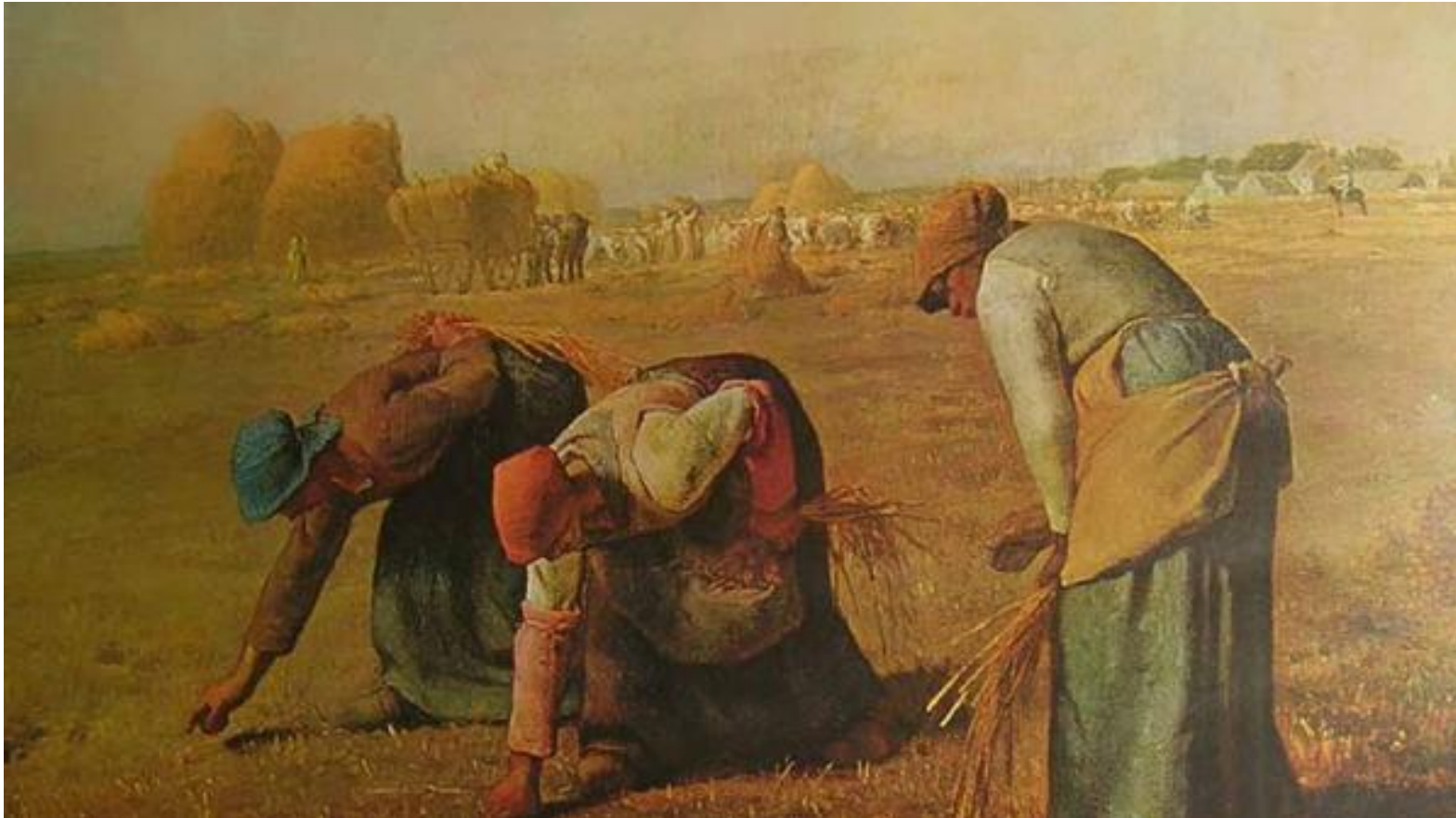




IL PAZIENTE COINFETTO TRATTATO CON I NUOVI DAA: l'esperienza della coorte SCOLTA.



Francois Millet, "Le spigolatrici" (1857)

Barbara Menzaghi



BACKGROUND

There are few data on the real-world experience of oral hepatitis C virus (HCV) direct-acting antiviral (DAA) drug combinations in HIV/HCV coinfecting patients (pts).

The aim of this study is to evaluate the safety and the efficacy of DAA therapies in a cohort of HIV/HCV patients.



METHODS

The SCOLTA project (Surveillance Cohort Long-Term Toxicity of Antiretrovirals/Antivirals) was set up as an active pharmacovigilance system for new antiretroviral drugs.

This online recording system was established by the CISAI group (Italian Coordinators for the Study of Allergies and HIV Infection; <http://www.cisai.info>).

With the introduction of new drugs for the treatment of chronic hepatitis HCV, this system was modified to assess the DAA treatment efficacy, safety and tolerability in Italian HCV patients, both monoinfected and coinfecting with HIV.

METHODS

- *Twenty-five Italian infectious diseases centers enroll patients and collect their data through this on-line system*
- *As this is an observational study, the local physicians establish the antiviral therapy.*
- *Periodical evaluations of these patients, including physical examination and laboratory tests were performed.*
- *Any reason for treatment interruption was recorded in a standard form.*





RESULTS

Overall 1303 pts (30.7% HCV/HIV) were included in this study

68.7% were males; median age was 54.0 years.

Among them, 62.4% had F4 and 27.2% F3 fibrosis.

HCV/HIV and HCV pts were similar in terms of baseline fibrosis.

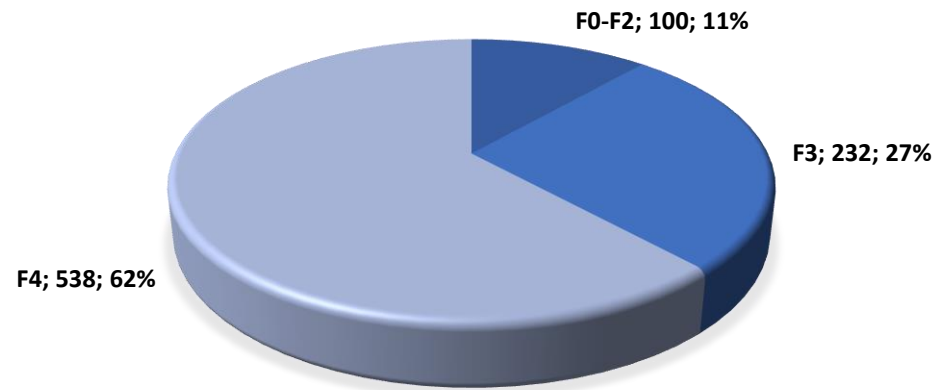
HCV/HIV subjects were younger, more frequently males and harbored more frequently HCV genotype 1a or 3.

Patients with HCV genotype 1 infection received more frequently SOF/LDV instead of 3D or SOF/SIM in case of HIV coinfection.

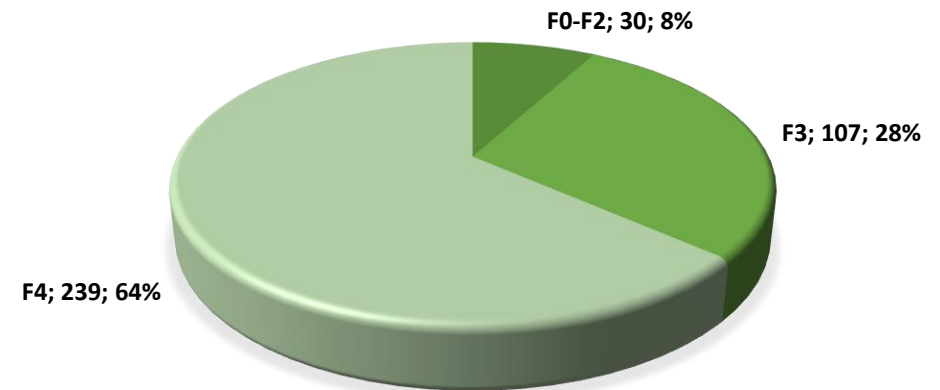
| Baseline characteristics | HCV n (902, 69.3%) | HCV/HIV n (400, 30.7%) | P |
|--|-----------------------|---------------------------|-------------------|
| Females | 316 (35.0) | 66 (16.5) | <0.0001 |
| Median age (min-max) | 56 (25-87) | 52 (34-76) | <0.0001 |
| METAVIR score | | | |
| F0-F2 | 100 (11.5) | 30 (8.0) | <u>0.17</u> |
| F3 | 232 (26.7) | 107 (28.5) | |
| F4 | 538 (61.8) | 239 (63.5) | |
| Genotype | | | |
| 1a | 164 (18.2) | 161 (40.2) | <u><0.0001</u> |
| 1b | 403 (44.6) | 55 (13.8) | |
| 2 | 107 (11.8) | 4 (1.0) | |
| 3 | 123 (13.6) | 118 (29.5) | |
| 4 | 106 (11.7) | 62 (15.5) | |
| HCV RNA T0 > 800,000 UI/mL | 422 (46.8) | 222 (30.7) | 0.004 |
| Undetectable HCV RNA 4 wks. | 373 (48.5) | 152 (49.0) | 0.86 |
| Ribavirin DAA regimen | 609 (67.5) | 273 (68.2) | 0.80 |
| 2D | 34 (3.8) | 22 (5.6) | <u><0.0001</u> |
| 3D | 189 (21.4) | 54 (13.7) | |
| SIM/DAC | 4 (0.4) | 0 | |
| SIM/PEG | 26 (2.9) | 3 (0.8) | |
| SOF/RIBA | 178 (20.1) | 82 (20.9) | |
| SOF/DAC | 108 (12.2) | 77 (19.6) | |
| SOF/LED | 124 (14.0) | 82 (20.9) | |
| SOF/SIM | 222 (25.1) | 73 (18.6) | |
| Outcome | | | |
| SVR 12 | 652 (94.1) | 302 (91.8) | 0.19 |
| Failure | 8 (1.2) | 8 (2.4) | |
| Relapse | 25 (3.6) | 17 (5.2) | |
| Interruption | 8 (1.2) | 2 (0.6) | |



HCV

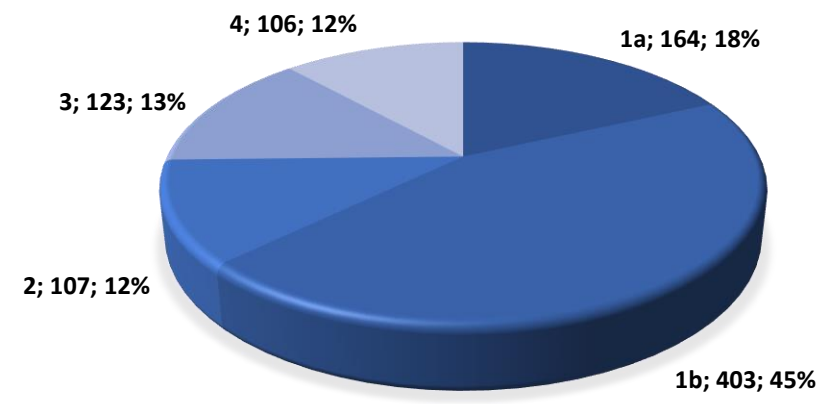


HCV/HIV

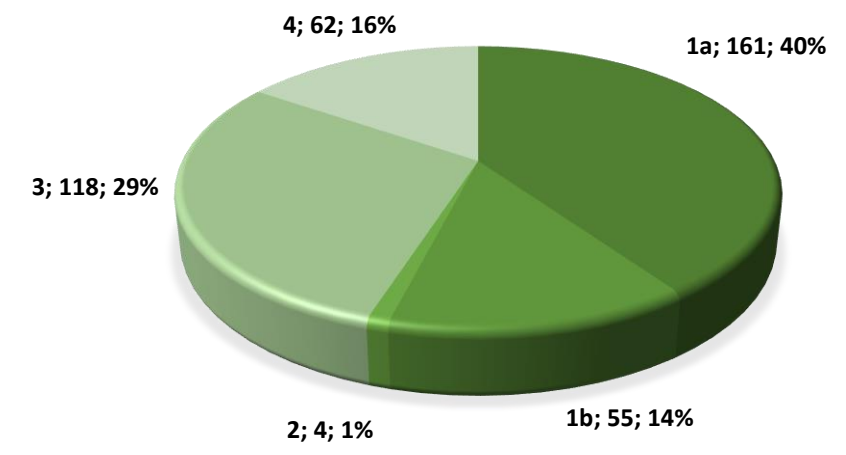




HCV

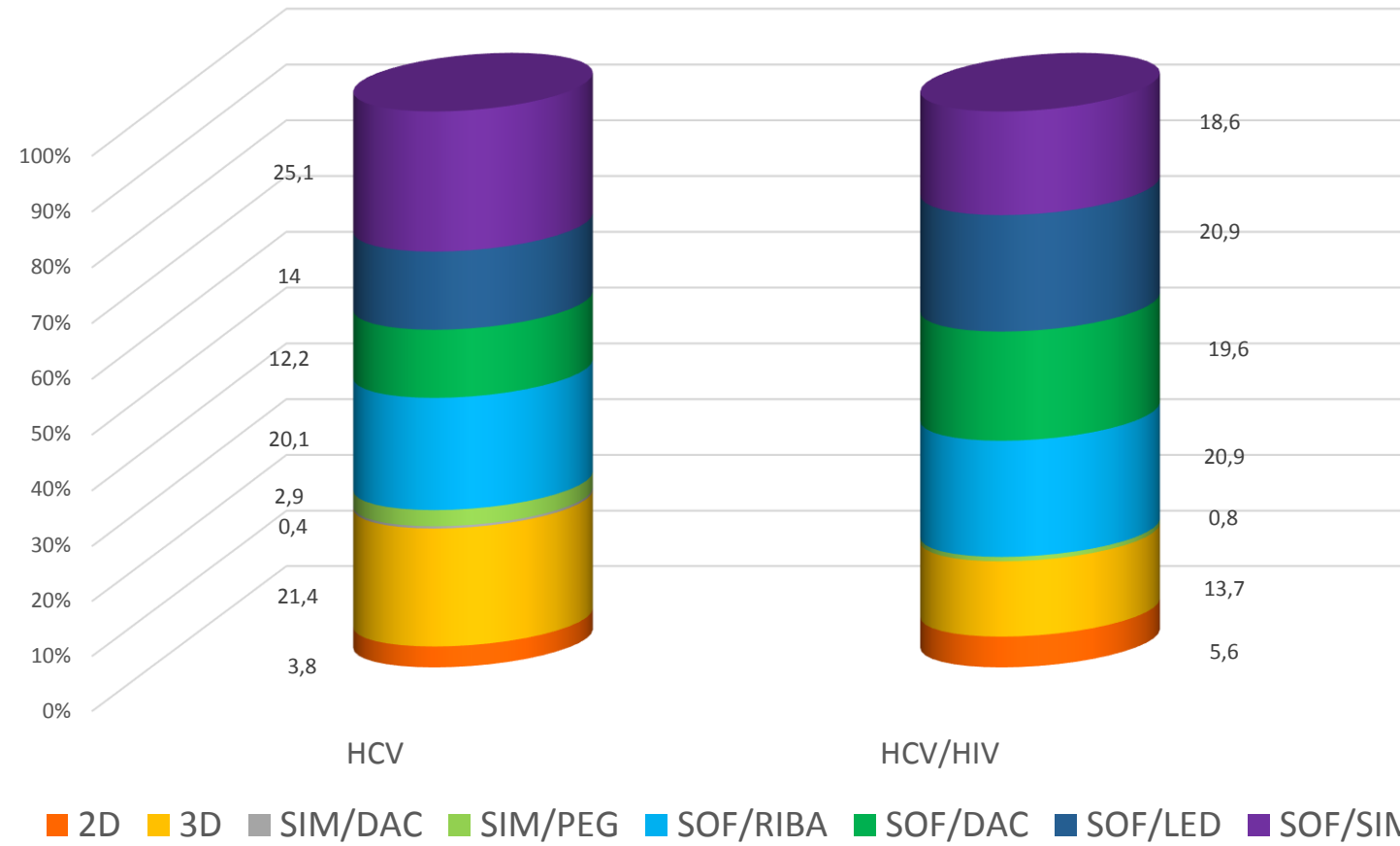


HCV/HIV





DAA regimen





HIV-HCV

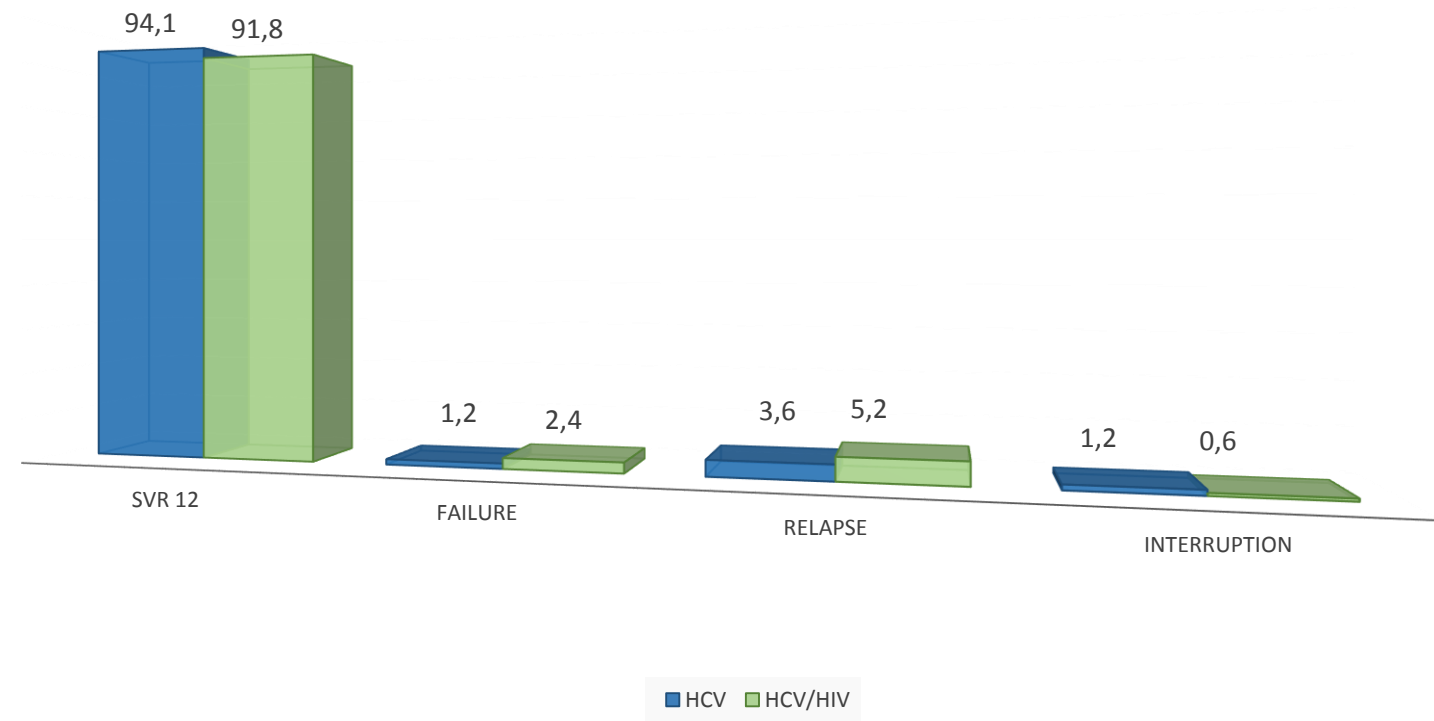
Excluding pts who prematurely discontinued treatment (n=10, 1.0%, 7 due to adverse events, 3 to death), SVR12 rate was 92.4% and 95.2% in HCV/HIV and HCV pts, respectively (p=0.07).

| Baseline characteristics | HCV | HCV/HIV | P |
|--------------------------|----------------|----------------|-------------|
| | n (902, 69.3%) | n (400, 30.7%) | |
| Outcome | | | <u>0.19</u> |
| SVR 12 | 652 (94.1) | 302 (91.8) | |
| Failure | 8 (1.2) | 8 (2.4) | |
| Relapse | 25 (3.6) | 17 (5.2) | |
| Interruption | 8 (1.2) | 2 (0.6) | |

The analyses were conducted in strata of HCV genotype. Including in the logistic regression terms for gender, age, baseline HCVRNA and DAA regimen, **HIV co-infection was not associated to a worse virological outcome (both in the per-protocol and intention-to-treat analysis).**



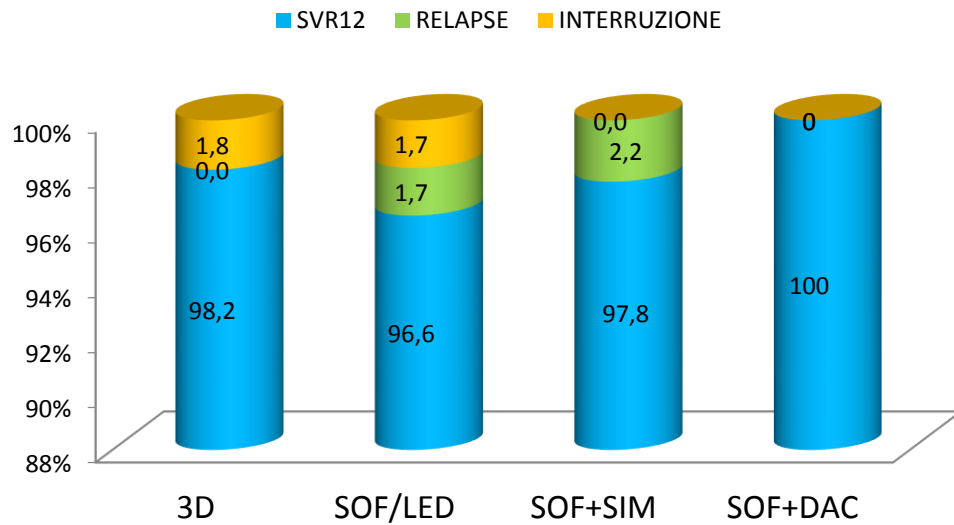
Outcome



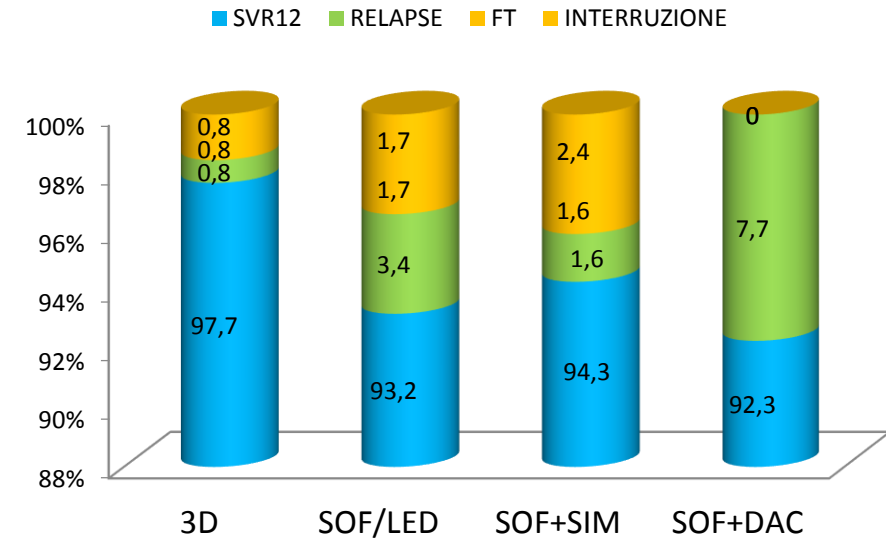


RESULTS

Genotipo 1a



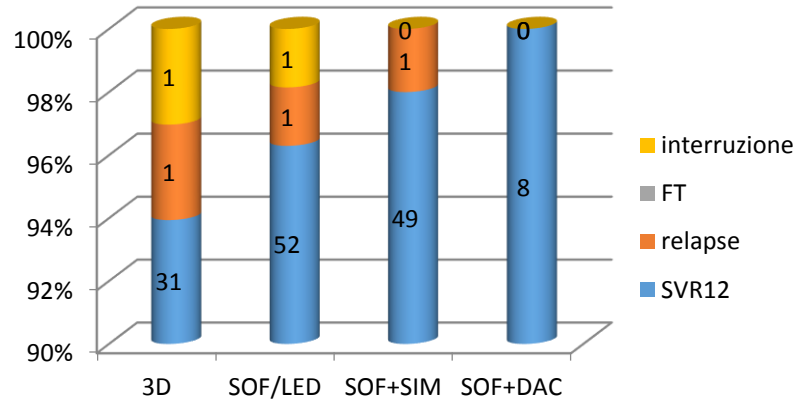
Genotipo 1b



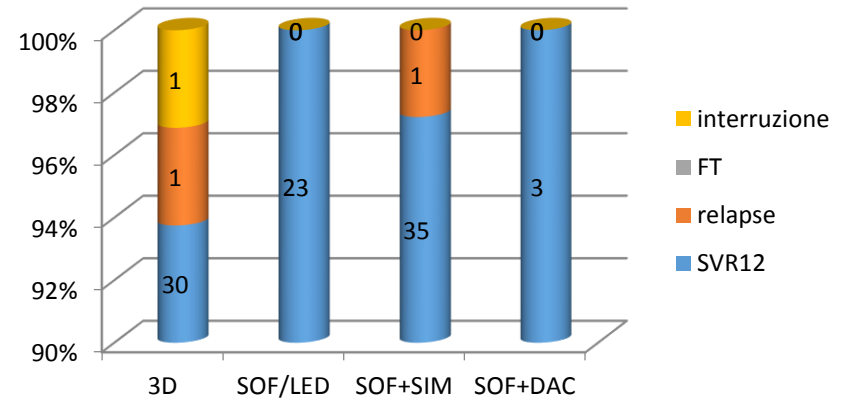


Genotipo 1a

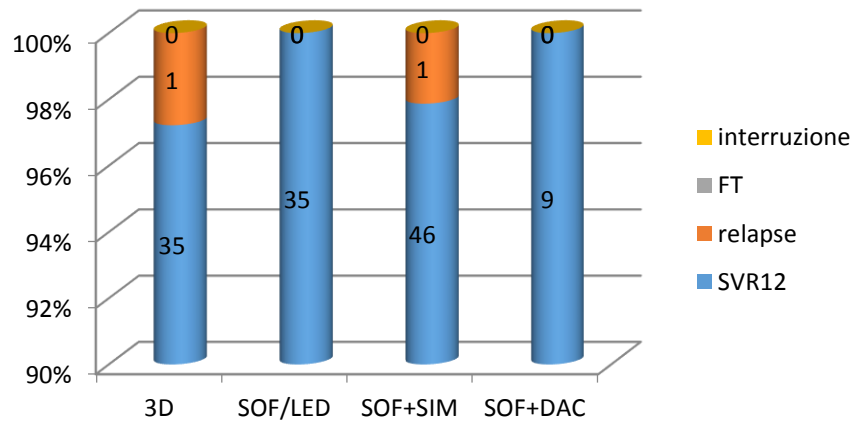
HCV/HIV



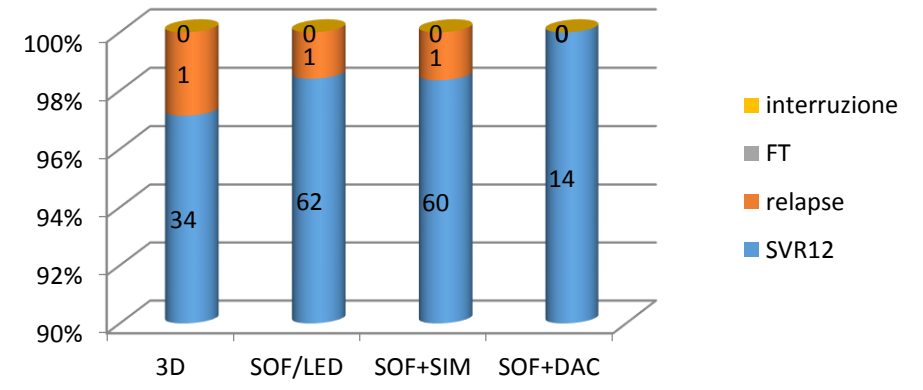
F3



HCV



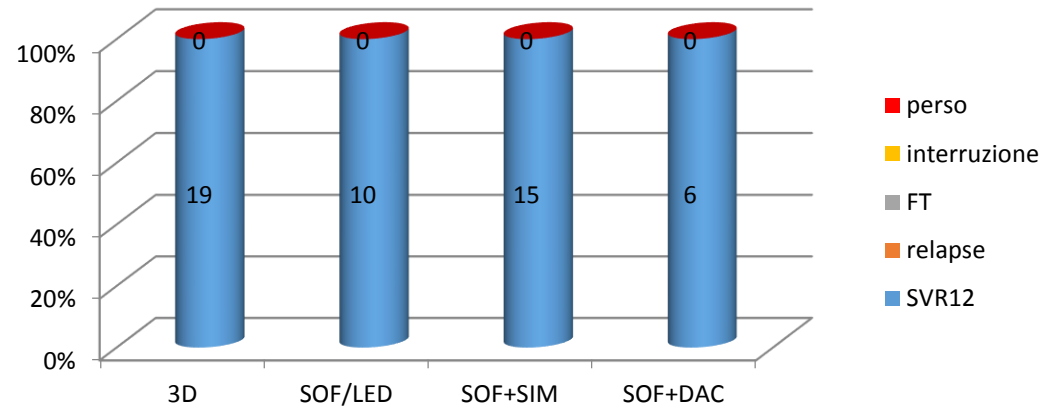
F4



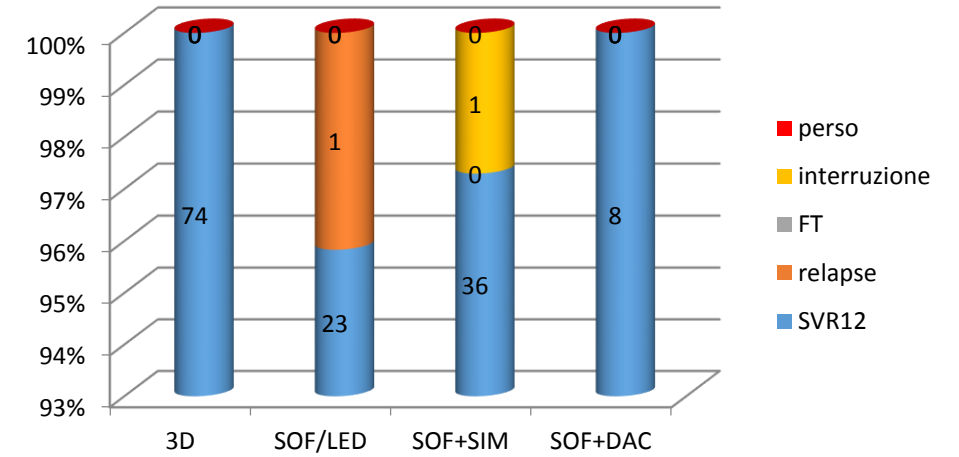


Genotipo 1b

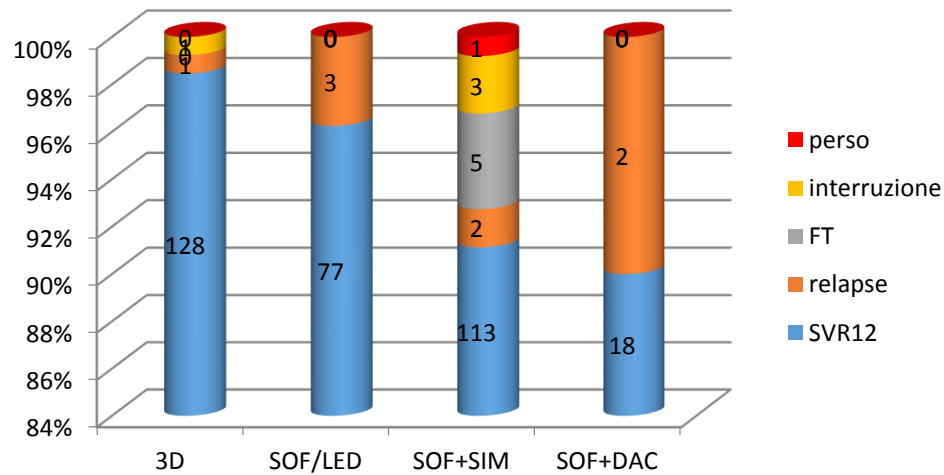
HCV/HIV



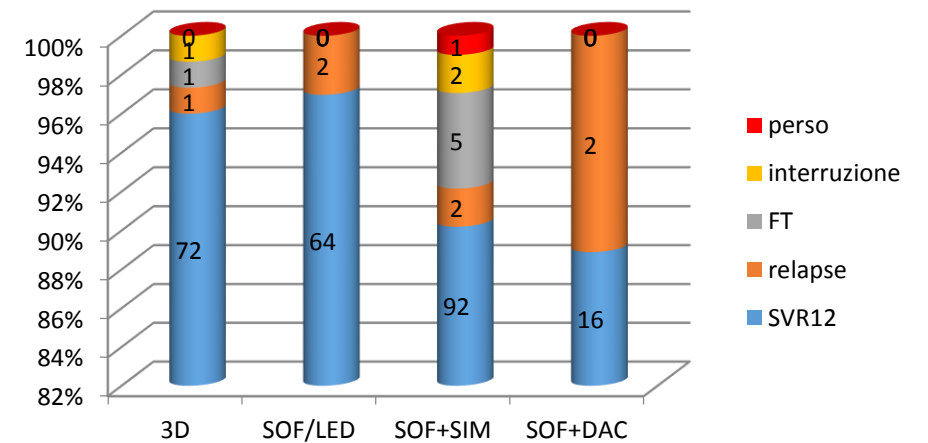
F3



HCV



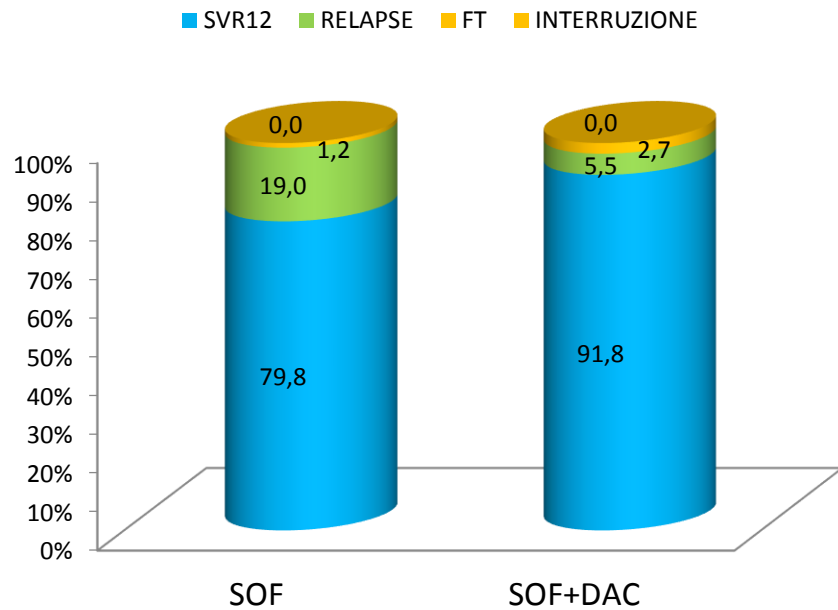
F4



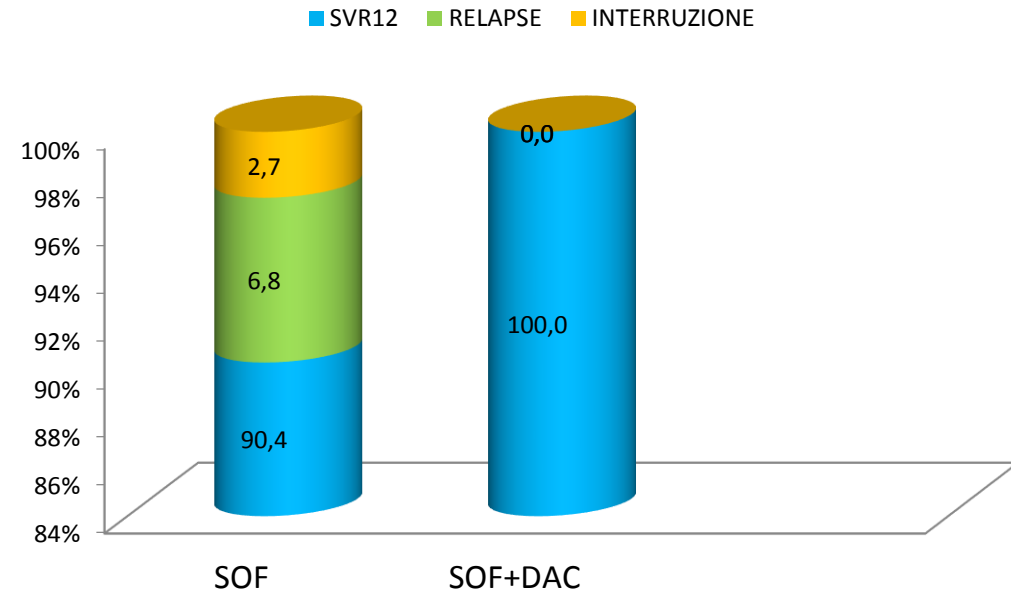


RESULTS

Genotipo 3



Genotipo 2

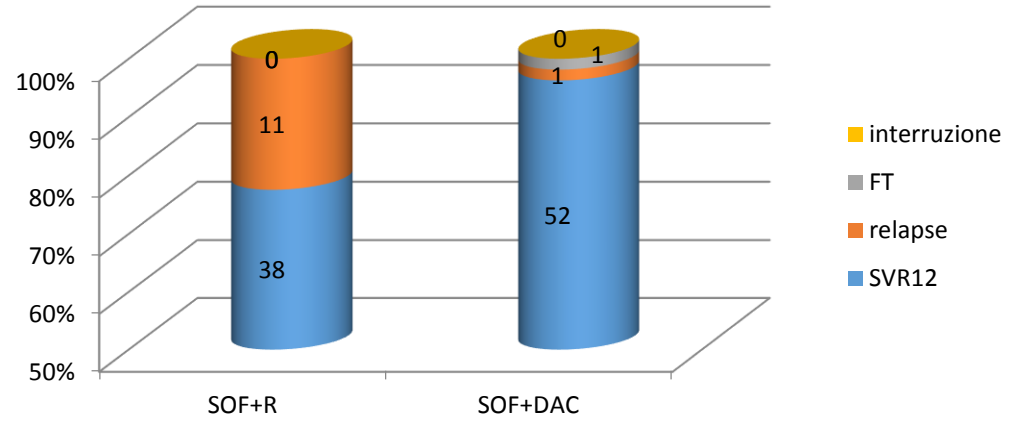


In genotype 3, SOF/DAC regimen (with or without ribavirin) was protective against failure (OR=0.32, 95% CI 0.11-0.96) as compared to SOF/RIBA.

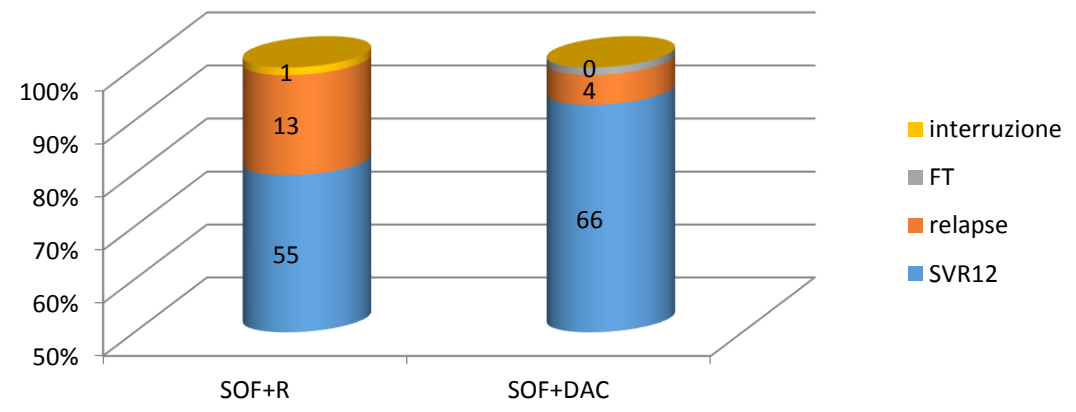


Genotipo 3

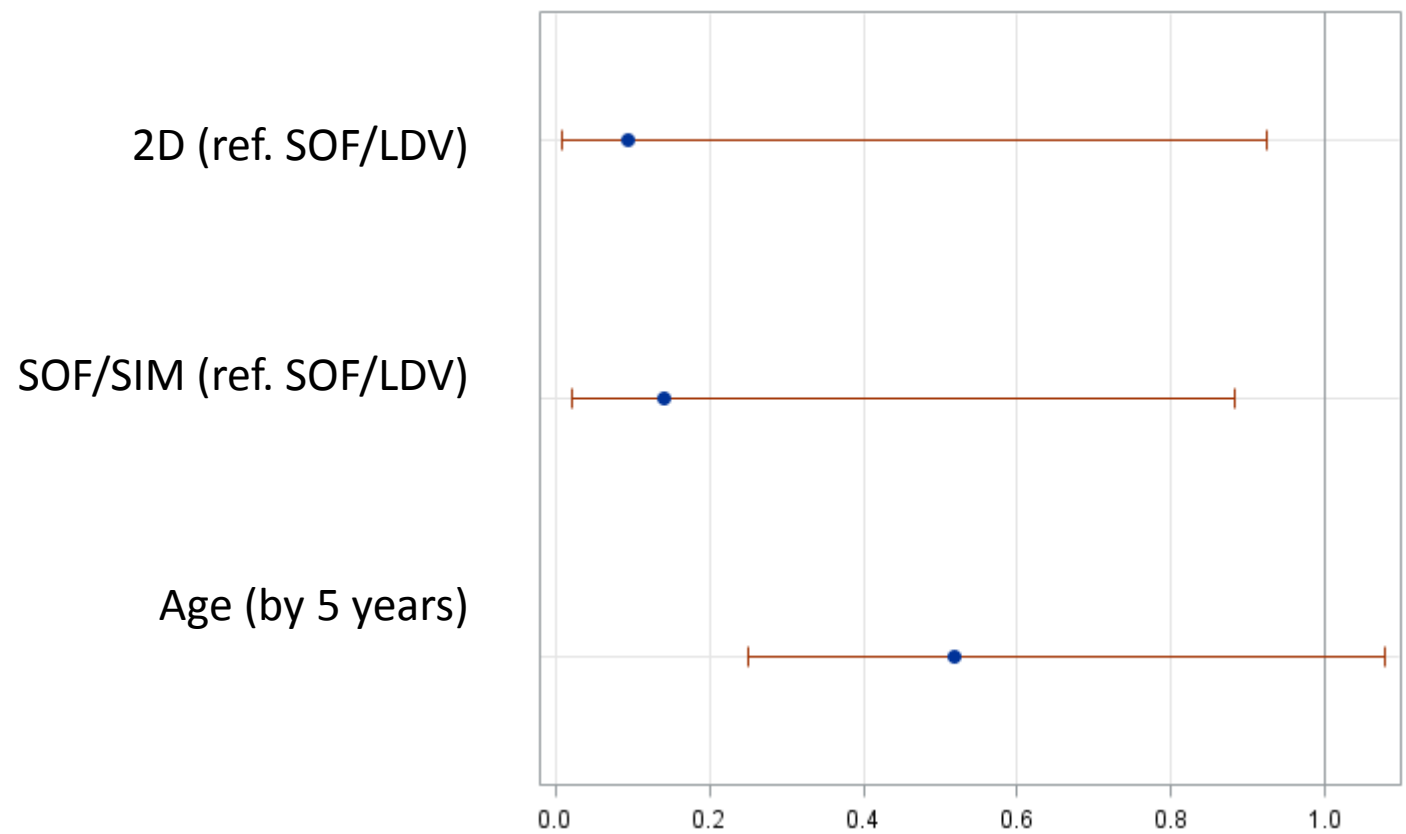
HCV/HIV



F4



RESULTS

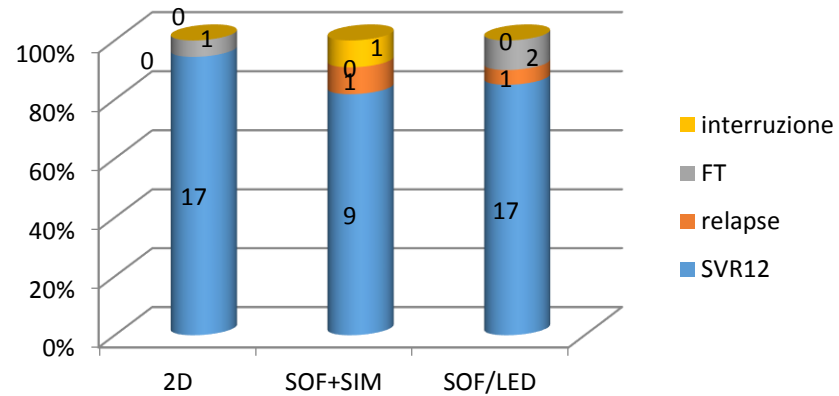


In genotype 4, as compared to SOF/LDV, 2D and SOF/SIM were less likely to experience failure (respectively OR 0.09, 95% CI 0.01-0.92 and OR 0.14, 95% CI 0.02-0.88).

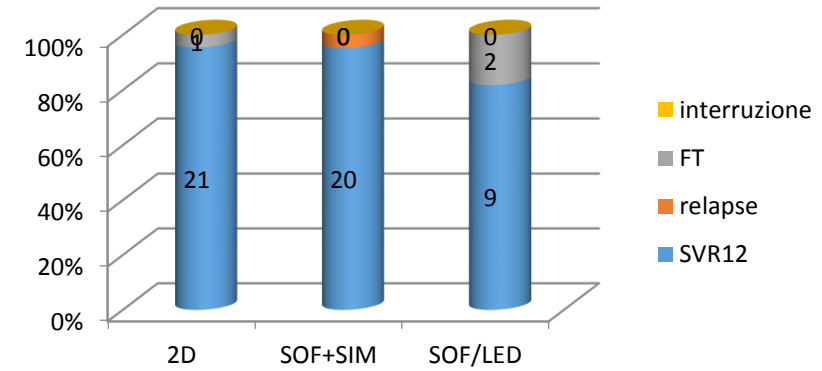


Genotipo 4

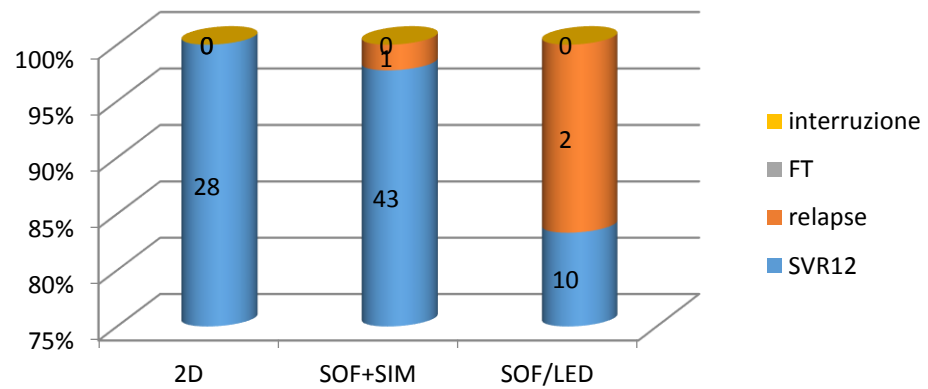
HCV/HIV



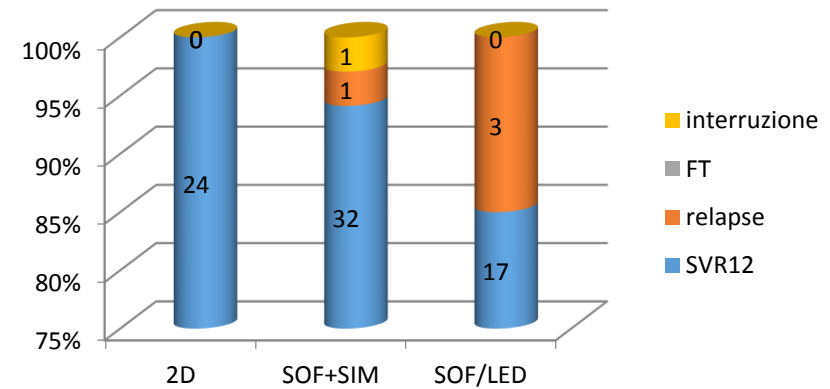
F3



HCV



F4





CONCLUSIONS

HIV co-infection did not affect the outcome of HCV treatment in a real life cohort.

In this study, we confirmed that genotype 3 HCV was associated with a lower SVR to DAA treatment.

In real life, high and similar rates of SVR were observed between mono- and co-infected pts.

SOF/DAC in genotype 3 and SOF/SIM and 2D in 4 were associated to better SVR, but because of the low number of failures this finding needs to be confirmed.



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