR compared with a comparator ritonavirboosted PI plus OBR in ART-experienced patients with at least one primary PI mutation (table II). Enfuvirtide, but not NNRTIs, were allowed in the OBR. PIs were not allowed in the elvitegravir arms. After 24 weeks, the mean change in HIV-1 viral load from baseline was 1.7 versus 1.4 log<sub>10</sub> copies/mL in the 125 mg elvitegravir and comparator PI arms, respectively. Further analysis found that elvitegravir without active antiretrovirals in the OBR reduced viral load by 0.7 versus 2.1 log<sub>10</sub> copies/mL in those with greater than one active NRTI or enfuvirtide in their OBR. [45] Elvitegravir is currently in phase III development.

The integrase gene is highly polymorphic, although, to date, no polymorphisms that have been found are associated with phenotypic integrase inhibitor resistance (table IV). Some polymorphisms are more common in ARTexperienced and others in ART-naive patients. Despite integrase polymorphisms, integrase inhibitors are active against HIV-2.[82,87-89] Raltegravir resistance was first described in the Protocol 005 study and later confirmed in other raltegravir studies. Two different pathways of integrase resistance mutations (N155 or Q148) have been described. Additional integrase mutations were frequently observed with both pathways and Q148 mutants were more common than N155. The most common mutation pattern was Q148H/G140S, which resulted in significantly higher raltegravir fold change in the concentration that produces 50% inhibition (IC<sub>50</sub>) relative to reference than Q148H alone. [43,44,83] At the time of raltegravir failure in BENCHMRK, 53% of patients had a Q148 mutation, 18% had N155 and 10% had a mixture. [90] Q148H in combination with secondary mutations was not associated with a decrease in replication capacity, while N155H was, which helps to explain why Q148H is the more common resistance pathway. [90] Virological failure of elvitegravir regimens also resulted from development of N155H, Q148R/H/K and, additionally, E92Q mutations. Multiple integrase mutations

Table IV. Integrase gene polymorphisms and resistance mutations<sup>[42,82-86]</sup>

Clinical situation	Polymorphism/mutation
Mutations and combinations associated with virological failure	With raltegravir: Q148R/H/K, Q148/G140S, N155H With elvitegravir: Q148R/H/K, N155H, E92Q
Additional mutations and combinations associated with phenotypic failure and increased fold change <i>in vitro</i>	With raltegravir: E92Q, T66l/E92Q, E92Q/N155H, G140S/Q148H, E138K/S147G/Q148R, Q148K/E138A/G140A With elvitegravir: T66l/E92Q, E92Q/N155H, G140S/Q148H, E138K/S147G/Q148R
Polymorphisms and prevalence in antiretroviral-naive patients	80%: V201I 46%: V72I >5%: L74M/I, T97A, T125A/V, V151I, M154I, K156N, V165I, I203M, T206S, S230N
Polymorphisms in antiretroviral-treated patients	M154I, V165I, M185L

(table IV) significantly increased the fold change in  $IC_{50}$  relative to reference, resulting in high level phenotypic resistance to both raltegravir and elvitegravir.<sup>[84]</sup>

Two patients in whom an elvitegravir plus ritonavir regimen failed were switched to a raltegravir-containing regimen, with no virological response after switching. After elvitegravir plus ritonavir virological failure, one patient was found to have a N155H mutation, and the other a Q148H mutation in addition to other INI mutations. One patient achieved an undetectable viral load (<50 copies/mL) after addition of darunavir plus ritonavir to raltegravir, whereas the second patient did not,[91] indicating that cross resistance occurs and virological response is unlikely after failure with one integrase inhibitor. Raltegravir-resistant HIV was recently tested against elvitegravir and found to display even greater resistance to elvitegravir, indicating cross resistance.<sup>[85]</sup> Two second-generation integrase inhibitors being developed by Merck (MD-0536 and MK-2048) demonstrated much less resistance to these HIV isolates than raltegravir or elvitegravir, and plans are to pursue development of these drugs for treatment of raltegravir- and elvitegravir-resistant HIV-1.

A phenotypic integrase assay (PhenoSense<sup>®</sup> Integrase Assay) that measures the susceptibility of HIV-1 to integrase inhibitors has been developed.<sup>[92]</sup> The biological cut-off for raltegravir

fold change in  $IC_{50}$  in this assay was established as 1.5.

## 2.1 Clinical Use of Integrase Inhibitors

The BENCHMRK and other trials led to the approval of raltegravir for treatment-experienced HIV-1 infected adults with detectable viral loads and resistance to multiple antiretroviral drugs. Integrase inhibitor-naive patients can be expected to respond to raltegravir; whereas, in the BENCHMRK trials, fewer than 3% of patients receiving raltegravir did not achieve a viral load of ≤400 copies/mL. [42] No clinical trials have addressed which of the antiretroviral agents work best in combination with raltegravir, although in the BENCHMRK studies, the best virological outcomes were seen with patients receiving raltegravir, enfuvirtide and ritonavir boosted darunavir. Thus, the more active the antiretroviral drugs in a regimen, the better the outcome. Raltegravir has a neutral lipid profile and may play a role in patients with poor lipid profiles or in whom lipid-raising regimens should be avoided.[38] Phenotypic assays for measuring integase susceptibility are currently available and genotypic assays are available in some settings. Raltegravir-containing regimens may fail in a patient because of integrase resistance (table IV), resistance to other antiretroviral agents, or reasons such as poor ART adherence or poor

bioavailability. Because there is cross resistance between raltegravir and elvitegravir (table IV), switching agents within the integrase inhibitor class if resistance develops (once elvitegravir becomes available) is not recommended.<sup>[85]</sup> Studies indicate that raltegravir is likely to be an option for first-line ART.<sup>[40]</sup> Current dose administration of raltegravir is 400 mg twice daily without dosage adjustment when combined with other ART.

## 3. Other New HIV Agents in Development

#### 3.1 Attachment Inhibitors

Nonspecific attachment inhibitors, which block binding of HIV to CD4 cells in the initial step of HIV cell entry, such as PC-515 (Carraguard<sup>TM</sup>), cyanovirin-N and molecular umbrella compounds, are being developed as topical microbicides.[93-97] Specific attachment inhibitors include cyclotriazadisulfonamide (CADA), which downregulates expression of CD4.[98-100] and naphthalene sulfonate, which disrupts binding of gp120 to CD4.[101,102] A naphthalene sulfate polymer (PRO 2000)[103,104] is being tested as a vaginal gel.[105] Ibalizumab (TNX-355) is an anti-CD4 monoclonal antibody, [106,107] which has synergy with enfuvirtide, [108] and is in phase II clinical trials. [109]

#### 3.2 HIV Glycoprotein 120 Agents

Agents targeting gp120 are targeting the first step in HIV cell entry by preventing the interaction of CD4 with gp120. SPL7013 binds HIV gp120<sup>[102]</sup> and is being studied as a microbicide. [110,111] PRO 542 is a CD4 IgG2 that binds to HIV gp120 and was in a phase II open-label study recently completed but is no longer in development. [112] In previous studies, PRO 542 reduced viral load in adults with ART failure and in antiretroviral-naive children. [113-115] Several small molecule CD4 attachment inhibitors have been explored [116-119] and similar compounds are in development. [116]

#### 3.3 CCR5 Co-Receptor Agonists

As discussed in section 1, targeting the chemokine co-receptors for HIV entry, in particular CCR5, has been a successful strategy. Other ways to manipulate the CCR5 receptor are being explored. ESN-196 is a small-molecule CCR5 coreceptor agonist that causes internalization of the receptor and prevents HIV-1 infection in vitro.[120] Regulated on Activation, Normal T cell Expressed and Secreted (RANTES), the endogenous CCR5 ligand, [121] is being developed as a topical microbicide, with potent HIV-1 suppressive activity against maraviroc- and vicriviroc-resistant HIV-1 strains in vitro.[68,122] An anti-CD40 monoclonal IgG1 that induces RANTES and macrophage inflammatory protein- $1\alpha$  expression is also being developed. [123]

#### 3.4 CXCR4 Co-Receptor Antagonists

The change in HIV-1 tropism to CXCR4 while receiving CCR5 inhibitor therapy has prompted the search for CXCR4 co-receptor antagonists. Several CXCR4 co-receptor antagonists have reached clinical trials, but none have progressed beyond phase II because of problems with toxicity or lack of virological response. [124-126] POL3026 was found to prevent HIV replication and emergence of CXCR4-tropic HIV *in vitro*. [127-129] POL3026 may have the potential for use in combination with CCR5 inhibitors to prevent the on-treatment selection of CXCR4 populations.

#### 3.5 Fusion Inhibitors

Enfuvirtide (T20) is the first-in-class fusion inhibitor to be FDA-approved for clinical use, and it works by binding gp41 and thus preventing fusion of the HIV viral and human cellular membranes. Several fusion inhibitors are in preclinical development, while others are in phase I trials. [130,131] A peptide derived from the gp41 HR2 region, TRI-1144 (TR-029114), is a second-generation fusion inhibitor that has undergone preclinical development, [132] and has now completed a phase I, single-dose, placebo-controlled clinical trial (TRI 1144-101) in 24 healthy volunteers

with results expected later this year. Preliminary results have been announced indicating that TRI-1144 was well tolerated, but, to date, no data has been published. The pharmacokinetics suggest that it can be administered less frequently than enfuvirtide, is stable in solution and therefore does not have to be reconstituted, and has a higher genetic barrier to resistance. Another gp41 inhibitor developed in China, sifuvirtide (FS0101, Fusogen), was able to block fusion of enfuvirtide-resistant HIV strains and was well tolerated in a phase Ia clinical study of healthy volunteers. [134]

#### 3.6 Maturation Inhibitors

Bevirimat (PA103001-4, PA-457) is a gag maturation inhibitor that inhibits the final step in the processing of the HIV-1 gag polyprotein by binding to the HIV-1 capsid precursor CA-SP1 (p25) cleavage site. This prevents conversion to the capsid protein p25, which is required for the formation of infectious viral particles. Both in vitro and animals studies indicate that bevirimat impairs SP1-capsid cleavage, delays replication of PI-resistant HIV and reduces viral load in SCID-hu Thy/Liv mice.[135-138] Bevirimat is well tolerated and it significantly decreases viral load. [139,140] None of the mutations associated with bevirimat in vitro were found in vivo.[136] Bevirimat has activity against ART-resistant and wild-type HIV, and has shown synergy with antiretrovirals from all classes.[141] Results from a phase II clinical trial of bevirimat with OBR in patients with antiretroviral resistance and in whom current ART regimens had failed were recently reported.[142] Bevirimat reduced HIV viral load by a mean of 1.6 log<sub>10</sub>/mL in patients who achieved trough levels of ≥20 µg/mL and who did not have any of the key baseline gag polymorphisms at Q369, V370 or T371. All responders, defined as those who had a drop in viral load >0.5 log<sub>10</sub>/mL, had trough levels ≥20 µg/mL, while 60% of nonresponders had troughs levels <20 µg/mL.[143] Trough levels of 20 µg/mL were achieved with a liquid formulation of bevirimat at doses of 250-400 mg/day. Viral load reduction was significantly lower in patients with gag polymorphisms. Adverse advents were not more common in the bevirimat than the placebo arm. Phase III studies are planned.

#### 4. Immunotherapy

Exogenous interleukin (IL)-2 promotes CD4+ T-cell differentiation and proliferation, and has the potential to prevent progression to AIDS. It has increased CD4+ cell counts in clinical trials. but a clear effect on clinical outcome has not been seen.[144-146] In the UK-Vanguard study, IL-2 significantly increased CD4+ cell counts at 24 and 48 weeks, but there was no difference in CD4+ cell counts beyond 48 weeks, and no difference in time to initiation of ART.[144] The ARNS (French Agency for HIV Research) 199 trial found that patients who were given IL-2 were less likely to require ART (p<0.0001). [145] Patients with discordant responses to ART who were given IL-2 therapy and subsequently demonstrated a CD4+ cell count increase actually had a significantly decreased risk of death or AIDS-defining event.<sup>[146]</sup> IL-2 is being studied further in ESPRIT (Evaluation of Subcutaneous Proleukin in a Randomized International Trial), an international trial of subcutaneous recombinant IL-2 for patients on combination ART with CD4+ cell counts >300/mm<sup>3</sup> being followed-up for 5 years for disease progression and death.[147] Results are expected at the end of 2008.[148] Finally, in an attempt to improve T-cell memory, maturation and survival, exogenous IL-7 is also under investigation.[149]

## Gene Therapy

Current gene therapy agents include lexgenleucel-T (VRX496), an antisense HIV envelope gene that has been sequence transduced into CD4+ cells, [150-152] and OZ1 (Johnson & Johnson) an anti-HIV-1 ribozyme that has been transduced into CD34+ cells. [153] A phase I study of VRX496 in five patients found no adverse events and a sustained decrease in viral load was achieved in one of the patients. [154] Three phase II clinical trials are underway. [155-157] A phase II

trial of OZ1 has been completed, but no further development is currently underway.<sup>[153]</sup>

Other preclinical genetic work has found that HIV is suppressed by small interfering RNAs targeting the RNA helicase DDX3, which exports RNA from the nucleus to the cytoplasm. [158,159] Zinc-finger DNA-binding protein nucleases against CCR5 DNA sequences were found to protect against CCR5-tropic HIV-1 infection in mice. [64,160,161] Finally, fusion inhibitors are being expressed constitutively from viral vectors. [160]

#### 6. Conclusion

Two new classes of antiretroviral drugs (CCR5 co-receptor antagonists and integrase inhibitors), with novel targets, have come into clinical use within the last year and significantly increase drug treatment options for ARTexperienced patients. Maraviroc and raltegravir are now approved, and additional agents within these classes are in late clinical development. Maturation inhibitors may be the next class poised to enter clinical usage. Antibody therapy against steps in HIV cell entry is being productively explored, as well as small molecule therapy against new targets of entry, such as CXCR4 receptors and gp120. Gene therapy against many steps in the HIV lifecycle has demonstrated good preclinical results, but clinical effectiveness has vet to be demonstrated. Finally, innovative immunotherapies are being explored, although, to date, results are inconclusive.

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