

(20449)

<b>Medical Benefit</b>		<b>Effective Date:</b> 07/01/15	<b>Next Review Date:</b> 05/19
<b>Preauthorization</b>	No	<b>Review Dates:</b> 05/09, 03/10, 03/11, 03/12, 03/13, 03/14, 03/15, 05/15, 05/16, 05/17, 05/18	

**Preauthorization is not required.**

*The following protocol contains medical necessity criteria that apply for this service. The criteria are also applicable to services provided in the local Medicare Advantage operating area for those members, unless separate Medicare Advantage criteria are indicated. If the criteria are not met, reimbursement will be denied and the patient cannot be billed. Please note that payment for covered services is subject to eligibility and the limitations noted in the patient's contract at the time the services are rendered.*

Populations	Interventions	Comparators	Outcomes
Individuals: <ul style="list-style-type: none"> <li>With HIV infection who are being considered for HIV coreceptor antagonist therapy</li> </ul>	Interventions of interest are: <ul style="list-style-type: none"> <li>HIV tropism testing</li> </ul>	Comparators of interest are: <ul style="list-style-type: none"> <li>No HIV tropism testing</li> </ul>	Relevant outcomes include: <ul style="list-style-type: none"> <li>Overall survival</li> <li>Disease-specific survival</li> <li>Morbid events</li> <li>Quality of life</li> <li>Hospitalizations</li> <li>Medication use</li> <li>Treatment-related morbidity</li> </ul>
Individuals: <ul style="list-style-type: none"> <li>With HIV infection receiving HIV coreceptor antagonist therapy or who have failed coreceptor antagonist therapy</li> </ul>	Interventions of interest are: <ul style="list-style-type: none"> <li>HIV tropism testing</li> </ul>	Comparators of interest are: <ul style="list-style-type: none"> <li>No HIV tropism testing</li> </ul>	Relevant outcomes include: <ul style="list-style-type: none"> <li>Overall survival</li> <li>Disease-specific survival</li> <li>Morbid events</li> <li>Quality of life</li> <li>Hospitalizations</li> <li>Medication use</li> <li>Treatment-related mortality</li> <li>Treatment-related morbidity</li> </ul>
Individuals: <ul style="list-style-type: none"> <li>With HIV infection who are undergoing tests to predict disease progression</li> </ul>	Interventions of interest are: <ul style="list-style-type: none"> <li>HIV tropism testing</li> </ul>	Comparators of interest are: <ul style="list-style-type: none"> <li>Plasma HIV RNA</li> <li>CD4 count</li> </ul>	Relevant outcomes include: <ul style="list-style-type: none"> <li>Overall survival</li> <li>Disease-specific survival</li> <li>Morbid events</li> <li>Quality of life</li> <li>Hospitalizations</li> <li>Medication use</li> </ul>

### DESCRIPTION

HIV tropism testing can determine the predominant coreceptor protein used by HIV to infect target cells. Tropism testing can help select patients for treatment with HIV coreceptor antagonists (e.g., maraviroc), which block specific coreceptor proteins.

## SUMMARY OF EVIDENCE

For individuals who have HIV infection who are being considered for HIV coreceptor antagonist therapy who receive HIV tropism testing, the evidence includes randomized controlled trials (RCTs). Relevant outcomes are overall survival, disease-specific survival, morbid events, quality of life, hospitalizations, medication use, and treatment-related morbidity. RCTs on treatment-naïve and treatment-experienced HIV-infected patients have provided evidence that selection of candidates for HIV coreceptor antagonist therapy using HIV tropism testing results in higher rates of treatment success compared with HIV coreceptor antagonist therapy without HIV tropism testing. The evidence is sufficient to determine that the technology results in a meaningful improvement in the net health outcome.

For individuals with HIV infection receiving HIV coreceptor antagonist therapy or who have failed coreceptor antagonist therapy who receive HIV tropism testing, the evidence includes post hoc analysis of RCTs and observational studies. Relevant outcomes are overall survival, disease-specific survival, morbid events, quality of life, hospitalizations, medication use, and treatment-related mortality and morbidity. Current evidence does not indicate improved outcomes with additional tropism monitoring during treatment. The evidence is insufficient to determine the effects of the technology on health outcomes.

For individuals with HIV infection who are undergoing tests to predict disease progression who receive HIV tropism testing, the evidence includes observational studies. Relevant outcomes are overall survival, disease-specific survival, morbid events, quality of life, hospitalizations, and medication use. Current evidence is inconsistent in as relates to whether HIV tropism testing independently predicts disease progression among HIV-infected patients. The evidence is insufficient to determine the effects of the technology on health outcomes.

## POLICY

HIV tropism testing (see Policy Guidelines for testing methods) may be considered **medically necessary** for selecting patients for treatment with HIV coreceptor antagonists, such as maraviroc, when there is an immediate plan to prescribe a coreceptor antagonist.

HIV tropism testing without immediate plans to prescribe HIV coreceptor antagonists such as maraviroc is **not medically necessary**.

Repeat HIV tropism testing during coreceptor antagonist treatment or after failure with coreceptor antagonists is **investigational**.

HIV tropism testing to predict disease progression (irrespective of co-receptor antagonist treatment) is **investigational**.

## POLICY GUIDELINES

Testing should be conducted immediately before intended prescribed use of maraviroc to obtain the most accurate prediction of tropism at the start of treatment.

Either phenotypic or V3 population genotypic testing may be used to determine HIV tropism; both are not necessary.

V3 population genotypic testing may be conducted by either standard V3 sequencing via Sanger methods (amplification and population sequence analysis of patient-derived V3 region) OR V3 deep sequencing methods (synonyms: ultra-deep sequencing; pyrosequencing; next-generation sequencing). In the U.S., the only currently commercially available plasma HIV DNA coreceptor genotypic test (requires HIV viral load of 1000 copies/mL or

more) includes step-wise testing, with an initial standard sequencing with reflex to V3 deep sequencing if standard sequencing detects only CCR5-tropic virus.

#### GENETIC COUNSELING

Genetic counseling is primarily aimed at patients who are at risk for inherited disorders, and experts recommend formal genetic counseling in most cases when genetic testing for an inherited condition is considered. The interpretation of the results of genetic tests and the understanding of risk factors can be very difficult and complex. Therefore, genetic counseling will assist individuals in understanding the possible benefits and harms of genetic testing, including the possible impact of the information on the individual's family. Genetic counseling may alter the utilization of genetic testing substantially and may reduce inappropriate testing. Genetic counseling should be performed by an individual with experience and expertise in genetic medicine and genetic testing methods.

#### BACKGROUND

##### HIV

HIV-1, which causes AIDS, uses coreceptor proteins (either CCR5 or CXCR4) on the surface of target cells to enter and infect the cells. The most commonly transmitted strains of HIV-1 bind to CCR5 and are said to have "tropism" for CCR5-expressing cells. Dual or mixed (D/M) tropic viruses can bind to either receptor type. It is estimated that around 85% of treatment-naive patients harbor CCR5-tropic virus only, around 15% harbor D/M virus, and less than 1% are infected with CXCR4-tropic virus alone. CXCR4-tropic virus is associated with immunosuppression and later stages of disease. Coreceptor antagonists have been designed to interfere with the interaction between HIV-1 and its coreceptors.

##### HIV Coreceptor Antagonists

Maraviroc (Selzentry) was the first coreceptor antagonist to be approved by the U.S. Food and Drug Administration (FDA). Maraviroc is a selective, slowly reversible, small-molecule antagonist of the interaction between human cell surface CCR5 and HIV-1 gp120, also necessary for HIV-1 cell infection. Blocking this interaction prevents CCR5-tropic HIV-1 entry into cells. However, CXCR4-tropic HIV-1 entry is not prevented. According to the drug's original label, maraviroc, in combination with other antiretroviral agents, is indicated for adults who are infected with only CCR5-tropic detectable HIV-1, who have evidence of viral replication and HIV-1 strains resistant to multiple antiretroviral agents.<sup>1</sup>

The currently approved maraviroc label indicates that maraviroc is indicated for combination antiretroviral treatment for adults infected with only CCR5-tropic HIV-1, without discussion of the presence of viral replication.<sup>2</sup> The FDA-approved full prescribing information for the drug states: "Tropism testing must be conducted on a current sample with a highly sensitive tropism assay that has demonstrated the ability to identify patients appropriate for use of SELZENTRY." This is because efficacy was not demonstrated in a phase 2 study of maraviroc in patients with D/M or CXCR4-tropic HIV-1. Due to potential adverse events (hepatic and cardiac toxicity), maraviroc should only be used in indicated patients.

Other HIV coreceptor antagonists are in the drug development pipeline. Cenicriviroc (Tobira Therapeutics) is a small-molecule antagonist of both CCR5 and CCR2, a receptor involved in a number of inflammatory diseases, that is currently being investigated for treatment of CCR5-tropic HIV.<sup>3</sup> In January 2015, cenicriviroc was granted fast track designation by the FDA for the treatment of nonalcoholic steatohepatitis in patients with liver fibrosis, but the drug does not yet have FDA approval.

##### HIV Tropism Testing

HIV tropism testing is available by either phenotypic or genotypic methods. Tropism testing with a phenotypic assay, a cellular-based assay that functionally determines tropism, is available with the enhanced sensitivity

Trofile<sup>®</sup> assay (ESTA; Monogram Biosciences, South San Francisco, CA). This phenotypic assay uses virus stocks pseudotyped with envelope sequences derived from patient plasma to infect cell lines engineered to express CCR5 or CXCR4 HIV-2 coreceptors. Genotypic tropism testing is based on sequencing the third variable (V3) loop of the HIV glycoprotein 120 gene; this is because the V3 loop interacts with the HIV co-receptor, and variants in V3 are associated with measurable changes in HIV tropism. Tropism assignment is derived from the sequence data using a bioinformatic algorithm such as geno2pheno. In the United States, Quest Diagnostics (Madison, NJ) offers the only commercially available genotypic HIV coreceptor tropism assay, which uses triplicate population sequencing with reflex to ultra-deep sequencing if only CCR5-tropic virus is detected. Quest Diagnostics also offers a proviral DNA tropism test (Trofile<sup>®</sup> DNA), which sequences the tropism of HIV-1 DNA that has integrated into the host genome of infected T lymphocytes via triplicate population sequencing, without the use of ultra-deep sequencing.

### REGULATORY STATUS

Clinical laboratories may develop and validate tests in-house and market them as a laboratory service; laboratory-developed tests (LDTs) must meet the general regulatory standards of the Clinical Laboratory Improvement Amendments (CLIA). HIV tropism tests are available under the auspices of the CLIA. Laboratories that offer LDTs must be licensed by the CLIA for high-complexity testing. To date, the FDA has chosen not to require any regulatory review of this test.

The FDA-approved full prescribing information for maraviroc (Selzentry<sup>™</sup>, Pfizer) states that: “Tropism testing must be conducted with a highly sensitive and specific tropism assay that has demonstrated the ability to identify patients appropriate for [maraviroc] use.”<sup>4</sup>

---

Services that are the subject of a clinical trial do not meet our Technology Assessment Protocol criteria and are considered investigational. *For explanation of experimental and investigational, please refer to the Technology Assessment Protocol.*

It is expected that only appropriate and medically necessary services will be rendered. We reserve the right to conduct prepayment and postpayment reviews to assess the medical appropriateness of the above-referenced procedures. **Some of this protocol may not pertain to the patients you provide care to, as it may relate to products that are not available in your geographic area.**

### REFERENCES

We are not responsible for the continuing viability of web site addresses that may be listed in any references below.

1. Food and Drug Administration. Full Prescribing Information: SELZENTRY. 2007; [http://www.accessdata.fda.gov/drugsatfda\\_docs/label/2007/022128lbl.pdf](http://www.accessdata.fda.gov/drugsatfda_docs/label/2007/022128lbl.pdf). Accessed November 28, 2017.
2. Food and Drug Administration. Full Prescribing Information: SELZENTRY. 2014; [http://www.accessdata.fda.gov/drugsatfda\\_docs/label/2014/022128s012lbl.pdf](http://www.accessdata.fda.gov/drugsatfda_docs/label/2014/022128s012lbl.pdf). Accessed November 28, 2017.
3. Tobira Therapeutics Inc. Efficacy, Safety, and Tolerability of Cenicriviroc (CVC) in Combination With Truvada or Sustiva Plus Truvada in HIV 1-infected, Antiretroviral Treatment-naïve, Adult Patients Infected With Only CCR5-tropic Virus. 2013; <http://clinicaltrials.gov/ct2/show/NCT01338883?term=NCT01338883&rank=1>. Accessed November 28, 2017.

4. Pfizer Inc. Selzentry™ (maraviroc) prescribing information. 2009; [http://www.accessdata.fda.gov/drugsatfda\\_docs/label/2009/022128s002lbl.pdf](http://www.accessdata.fda.gov/drugsatfda_docs/label/2009/022128s002lbl.pdf). Accessed November 28, 2017.
5. Whitcomb JM, Huang W, Fransen S, et al. Development and characterization of a novel single-cycle recombinant-virus assay to determine human immunodeficiency virus type 1 coreceptor tropism. *Antimicrob Agents Chemother*. Feb 2007;51(2):566-575. PMID 17116663
6. Reeves JD, Coakley E, Petropoulos CJ, et al. An enhanced-sensitivity Trofile HIV coreceptor tropism assay for selecting patients for therapy with entry inhibitors targeting CCR5: A review of analytical and clinical studies. *J Viral Entry*. 2009;3:94-102.
7. Panel on Antiretroviral Guidelines for Adults and Adolescents, Department of Health and Human Services. Guidelines for the Use of Antiretroviral Agents in Adults and Adolescents Living with HIV. 2017; <https://aidsinfo.nih.gov/contentfiles/lvguidelines/AdultandAdolescentGL.pdf>. Accessed November 28, 2017.
8. Cooper DA, Heera J, Goodrich J, et al. Maraviroc versus efavirenz, both in combination with zidovudine-lamivudine, for the treatment of antiretroviral-naïve subjects with CCR5-tropic HIV-1 infection. *J Infect Dis*. Mar 15 2010;201(6):803-813. PMID 20151839
9. Wilkin TJ, Goetz MB, Leduc R, et al. Reanalysis of coreceptor tropism in HIV-1-infected adults using a phenotypic assay with enhanced sensitivity. *Clin Infect Dis*. Apr 1 2011;52(7):925-928. PMID 21427401
10. Lengauer T, Sander O, Sierra S, et al. Bioinformatics prediction of HIV coreceptor usage. *Nat Biotechnol*. Dec 2007;25(12):1407-1410. PMID 18066037
11. Max Planck Institut Informatik. Geno2pheno [coreceptor] 2.5. 2014; <http://coreceptor.bioinf.mpi-inf.mpg.de/index.php>. Accessed November 28, 2017.
12. Jensen MA, Li FS, van't Wout AB, et al. Improved coreceptor usage prediction and genotypic monitoring of R5-to-X4 transition by motif analysis of human immunodeficiency virus type 1 env V3 loop sequences. *J Virol*. Dec 2003;77(24):13376-13388. PMID 14645592
13. Mullins Lab, University of Washington. Web PSSM. 2009; <https://indra.mullins.microbiol.washington.edu/webpssm/>. Accessed November 28, 2017.
14. Diez-Fuertes F, Delgado E, Vega Y, et al. Improvement of HIV-1 coreceptor tropism prediction by employing selected nucleotide positions of the env gene in a Bayesian network classifier. *J Antimicrob Chemother*. Jul 2013;68(7):1471-1485. PMID 23511232
15. Svicher V, D'Arrigo R, Alteri C, et al. Performance of genotypic tropism testing in clinical practice using the enhanced sensitivity version of Trofile as reference assay: results from the OSCAR Study Group. *New Microbiol*. Jul 2010;33(3):195-206. PMID 20954437
16. Prosperi MC, Bracciale L, Fabbiani M, et al. Comparative determination of HIV-1 co-receptor tropism by Enhanced Sensitivity Trofile, gp120 V3-loop RNA and DNA genotyping. *Retrovirology*. Jun 30 2010;7:56. PMID 20591141
17. Swenson LC, Moores A, Low AJ, et al. Improved detection of CXCR4-using HIV by V3 genotyping: application of population-based and "deep" sequencing to plasma RNA and proviral DNA. *J Acquir Immune Defic Syndr*. Aug 2010;54(5):506-510. PMID 20512044
18. Bartlett AD, MaCartney MJ, Conibear TC, et al. The utility of different bioinformatics algorithms for genotypic HIV-1 tropism testing in a large clinical cohort with multiple subtypes. *AIDS*. Jul 17 2014;28(11):1611-1617. PMID 24785955
19. Ceresola ER, Nozza S, Sampaolo M, et al. Performance of commonly used genotypic assays and comparison with phenotypic assays of HIV-1 coreceptor tropism in acutely HIV-1-infected patients. *J Antimicrob Chemother*. May 2015;70(5):1391-1395. PMID 25608585
20. Sanchez V, Masia M, Robledano C, et al. Performance of genotypic algorithms for predicting HIV-1 tropism measured against the enhanced-sensitivity Trofile coreceptor tropism assay. *J Clin Microbiol*. Nov 2010; 48(11):4135-4139. PMID 20861336
21. Strang AL, Cameron J, Booth CL, et al. Genotypic prediction of viral co-receptor tropism: correlation with enhanced Trofile. 15th Annual Conference of the British HIV Association 1-3 April 2009; 2009;Liverpool, UK.

22. Pou C, Cabrera C, Dalmau J, et al. Co-Receptor Tropism Prediction in Chronically HIV-1-Infected Subjects with Suppressed Viremia. 7th European HIV Drug Resistance Workshop; March 27–29, 2009;2009; Stockholm, Sweden.
23. Harrigan PR. MOTIVATE tropism study group. Optimization of clinical cutoffs for determining HIV co-receptor use by population and “deep” sequencing methods Infectious Diseases Society of America, 29 October-1 November 2009; 2009;Philadelphia, PA.
24. Saliou A, Delobel P, Dubois M, et al. Concordance between two phenotypic assays and ultradeep pyrosequencing for determining HIV-1 tropism. *Antimicrob Agents Chemother.* Jun 2011;55(6):2831-2836. PMID 21464245
25. Archer J, Weber J, Henry K, et al. Use of four next-generation sequencing platforms to determine HIV-1 coreceptor tropism. *PLoS One.* Nov 2012;7(11):e49602. PMID 23166726
26. Gibson RM, Meyer AM, Winner D, et al. Sensitive deep-sequencing-based HIV-1 genotyping assay to simultaneously determine susceptibility to protease, reverse transcriptase, integrase, and maturation inhibitors, as well as HIV-1 coreceptor tropism. *Antimicrob Agents Chemother.* Jan 2014;58(4):2167-2185. PMID 24468782
27. Svicher V, Alteri C, and Montano M, et al. Performance of genotypic tropism testing on proviral DNA in clinical practice: results from the DIVA study group. *New Microbiol.* Jan 2012;35(1):17-25. PMID 22378549
28. Svicher V, Alteri C, and Montano M, et al. Genotypic testing on HIV-1 DNA as a tool to assess HIV-1 co-receptor usage in clinical practice: results from the DIVA study group. *Infection.* Feb 2014;42(1):61-71. PMID 24146352
29. Brown J, Burger H, Weiser B, et al. A genotypic HIV-1 proviral DNA coreceptor tropism assay: characterization in viremic subjects. *AIDS Res Ther.* Jun 2014;11:14. PMID 24904682
30. Gulick RM, Lalezari J, Goodrich J, et al. Maraviroc for previously treated patients with R5 HIV-1 infection. *N Engl J Med.* Oct 2 2008;359(14):1429-1441. PMID 18832244
31. Katzenstein DA, Hammer SM, Hughes MD, et al. The relation of virologic and immunologic markers to clinical outcomes after nucleoside therapy in HIV-infected adults with 200 to 500 CD4 cells per cubic millimeter. AIDS Clinical Trials Group Study 175 Virology Study Team. *N Engl J Med.* Oct 10 1996;335(15):1091-1098. PMID 8813039
32. Hardy WD, Gulick RM, Mayer H, et al. Two-year safety and virologic efficacy of maraviroc in treatment-experienced patients with CCR5-tropic HIV-1 infection: 96-week combined analysis of MOTIVATE 1 and 2. *J Acquir Immune Defic Syndr.* Dec 15 2010;55(5):558-564. PMID 20703158
33. Gulick RM, Fatkenheuer G, Burnside R, et al. Five-Year Safety Evaluation of Maraviroc in HIV-1-Infected Treatment-Experienced Patients. *J Acquir Immune Defic Syndr.* Jan 1 2014;65(1):78-81. PMID 24419064
34. Saag M, Goodrich J, Fatkenheuer G, et al. A double-blind, placebo-controlled trial of maraviroc in treatment-experienced patients infected with non-R5 HIV-1. *J Infect Dis.* Jun 1 2009;199(11):1638-1647. PMID 19432546
35. Huang W, Toma J, Stawiski E, et al. Characterization of human immunodeficiency virus type 1 populations containing CXCR4-using variants from recently infected individuals. *AIDS Res Hum Retroviruses.* Aug 2009; 25(8):795-802. PMID 19678765
36. Gonzalez-Serna A, McGovern RA, Harrigan PR, et al. Correlation of the virological response to short-term Maraviroc monotherapy with standard and deep sequencing-based genotypic tropism methods. *Antimicrob Agents Chemother.* Dec 5 2012;56(3):1202-1207. PMID 22143533
37. McGovern RA, Thielen A, Mo T, et al. Population-based V3 genotypic tropism assay: a retrospective analysis using screening samples from the A4001029 and MOTIVATE studies. *AIDS.* Oct 23 2010;24(16):2517-2525. PMID 20736814
38. Harrigan PR, McGovern R, Dong W, et al. Screening for HIV tropism using population-based V3 genotypic analysis: a retrospective virological outcome analysis using stored plasma screening samples from MOTIVATE-1. 5th International AIDS Society Symposium, 19-23 July 2009; 2009; Cape Town, South Africa.

39. MCGovern R, Dong W, Mo T, et al. Optimization of clinically relevant cut-points for the determination of HIV co-receptor usage to predict maraviroc responses in treatment experienced (TE) patients using population V3 genotyping. 12th European AIDS Conference; 11-14 November 2009; 2009; Cologne, Germany.
40. Vandekerckhove LP, Wensing AM, Kaiser R, et al. European guidelines on the clinical management of HIV-1 tropism testing. *Lancet Infect Dis*. May 2011;11(5):394-407. PMID 21429803
41. McGovern RA, Thielen A, Portsmouth S, et al. Population-based sequencing of the V3-loop can predict the virological response to maraviroc in treatment-naive patients of the MERIT trial. *J Acquir Immune Defic Syndr*. Nov 1 2012;61(3):279-286. PMID 23095934
42. Swenson LC, Mo T, Dong WW, et al. Deep sequencing to infer HIV-1 co-receptor usage: application to three clinical trials of maraviroc in treatment-experienced patients. *J Infect Dis*. Jan 15 2011;203(2):237-245. PMID 21288824
43. Kagan RM, Johnson EP, Siaw M, et al. A genotypic test for HIV-1 tropism combining Sanger sequencing with ultradeep sequencing predicts virologic response in treatment-experienced patients. *PLoS One*. Oct 2012; 7(9):e46334. PMID 23029482
44. Swenson LC, Mo T, Dong WW, et al. Deep V3 sequencing for HIV type 1 tropism in treatment-naive patients: a reanalysis of the MERIT trial of maraviroc. *Clin Infect Dis*. Oct 2011;53(7):732-742. PMID 21890778
45. Heera J, Valluri S, Craig C, et al. First prospective comparison of genotypic vs. phenotypic tropism assays in predicting virologic responses to Maraviroc (MVC) in a phase 3 study: MODERN. *J Int AIDS Soc*. Nov 2014; 17(4 Suppl 3):19519. PMID 25394028
46. Poveda E, Paredes R, Moreno S, et al. Update on clinical and methodological recommendations for genotypic determination of HIV tropism to guide the usage of CCR5 antagonists. *AIDS Rev*. Jul 2012;14(3):208-217. PMID 22833064
47. Nozza S, Pignataro AR, Galli L, et al. 48 week outcomes of maraviroc-containing regimens following the genotypic or Trofile assay in HIV-1 failing subjects: the OSCAR Study. *New Microbiol*. Jul 2016;39(3):192-196. PMID 27704143
48. Garcia F, Poveda E, Perez-Elias MJ, et al. Genotypic tropism testing in proviral DNA to guide maraviroc initiation in aviremic subjects: 48-week analysis of the PROTEST study. *J Int AIDS Soc*. Nov 2014;17(4 Suppl 3):19520. PMID 25394029
49. Mellors JW, Rinaldo CR, Jr., Gupta P, et al. Prognosis in HIV-1 infection predicted by the quantity of virus in plasma. *Science*. May 24 1996;272(5265):1167-1170. PMID 8638160
50. Mellors JW, Munoz A, Giorgi JV, et al. Plasma viral load and CD4+ lymphocytes as prognostic markers of HIV-1 infection. *Ann Intern Med*. Jun 15 1997;126(12):946-954. PMID 9182471
51. Philpott SM. HIV-1 coreceptor usage, transmission, and disease progression. *Curr HIV Res*. Apr 2003; 1(2):217-227. PMID 15043204
52. Moyle GJ, Wildfire A, Mandalia S, et al. Epidemiology and predictive factors for chemokine receptor use in HIV-1 infection. *J Infect Dis*. Mar 15 2005;191(6):866-872. PMID 15717260
53. Weber J, Piontkivska H, Quinones-Mateu ME. HIV type 1 tropism and inhibitors of viral entry: clinical implications. *AIDS Rev*. Apr-Jun 2006;8(2):60-77. PMID 16848274
54. Daar ES, Kesler KL, Petropoulos CJ, et al. Baseline HIV type 1 coreceptor tropism predicts disease progression. *Clin Infect Dis*. Sep 1 2007;45(5):643-649. PMID 17683002
55. Fatkenheuer G, Nelson M, Lazzarin A, et al. Subgroup analyses of maraviroc in previously treated R5 HIV-1 infection. *N Engl J Med*. Oct 2 2008;359(14):1442-1455. PMID 18832245
56. Raymond S, Maillard A, Amiel C, et al. Virological failure of patients on maraviroc-based antiretroviral therapy. *J Antimicrob Chemother*. Feb 2015;70(6):1858-1864. PMID 25700719
57. Wilkin TJ, Gulick RM. CCR5 Antagonism in HIV Infection: Current Concepts and Future Opportunities. *Annu Rev Med*. Feb 18 2012;63:81-93. PMID 22034870
58. Almeida FJ, Zapparoli MS, Moreira DH, et al. Association of X4 tropism with disease progression in antiretroviral-treated children and adolescents living with HIV/AIDS in Sao Paulo, Brazil. *Braz J Infect Dis*. May-Jun 2014;18(3):300-307. PMID 24275366

59. Visseaux B, Charpentier C, Rouard C, et al. HIV-2 X4 tropism is associated with lower CD4+ cell count in treatment-experienced patients. *AIDS*. Sep 10 2014;28(14):2160-2162. PMID 25265081
60. Fontdevila MC, Cozzi-Lepri A, Phillips A, et al. Plasma HIV-1 tropism and risk of short-term clinical progression to AIDS or death. *J Int AIDS Soc*. Nov 2014;17(4 Suppl 3):19685. PMID 25397435
61. Saracino A, Bruno G, Scudeller L, et al. Does HIV-1 co-receptor tropism correlate with fibrosis progression in HIV/HCV co-infected patients? *J Clin Virol*. Mar 2014;59(3):167-171. PMID 24461764
62. Casadella M, Cozzi-Lepri A, Phillips A, et al. Plasma HIV-1 Tropism and the Risk of Short-Term Clinical Progression to AIDS or Death. *PLoS One*. 2017;12(1):e0166613. PMID 28129343
63. Castagna A, Monno L, Carta S, et al. Switch of predicted HIV-1 tropism in treated subjects and its association with disease progression. *Medicine (Baltimore)*. Nov 2016;95(44):e5222. PMID 27858869
64. Aberg JA, Gallant JE, Ghanem KG, et al. Primary care guidelines for the management of persons infected with HIV: 2013 update by the HIV Medicine Association of the Infectious Diseases Society of America. *Clin Infect Dis*. Jan 2014;58(1):1-10. PMID 24343580
65. Department of Health and Human Services. Guidelines for the Use of Antiretroviral Agents in HIV-1-Infected Adults and Adolescents: Considerations for Antiretroviral Use in Special Patient Populations. 2016; <http://aidsinfo.nih.gov/guidelines/html/1/adult-and-adolescent-arv-guidelines/21/hiv-infected-adolescents-and-young-adults>. Accessed November 28, 2017.