



This Medication Guide has been approved by the U.S. Food and Drug Administration.

Medication Guide available at  
<http://camberpharma.com/medication-guides>



Manufactured for:  
Camber Pharmaceuticals, Inc.,  
Piscataway, NJ 08854

Manufactured by:  
**HETERO™**  
Hetero Labs Limited  
Jeddahmedta, Hyderabad - 500 055,  
India

Issued: 01/2022

## 12.3 Pharmacokinetics

Table 11. Mean Maraviroc Pharmacokinetic Parameters in Adults

Patient Population	Maraviroc Dose	n	AUC <sub>0-24</sub> (ng·h/mL)	C <sub>max</sub> (ng/mL)	C <sub>min</sub> (ng/mL)
Healthy volunteers (Phase 1a)	300 mg twice daily	64	2,500	888	42.1
Asymptomatic HIV subjects (Phase 2a)	300 mg twice daily	8	2,560	818	33.8
Treatment-experienced HIV subjects (Phase 3)	300 mg twice daily	94	1,513	268	37.2
1 = CYP2A1 inhibitor	150 mg twice daily	375	2,463	332	101
Treatment-naïve HIV subjects (Phase 2b/3)	300 mg twice daily	344	1,866	287	60

The estimated exposure is lower compared with other trials possibly due to sparse sampling, food effect, compliance, and concomitant medications. Absorption Peak maraviroc plasma concentrations are attained 0.5 to 4 hours following single oral doses of 1 to 1,200 mg administered to uninfected volunteers. The pharmacokinetics of oral maraviroc are not dose proportional over the dose range. The absolute bioavailability of 100 mg dose is 23% and is predicted to be 33% at 300 mg. Maraviroc is a substrate for the efflux transporter P-gp. Effect of Food on Oral Absorption: Co-administration of a 300-mg tablet with a high-fat breakfast reduced maraviroc C<sub>max</sub> and AUC by 33% and co-administration of 75 mg of oral solution with a high-fat breakfast reduced maraviroc AUC by 73% in healthy adult volunteers. Studies with the tablet formulation demonstrated no effect of food on the pharmacokinetics of maraviroc.

There were no food restrictions in the adult trials using the tablet formulation or in the pediatric trial using both tablet and oral solution formulations that demonstrated the efficacy/safety/activity and safety of maraviroc. (See Clinical Studies 14.1, 14.2).

Distribution Maraviroc is bound (approximately 76%) to human plasma proteins, and shows moderate affinity for albumin and alpha-1 acid glycoprotein. The volume of distribution of maraviroc is approximately 104L.

Elimination Trials in humans and *in vitro* studies using human liver microsomes and expressed enzymes have demonstrated that maraviroc is principally metabolized by the cytochrome P450 system to metabolites that are essentially inactive against HIV-1. *In vitro* studies indicate that CYP2A1 is the major enzyme responsible for maraviroc metabolism. *In vitro* studies also indicate that polymorphic enzymes CYP2C8, CYP2C9, and CYP2C19 do not contribute significantly to the metabolism of maraviroc.

Maraviroc is the major circulating component (~42% drug-related radioactivity) following a single oral dose of 300 mg <sup>14</sup>C-maraviroc. The most significant circulating metabolite in humans is a secondary amine (~22% radioactivity) formed by N-dealkylation. This polar metabolite has no significant pharmacological activity. Other metabolites are products of *in-vitro* oxidation and are only minor components of plasma drug-related radioactivity.

Excretion: The terminal half-life of maraviroc following oral dosing to steady state in healthy subjects was 14 to 18 hours. A mass balance/excretion trial was conducted using a single 200-mg dose of <sup>14</sup>C-labeled maraviroc. Approximately 20% of the radioactivity was recovered in the urine and 76% was recovered in the feces over 180 hours. Maraviroc was the major component present in urine (mean of 8% dose) and feces (mean of 25% dose). The remainder was excreted as metabolites.

## Specific Populations

**Patients with Hepatic Impairment:** Maraviroc is primarily metabolized and eliminated by the liver. A trial compared the pharmacokinetics of a single 300-mg dose of maraviroc in subjects with mild Child-Pugh Class A, n = 8) and moderate (Child-Pugh Class B, n = 8) hepatic impairment with pharmacokinetics in healthy subjects (n = 8). The mean C<sub>max</sub> and AUC were 15% and 25% higher, respectively, for subjects with mild hepatic impairment, and 32% and 46%, respectively, for subjects with moderate hepatic impairment compared with subjects with normal hepatic function. These changes do not warrant a dose adjustment. Maraviroc concentrations are higher when maraviroc 150 mg is administered with a potent CYP2A1 inhibitor compared with following administration of 300 mg without a CYP2A1 inhibitor, so patients with moderate hepatic impairment who receive maraviroc 150 mg with a potent CYP2A1 inhibitor should be monitored closely for maraviroc-associated adverse events. The pharmacokinetics of maraviroc have been studied in subjects with severe hepatic impairment (See Warnings and Precautions 5.5).

**Patients with Renal Impairment:** A trial compared the pharmacokinetics of a single 300-mg dose of maraviroc in adult subjects with severe renal impairment (GFR less than 30 mL per minute, n = 8) and ESRD (n = 6) with healthy volunteers (n = 6). Geometric mean values for maraviroc C<sub>max</sub> and AUC were 2.4-fold and 3.2-fold higher, respectively, for subjects with severe renal impairment, and 1.7-fold and 2.0-fold higher, respectively, for subjects with ESRD as compared with subjects with normal renal function in the trial. Hemodialysis had a minimal effect on maraviroc clearance and exposure in subjects with ESRD. Exposures observed in subjects with severe renal impairment (ESRD) were within the range observed in previous 300-mg single-dose trials of maraviroc in healthy volunteers with normal renal function. However, maraviroc exposures in the subjects with normal renal function in the trial were 50% lower than those observed in previous trials. Based on the results of this trial, no dose adjustment is recommended for patients with renal impairment receiving maraviroc without a potent CYP2A1 inhibitor or inducer. However, if patients with severe renal impairment (ESRD) experience any symptoms of potential hepatotoxicity while taking maraviroc 300 mg twice daily, their dose should be reduced to 150 mg twice daily (See Dosage and Administration 2.3, Warnings and Precautions 5.3).

In addition, the trial compared the pharmacokinetics of multiple dose maraviroc in combination with zalcitabine/ritonavir 1,000/100 mg twice daily in potent CYP2A1 inhibitor combination for 7 days in subjects with mild renal impairment (GFR greater than 50 mL per minute and less than or equal to 80 mL per minute, n = 8) and moderate renal impairment (GFR greater than or equal to 30 and less than or equal to 50 mL per minute, n = 8) with healthy volunteers with normal renal function (n = 8). Subjects received 150 mg of maraviroc at different dose frequencies (healthy volunteers – every 12 hours, mild renal impairment – every 24 hours, moderate renal impairment – every 48 hours). Compared with healthy volunteers, for every 12 hours, geometric mean rates for maraviroc AUC<sub>0-24</sub>, C<sub>max</sub>, and C<sub>min</sub> were 50% higher, 20% higher, and 43% lower, respectively, for subjects with mild renal impairment (dosed every 24 hours). Geometric mean rates for maraviroc AUC<sub>0-24</sub>, C<sub>max</sub>, and C<sub>min</sub> were 10% higher, 28% lower, and 85% lower, respectively, for subjects with moderate renal impairment (dosed every 48 hours) compared with healthy volunteers (dosed every 12 hours). Based on the data from this trial, no adjustment in dose is recommended for patients with mild or moderate renal impairment (See Dosage and Administration 2.3).

**Pediatric Patients: Age 2 to Less Than 18 Years** The pharmacokinetics of maraviroc were evaluated in CCR5-tropic, HIV-1-infected, treatment-experienced pediatric subjects aged 2 to less than 18 years in the dose-finding study of Trial A4001031, doses were administered with food on intensive pharmacokinetic evaluation days and optimized to achieve an average concentration over the dosing interval (C<sub>avg</sub>) of greater than 100 ng/mL. Throughout the trial, no non-intensive pharmacokinetic evaluation days were taken with or without food. The initial dose of maraviroc was based on BSA and concomitant medication exposure (i.e., presence of CYP2A1 inhibitors and/or inducers). The conversion of dosing to a weight (kg) band-based in children provided comparable exposures with those observed in the trial at the corresponding BSA.

Maraviroc pharmacokinetic parameters in pediatric subjects aged 2 to less than 18 years receiving potent CYP2A1 inhibitors with or without a potent CYP2A1 inducer were similar to those observed in adults (Table 12).

Table 12. Maraviroc Pharmacokinetic Parameters in Treatment-Experienced Pediatric Patients Receiving Maraviroc with Potent CYP2A1 Inhibitors With or Without a Potent CYP2A1 Inducer					
Weight	Dose of Maraviroc	Maraviroc Pharmacokinetic Parameters* Geometric Mean			
		AUC <sub>0-24</sub> (ng·h/mL)	C <sub>max</sub> (ng/mL)	C <sub>min</sub> (ng/mL)	C <sub>min</sub> (ng/mL)
10 kg to < 20 kg	50 mg twice daily	2,349	108	324	73
20 kg to < 30 kg	75 mg twice daily	3,020	252	394	118
30 kg to < 40 kg	100 mg twice daily	3,229	269	430	128
> 40 kg	150 mg twice daily	4,044	337	563	152

\*Model-predicted steady-state pharmacokinetic parameters are presented.

Clinical pharmacokinetic data in pediatric patients aged 2 to less than 18 years receiving noninteracting concomitant medications are limited. Based on population pharmacokinetic modeling and simulation, the recommended dosing regimen of maraviroc for this population is predicted to result in similar maraviroc exposures when administered with or without food. The recommended dosing regimen of maraviroc for this population is predicted to result in similar maraviroc exposures when administered with or without food. The initial dose of maraviroc was based on BSA and concomitant medication exposure (i.e., presence of CYP2A1 inhibitors and/or inducers). The conversion of dosing to a weight (kg) band-based in children provided comparable exposures with those observed in the trial at the corresponding BSA.

**Race and Gender:** Based on population pharmacokinetics and 2 clinical CYP2A1 genotype analyses for race, no dosage adjustment is recommended based on race or gender.

## Drug Interactions

**Effect of Concomitant Drugs on the Pharmacokinetics of Maraviroc:** Maraviroc is a substrate of CYP2A1 and P-gp and hence its pharmacokinetics are likely to be modulated by inhibitors and inducers of these enzymatic transporters. The CYP2A1 inhibitors ketoconazole, itraconazole, voriconazole, danazol, and efavirenz are known to increase the C<sub>max</sub> and AUC of maraviroc (Table 14). While not studied, potent CYP2A1 and/or P-gp inducers rifampin, rifabutin, and efavirenz decreased the C<sub>max</sub> and AUC of maraviroc (Table 14). While not studied, potent CYP2A1 and/or P-gp inducers carbamazepine, phenytoin, and phenylethylamine decreased the C<sub>max</sub> and AUC of maraviroc (Table 14). In an *in vitro* study results, maraviroc is also a substrate of DATP1B1 and MRP2; its pharmacokinetics may be modulated by inhibitors of these transporters.

Tenofovir/ritonavir (not CYP2A1 inhibitor P-gp inducer) did not affect the steady-state pharmacokinetics of maraviroc (Table 14). Zalcitabine and lamivudine did not affect the pharmacokinetics of maraviroc.

**Table 14. Effect of Co-administered Agents on the Pharmacokinetics of Maraviroc**

Table 14. Effect of Co-administered Agents on the Pharmacokinetics of Maraviroc					
Co-administered Drug and Dose	n	Dose of maraviroc	Ratio (90% CI) of Maraviroc Pharmacokinetic Parameters without/with Co-administered Drug (No Effect = 1.00)		
			C <sub>max</sub>	AUC <sub>0-24</sub>	C <sub>min</sub>
<b>CYP2A1 and/or P-gp Inhibitors</b>					
Ketoconazole 400 mg b.i.d.	12	100 mg b.i.d.	3.75 (3.01, 4.68)	5.00 (3.98, 6.29)	3.38 (2.38, 4.78)
Rifampin 100 mg b.i.d.	8	100 mg b.i.d.	4.55 (3.37, 6.13)	4.61 (3.52, 5.98)	1.28 (0.79, 2.08)
Sequinor (soft gel capsules) 100 mg b.i.d.	11	100 mg b.i.d.	11.3 (8.86, 14.1)	7.77 (7.87, 12.14)	4.78 (3.41, 6.71)
Lopinavir/ritonavir 400 mg/100 mg b.i.d.	11	300 mg b.i.d.	9.24 (7.88, 10.7)	3.96 (3.43, 4.58)	1.97 (1.68, 2.34)
Atazanavir 400 mg q.d.	12	300 mg b.i.d.	4.18 (3.65, 4.80)	4.57 (3.30, 6.07)	1.59 (1.72, 2.58)
Atazanavir/ritonavir 300 mg/100 mg q.d.	12	300 mg b.i.d.	6.67 (5.78, 7.78)	6.87 (4.60, 5.41)	2.67 (2.32, 3.08)
Darunavir/ritonavir 600 mg/100 mg b.i.d.	12	150 mg b.i.d.	8.00 (6.35, 10.1)	4.05 (2.94, 5.58)	2.29 (1.46, 3.58)
Etravirine/ritonavir 150 mg/100 mg q.d.	11	150 mg b.i.d.	4.23 (3.47, 5.16)	2.86 (2.33, 3.51)	2.15 (1.71, 2.68)
<b>CYP2A1 and/or P-gp Inducers</b>					
Efavirenz 600 mg q.d.	12	100 mg b.i.d.	0.55 (0.43, 0.72)	0.55 (0.40, 0.62)	0.48 (0.38, 0.63)
Etravirine 600 mg q.d.	12	200 mg b.i.d. (+ efavirenz) 100 mg b.i.d. (alone)	1.08 (0.88, 1.30)	1.15 (0.88, 1.30)	1.16 (0.87, 1.59)
Rifampin 600 mg q.d.	12	100 mg b.i.d.	0.22 (0.17, 0.28)	0.37 (0.33, 0.41)	0.34 (0.26, 0.43)
Rifampin 600 mg q.d.	12	200 mg b.i.d. (+ rifampin) 100 mg b.i.d. (alone)	0.88 (0.54, 0.82)	1.04 (0.88, 1.22)	0.97 (0.72, 1.28)
Etravirine 200 mg b.i.d.	14	300 mg b.i.d.	0.61 (0.53, 0.71)	0.47 (0.38, 0.58)	0.40 (0.28, 0.57)
<b>Neurolept</b>					
Haloperidol 150 mg b.i.d.	8	300 mg single dose	0.51 (0.35, 0.75)	1.01 (0.65, 1.59)	1.54 (0.86, 2.31)
<b>CYP2A1 and/or P-gp Inhibitors and Inducers</b>					
Lopinavir/ritonavir + efavirenz 400 mg/100 mg b.i.d. + 600 mg q.d.	11	300 mg b.i.d.	6.28 (4.72, 8.38)	2.53 (2.24, 2.87)	1.01 (0.75, 1.35)
Sequinor (soft gel capsules) 100 mg b.i.d. + efavirenz 600 mg q.d.	11	100 mg b.i.d.	8.42 (6.48, 10.97)	5.00 (4.26, 5.87)	2.28 (1.64, 3.11)
Darunavir/ritonavir + efavirenz 600 mg/100 mg b.i.d. + 200 mg q.d.	10	150 mg b.i.d.	5.27 (4.51, 6.16)	3.10 (2.57, 3.74)	1.77 (1.20, 2.60)
Fosamprenavir/ritonavir 700 mg/100 mg b.i.d.	14	300 mg b.i.d.	4.74 (4.03, 5.57)	2.49 (2.18, 2.82)	1.52 (1.27, 1.82)
Fosamprenavir/ritonavir 1,400 mg/100 mg b.i.d.	14	300 mg q.d.	1.80 (1.53, 2.13)	2.28 (1.98, 2.58)	1.45 (1.20, 1.74)
Tenofovir/ritonavir 300 mg/100 mg b.i.d.	12	150 mg b.i.d.	1.80 (1.55, 2.08)	1.02 (0.85, 1.23)	0.86 (0.61, 1.21)
<b>Other</b>					
Raltegravir 400 mg b.i.d.	17	300 mg b.i.d.	0.90 (0.85, 0.96)	0.86 (0.80, 0.92)	0.78 (0.67, 0.94)

\*Compared with historical data.

**Effect of Maraviroc on the Pharmacokinetics of Concomitant Drugs:** Maraviroc is unlikely to inhibit the metabolism of co-administered drugs metabolized by the following cytochrome P enzymes (CYP1A2, CYP2B8, CYP2C8, CYP2C9, CYP2C19, and CYP3A4) or to inhibit the uptake of DATP1B1 or the export of MRP2 because maraviroc did not inhibit activity of these enzymes or transporters at clinically relevant concentrations *in vitro*. Maraviroc does not induce CYP1A2 *in vitro*. Additionally, *in vitro* studies have shown that maraviroc is not a substrate for, and does not inhibit, any of the major renal uptake inhibitors (organic anion transporter (OAT1), OAT2, organic cation transporter (OCT2), novel organic cation transporter (NCTN), and OCTN2) at clinically relevant concentrations.

*In vitro* results suggest that maraviroc could inhibit P-gp in the gut. However, maraviroc did not significantly affect the pharmacokinetics of digoxin *in vivo*, indicating maraviroc may not significantly inhibit or induce P-gp clinically.

Drug interaction trials were performed with maraviroc and other drugs likely to be co-administered or commonly used as probes for pharmacokinetic interactions (Table 14).

Co-administration of fosamprenavir 700 mg/ritonavir 100 mg twice daily and maraviroc 300 mg twice daily decreased the C<sub>max</sub> and AUC of maraviroc by 38% and 35%, respectively. Co-administration of fosamprenavir 1,400 mg/ritonavir 100 mg once daily and maraviroc 300 mg once daily decreased the C<sub>max</sub> and AUC of maraviroc by 15% and 20%, respectively. No dose adjustment is necessary when maraviroc tablets are administered 150 mg twice daily in combination with fosamprenavir/ritonavir dosed once or twice daily. Fosamprenavir should be given with ritonavir when co-administered with maraviroc tablets.

Maraviroc had no significant effect on the pharmacokinetics of zalcitabine, zidovudine, or lamivudine. Maraviroc decreased the C<sub>max</sub> and AUC of raltegravir by 27% and 37%, respectively, which is not clinically relevant. Maraviroc had no clinically relevant effect on the pharmacokinetics of maribavir, the oral contraceptive ethinylloestradiol and norgestrel, or nevirapine. No effect on the urinary 8β-hydroxycholesterol ratio, suggesting no induction of CYP2A1 *in vivo*. Maraviroc had no effect on the dehydroepiandrosterone (DHEA) to 300 mg twice daily or less than 100 mg and did not cause inhibition of CYP2B6 *in vitro* until concentrations greater than 100 μM. However, there was 234% increase in dehydroepiandrosterone (DHEA) when compared with baseline at 600 mg once daily, suggesting potential inhibition of CYP2B6 at higher doses.

## 12.4 Microbiology

## Mechanism of Action

Maraviroc is a member of a therapeutic class called CCR5 co-receptor antagonists. Maraviroc selectively binds to the human chemokine receptor CCR5 present on the cell membrane, preventing the interaction of HIV-1 gp120 and CCR5 necessary for CCR5-tropic HIV-1 to enter cells. CCR5-tropic and dual-tropic HIV-1 entry is not inhibited by maraviroc.

## Antiviral Activity in Cell Culture

Maraviroc inhibits the replication of CCR5-tropic laboratory strains and primary isolates of HIV-1 in models of acute peripheral blood monocyte infection. The mean EC<sub>50</sub> value (50% effective concentration) for maraviroc against HIV-1 group M isolates and subtype A1 and C circulating recombinant from AEl and group D isolates ranged from 0.1 to 4.5 nM (0.05 to 2.3 μg per mL) in cell culture.

When used with other antiretroviral agents in cell culture, the combination of maraviroc was not antagonistic with non-nucleoside reverse transcriptase inhibitors (NNRTIs: efavirenz and nevirapine), NRTIs (abacavir, didanosine, emtricitabine, lamivudine, stavudine, zalcitabine, and zidovudine), or protease inhibitors (PIs: amprenavir, atazanavir, darunavir, indinavir, lopinavir, nelfinavir, ritonavir, saquinavir, and zalcitabine). Maraviroc was not antagonistic with the HIV-1 gp41 fusion inhibitor enfuvirtide. Maraviroc was not active against CXCR4-tropic and dual-tropic viruses EC<sub>50</sub> values greater than 10 μM. The antiviral activity of maraviroc against HIV-1 has been well evaluated.

Fifteen of these viruses were sequenced in the gp120 encoding region and multiple amino acid substitutions with unique patterns in the R570N isolates 3 amino acid residues deleted in the V3 loop, GSA (HXB2 position 315 to 317), were associated with maraviroc resistance. The relevance of the specific gp120 substitutions observed in maraviroc-resistant isolates selected in cell culture to clinical maraviroc resistance is not known. Maraviroc-resistant viruses were characterized phenotypically by concentration-response curves that did not reach 100% inhibition in phenotypic drug assays, rather than increases in EC<sub>50</sub> values.

**Cross-Resistance in Cell Culture:** Maraviroc had antiviral activity against HIV-1 clinical isolates resistant to NNRTIs, NRTIs, PIs, and the gp41 fusion inhibitor enfuvirtide in cell culture (EC<sub>50</sub> values ranged from 0.7 to 8.9 nM (0.38 to 4.57 μg per mL) in cell culture). Maraviroc-resistant viruses that emerged in cell culture remained susceptible to enfuvirtide and the protease inhibitor saquinavir.

**Clinical Resistance:** Virologic failure on maraviroc can result from genotypic and phenotypic resistance to maraviroc, through outgrowth of undetectable CXCR4-tropic virus prior to maraviroc treatment (see *Warnings and Precautions*), through resistance to background therapy drugs (Table 15), or to drug exposure to maraviroc (See *Clinical Pharmacology* 12.2.3).

**Antiretroviral Treatment-Experienced Adult Subjects (Trial A4000227 and A4000228):** Week 48 data from treatment-experienced subjects failing maraviroc-containing regimens with CCR5-tropic virus (n = 58) have identified 22 viruses that had decreased susceptibility to maraviroc characterized in phenotypic drug assays by concentration-response curves that did not reach 100% inhibition. Additionally, CCR5-tropic virus from 2 of these treatment-failure subjects had greater than equal to 3-fold shift in EC<sub>50</sub> values for maraviroc at the time of failure.

Fifteen of these viruses were sequenced in the gp120 encoding region and multiple amino acid substitutions with unique patterns in the R570N isolates 3 amino acid residues deleted in the V3 loop, GSA (HXB2 position 315 to 317), were associated with maraviroc resistance. The relevance of the specific gp120 substitutions observed in maraviroc-resistant isolates selected in cell culture to clinical maraviroc resistance is not known. Maraviroc-resistant viruses were characterized phenotypically by concentration-response curves that did not reach 100% inhibition in phenotypic drug assays, rather than increases in EC<sub>50</sub> values.

**Cross-Resistance in Cell Culture:** Maraviroc had antiviral activity against HIV-1 clinical isolates resistant to NNRTIs, NRTIs, PIs, and the gp41 fusion inhibitor enfuvirtide in cell culture (EC<sub>50</sub> values ranged from 0.7 to 8.9 nM (0.38 to 4.57 μg per mL) in cell culture). Maraviroc-resistant viruses that emerged in cell culture remained susceptible to enfuvirtide and the protease inhibitor saquinavir.

**Clinical Resistance:** Virologic failure on maraviroc can result from genotypic and phenotypic resistance to maraviroc, through outgrowth of undetectable CXCR4-tropic virus prior to maraviroc treatment (see *Warnings and Precautions*), through resistance to background therapy drugs (Table 15), or to drug exposure to maraviroc (See *Clinical Pharmacology* 12.2.3).

**Antiretroviral Treatment-Experienced Adult Subjects (Trial A4000227 and A4000228):** Week 48 data from treatment-experienced subjects failing maraviroc-containing regimens with CCR5-tropic virus (n = 58) have identified 22 viruses that had decreased susceptibility to maraviroc characterized in phenotypic drug assays by concentration-response curves that did not reach 100% inhibition. Additionally, CCR5-tropic virus from 2 of these treatment-failure subjects had greater than equal to 3-fold shift in EC<sub>50</sub> values for maraviroc at the time of failure.

Fifteen of these viruses were sequenced in the gp120 encoding region and multiple amino acid substitutions with unique patterns in the R570N isolates 3 amino acid residues deleted in the V3 loop, GSA (HXB2 position 315 to 317), were associated with maraviroc resistance. The relevance of the specific gp120 substitutions observed in maraviroc-resistant isolates selected in cell culture to clinical maraviroc resistance is not known. Maraviroc-resistant viruses were characterized phenotypically by concentration-response curves that did not reach 100% inhibition in phenotypic drug assays, rather than increases in EC<sub>50</sub> values.

**Cross-Resistance in Cell Culture:** Maraviroc had antiviral activity against HIV-1 clinical isolates resistant to NNRTIs, NRTIs, PIs, and the gp41 fusion inhibitor enfuvirtide in cell culture (EC<sub>50</sub> values ranged from 0.7 to 8.9 nM (0.38 to 4.57 μg per mL) in cell culture). Maraviroc-resistant viruses that emerged in cell culture remained susceptible to enfuvirtide and the protease inhibitor saquinavir.

**Clinical Resistance:** Virologic failure on maraviroc can result from genotypic and phenotypic resistance to maraviroc, through outgrowth of undetectable CXCR4-tropic virus prior to maraviroc treatment (see *Warnings and Precautions*), through resistance to background therapy drugs (Table 15), or to drug exposure to maraviroc (See *Clinical Pharmacology* 12.2.3).

**Antiretroviral Treatment-Experienced Adult Subjects (Trial A4000227 and A4000228):** Week 48 data from treatment-experienced subjects failing maraviroc-containing regimens with CCR5-tropic virus (n = 58) have identified 22 viruses that had decreased susceptibility to maraviroc characterized in phenotypic drug assays by concentration-response curves that did not reach 100% inhibition. Additionally, CCR5-tropic virus from 2 of these treatment-failure subjects had greater than equal to 3-fold shift in EC<sub>50</sub> values for maraviroc at the time of failure.

Fifteen of these viruses were sequenced in the gp120 encoding region and multiple amino acid substitutions with unique patterns in the R570N isolates 3 amino acid residues deleted in the V3 loop, GSA (HXB2 position 315 to 317), were associated with maraviroc resistance. The relevance of the specific gp120 substitutions observed in maraviroc-resistant isolates selected in cell culture to clinical maraviroc resistance is not known. Maraviroc-resistant viruses were characterized phenotypically by concentration-response curves that did not reach 100% inhibition in phenotypic drug assays, rather than increases in EC<sub>50</sub> values.

**Cross-Resistance in Cell Culture:** Maraviroc had antiviral activity against HIV-1 clinical isolates resistant to NNRTIs, NRTIs, PIs, and the gp41 fusion inhibitor enfuvirtide in cell culture (EC<sub>50</sub> values ranged from 0.7 to 8.9 nM (0.38 to 4.57 μg per mL) in cell culture). Maraviroc-resistant viruses that emerged in cell culture remained susceptible to enfuvirtide and the protease inhibitor saquinavir.

**Clinical Resistance:** Virologic failure on maraviroc can result from genotypic and phenotypic resistance to maraviroc, through outgrowth of undetectable CXCR4-tropic virus prior to maraviroc treatment (see *Warnings and Precautions*), through resistance to background therapy drugs (Table 15), or to drug exposure to maraviroc (See *Clinical Pharmacology* 12.2.3).

**Antiretroviral Treatment-Experienced Adult Subjects (Trial A4000227 and A4000228):** Week 48 data from treatment-experienced subjects failing maraviroc-containing regimens with CCR5-tropic virus (n = 58) have identified 22 viruses that had decreased susceptibility to maraviroc characterized in phenotypic drug assays by concentration-response curves that did not reach 100% inhibition. Additionally, CCR5-tropic virus from 2 of these treatment-failure subjects had greater than equal to 3-fold shift in EC<sub>50</sub> values for maraviroc at the time of failure.

Fifteen of these viruses were sequenced in the gp120 encoding region and multiple amino acid substitutions with unique patterns in the R570N isolates 3 amino acid residues deleted in the V3 loop, GSA (HXB2 position 315 to 317), were associated with maraviroc resistance. The relevance of the specific gp120 substitutions observed in maraviroc-resistant isolates selected in cell culture to clinical maraviroc resistance is not known. Maraviroc-resistant viruses were characterized phenotypically by concentration-response curves that did not reach 100% inhibition in phenotypic drug assays, rather than increases in EC<sub>50</sub> values.

**Cross-Resistance in Cell Culture:** Maraviroc had antiviral activity against HIV-1 clinical isolates resistant to NNRTIs, NRTIs, PIs, and the gp41 fusion inhibitor enfuvirtide in cell culture (EC<sub>50</sub> values ranged from 0.7 to 8.9 nM (0.38 to 4.57 μg per mL) in cell culture). Maraviroc-resistant viruses that emerged in cell culture remained susceptible to enfuvirtide and the protease inhibitor saquinavir.

**Clinical Resistance:** Virologic failure on maraviroc can result from genotypic and phenotypic resistance to maraviroc, through outgrowth of undetectable CXCR4-tropic virus prior to maraviroc treatment (see *Warnings and Precautions*), through resistance to background therapy drugs (Table 15), or to drug exposure to maraviroc (See *Clinical Pharmacology* 12.2.3).

\*Includes subjects failing with CXCR4 or dual-tropic viruses because these viruses are not intrinsically susceptible to maraviroc. *In vitro* studies of treatment-naïve subjects at 18 weeks, 37 subjects failed a maraviroc-containing regimen with CCR5-tropic virus and had a tropism result at failure; 7 of these subjects had evidence of maraviroc phenotypic resistance defined as concentration-response curves that did not reach 85% inhibition. One additional subject had a greater than or equal to 3-fold shift in the EC<sub>50</sub> value for maraviroc at the time of failure. A dual analysis of the V3 loop amino acid sequence was performed from 8 of the 7 subjects. Changes in V3 loop amino acid sequence differed between each of these different subjects, even for those infected with the same virus clade, suggesting that there are multiple diverse