Lopinavir/ritonavir Pre-Registrational Usage in France (ATU): A 2000 to 2001 Insight into the Therapeutic Trends and Outcomes in HIV Treatment-Experienced Patients

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BACKGROUND

Lopinavir, co-formulated with a low dose of ritonavir, takes advantage of the P450 cytochrome inhibitor properties of ritonavir. Lopinavir is exquisitely sensitive to pharmacokinetic enhancement by ritonavir, resulting in substantially increased LPV drug exposure. The mean LPV ratio of C_{trough} (protein binding adjusted) to EC_{top} (Inhibitory Quotient or IO) for wild-type virus is 7-75 (at 400/100 mg BID dose), potentially providing a pharmacological barrier to the emergence of viral resistance and superior antiviral activity, contributing to durability.

METHODS

Selection criteria, initially restricted to patients with CD₄ < 200 cells/mm³ and viral load >4 log₁₀ copies/mL, were modified to authorize the use of LPV/r in a broader 1-experienced population.

At baseline, demographics, medical history, virology (HIV-RNA), immunology (CD_d), prior and current antiretroviral medications and genotype were collected. Follow-up data was provided on a voluntary basis for HIV-RNA, CD₄ and Serious Adverse Events.

The overall enrollment in this cohort was 3,819 patients. Patients excluded from this analysis included 193 patients previously enrolled in the Expanded Access Program (EAP) and 179 patients with "Nominative ATU status." "Nominative ATU status" was granted for patients who did not fulfill the selection criteria of the ATU, i.e., children.

Number (N) refers to number of subjects for whom specific data was available.

RESULTS

Summary of Baseline Demographic and Disease Characteristics

A total of 3,447 patients were included in this summary. The mean baseline HIV RNA and CD₄ count for these patients were 4.66 log₁

Demographic and disease characteristics are presented in Table 1.

Table 1. Baseline Demographic and Disease Characteristics

CD₄ Count (cells/mm³ Age (N = 1805) Mean Median Asymptomatic (Stage A)
Symptomatic (Stage B)
AIDS – Indicator (Stage C) 16.8% 31.9% 51.3% HIV Diagnosis Date (N = 2753) HIV RNA (log₁₀ copies/mL)

Figure 1. Baseline Patient Population by Age



50% of patients were \ge 40 years with the single most highly represented group being 35-40 years of age (30%).

Figure 2. Baseline Patient and Population by CDC Stage (N = 2465)

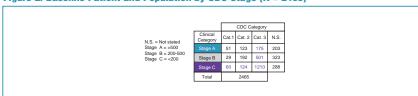


Figure 3. Distribution of Baseline CD4 Cell Count for Patients Initiating Kaletra Therapy (N = 3415)

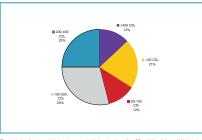


Figure 4. Distribution of Baseline Viral Load for Patients Initiating Kaletra Therapy (N = 3412)

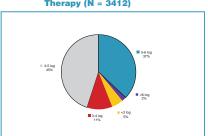


Figure 5. Trend in Baseline CD₄ Count During

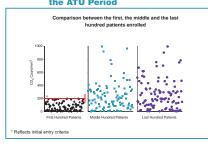
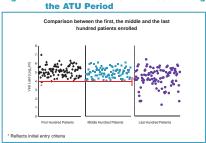


Figure 6. Trend in Baseline Viral Load During



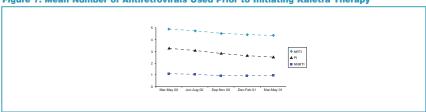
As the selection criteria became less restrictive, the mean baseline status of the patient population shifted to healthier patients, as reflected by Figures 5 and 6.

Median CD₄ count was 64, 180, and 192 for the first, middle and last hundred patients, respectively, in Figure 5.

Median viral load was 5.15, 4.85, and 4.55 log₁₀/mL for the first, middle and last hundred patients, respectively, in Figure 6.

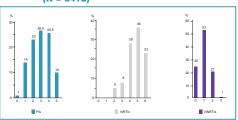
Prior Antiretroviral Experience

Figure 7. Mean Number of Antiretrovirals Used Prior to Initiating Kaletra Therapy



• Figure 7 demonstrates the mean number of prior NRTIs, PIs and NNRTIs used prior to the ATU experience.

Figure 8. Number of Previous Antiretroviral Drugs (N = 3412)



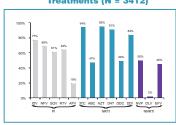
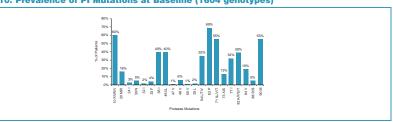


Figure 10. Prevalence of PI Mutations at Baseline (1604 genotypes)



Baseline Genotype

- *HIV RNA sequence was performed in this heavily ARV-experienced patient population according to the French guidelines (Rapport Delfraissy 2000) and were provided, when available, on a voluntary basis.
- *The prevalence of multiple baseline PI mutations associated with cross resistance within the PI class was high (Figure 10). The majority of isolates contained one or more primary mutations' at positions 82, 84 and 90, as well as secondary mutations at positions 10, 46, 54 and 71.

Figure 11. Number of PI-Associated Mutations at Baseline (1604 patients)

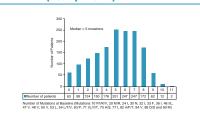
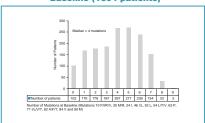


Figure 12. Number of Mutations from LPV



- The baseline median LPV mutation score (number of mutations associated with reduced susceptibility to LPV) was 4 (Figure 12).
- *Approximately half of the patients (777/1604, 48%) had a baseline LPV mutation score of 4-6 mutations (Figure 12).

Concomitant Antiretroviral Therapy

Figure 13. Description of LPV/r-Containing Regimens (N = 1828)

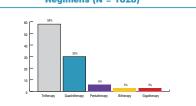
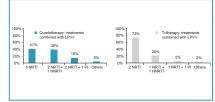


Figure 14. Description of LPV/r-Containing Regimens (N = 1828)



- A sizeable minority of patients (30%) received three additional drugs, consisting mainly of either three NRTIs (41%) or two NRTIs and one NNRTI (39%). Nine percent of patients received more than three concomitant ARVs. Concomitant use of another PI was uncommon in this production.

Figure 15. Associated Antiretroviral Treatments (N = 1828)



CONCLUSIONS

With 50% of patients in the >40 years of age group, a benefit of HAART therapy and its relation to the aging of the HIV population is suggested. This was further documented by comparing these results to the period of protease inhibitor introduction, where the most highly represented group (Norvir ATU experience, 1996) was

The ATU patient profile is that of highly antiretroviral experienced HIV-infected patients with an advanced stage of the disease. Prior antiretroviral experience for these patients typically included at least 3 Pls, 5 NRTis and 1 NNRTI.

REFERENCE

Hirsch, MS, et al. Antiretroviral drug resistance testing in adults with HIV infection: Implications for Clinical Management. International AIDS Society - USA Panel JAMA 1998; 279;1984-94

ACKNOWLEDGMENTS