

Goals for Treatment and Predicting Response



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Disclosure Information



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Outline



- Rationale and goals for treatment (SVR, disease progression, survival)
- Viral factors that predict response to therapy (HCV genotype, HCV RNA level)
- Host factors that predict response to therapy (race, age, sex, *IL28B* genotype)
- Combined factors (degree of fibrosis)

Rationale and Goals for Treatment (SVR, Improve Histology, Survival)

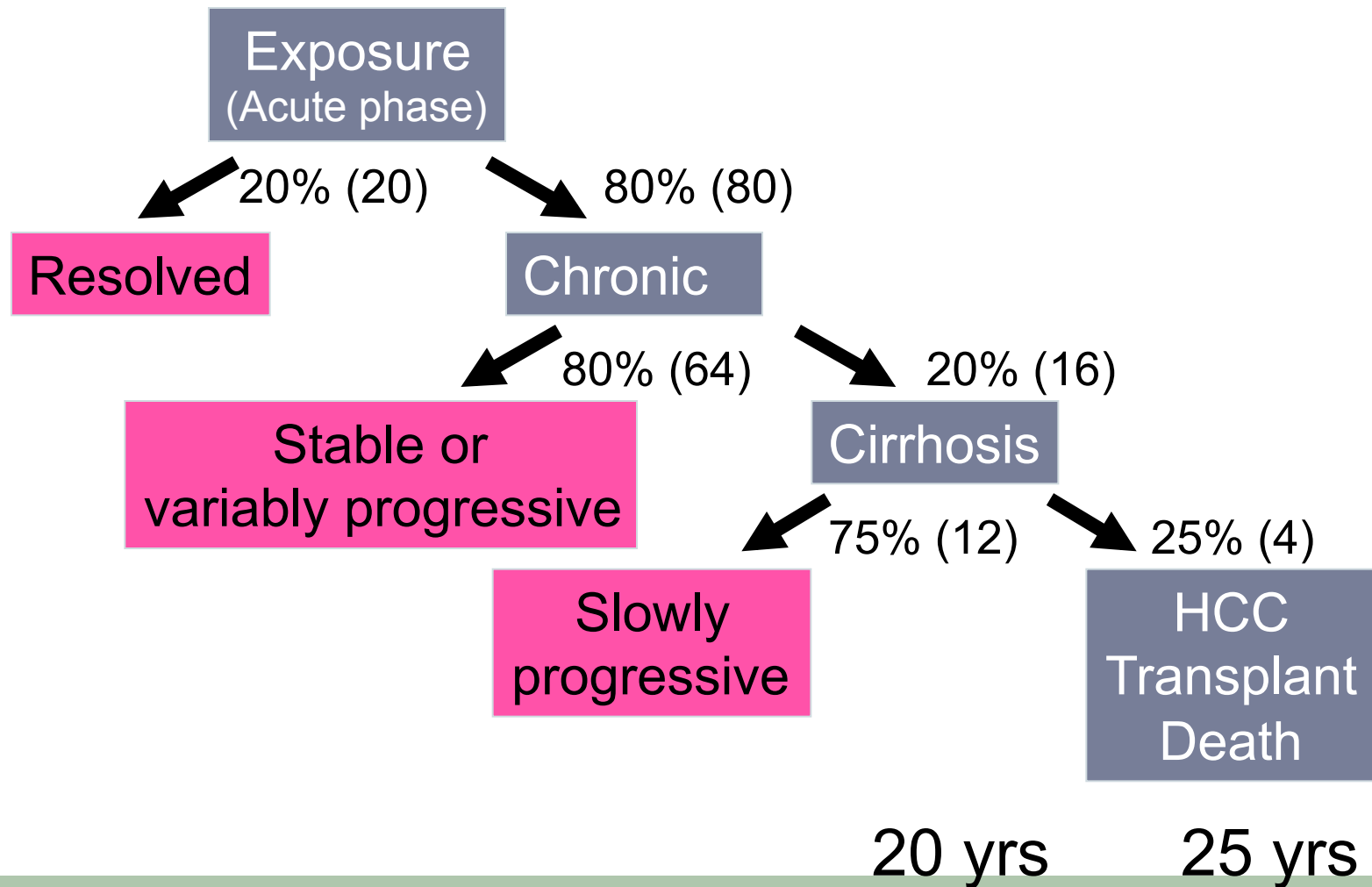


Rationale and Goals for Treatment



- Chronic HCV is a progressive disease that can produce significant morbidity and mortality if left untreated
- HCV RNA virus without latent, stable intermediate
- Requires continuous replication
- SVR is possible and connotes clinical cure
- Defined as clearance of HCV RNA 24 weeks after completion of a course of antiviral therapy
 - SVR 12 weeks after discontinuation of treatment equates to SVR 24 weeks after discontinuation of treatment

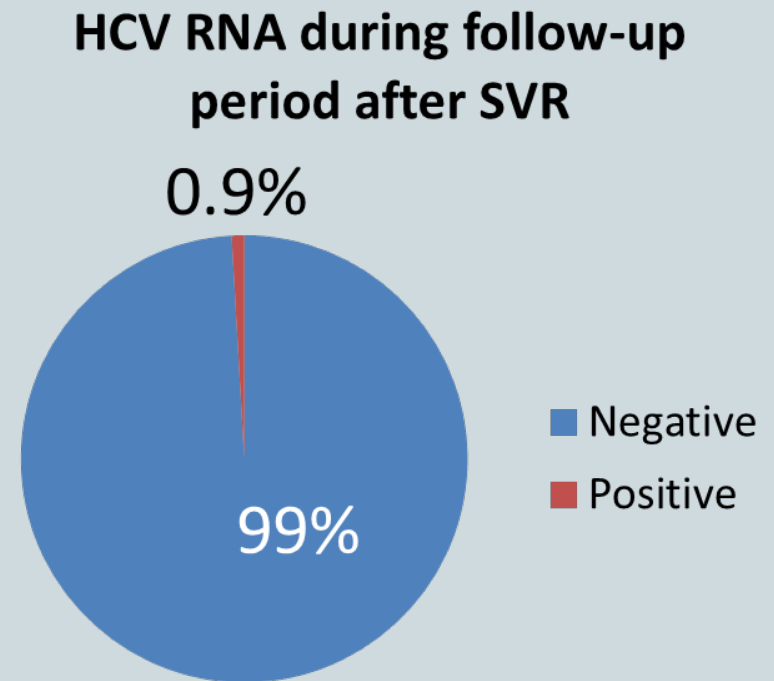
Natural History of Chronic HCV



An SVR is Truly Sustained



- Cohort study
- Patients from 9 trials
- HCV: 1243 patients
- HIV/HCV: 100 patients
- Follow-up: mean 3.9 yrs (range, 0.8-7.1 yrs)
- 0.9% HCV RNA positive after treatment ended



Impact of SVR on Natural History in Patients With Advanced Fibrosis

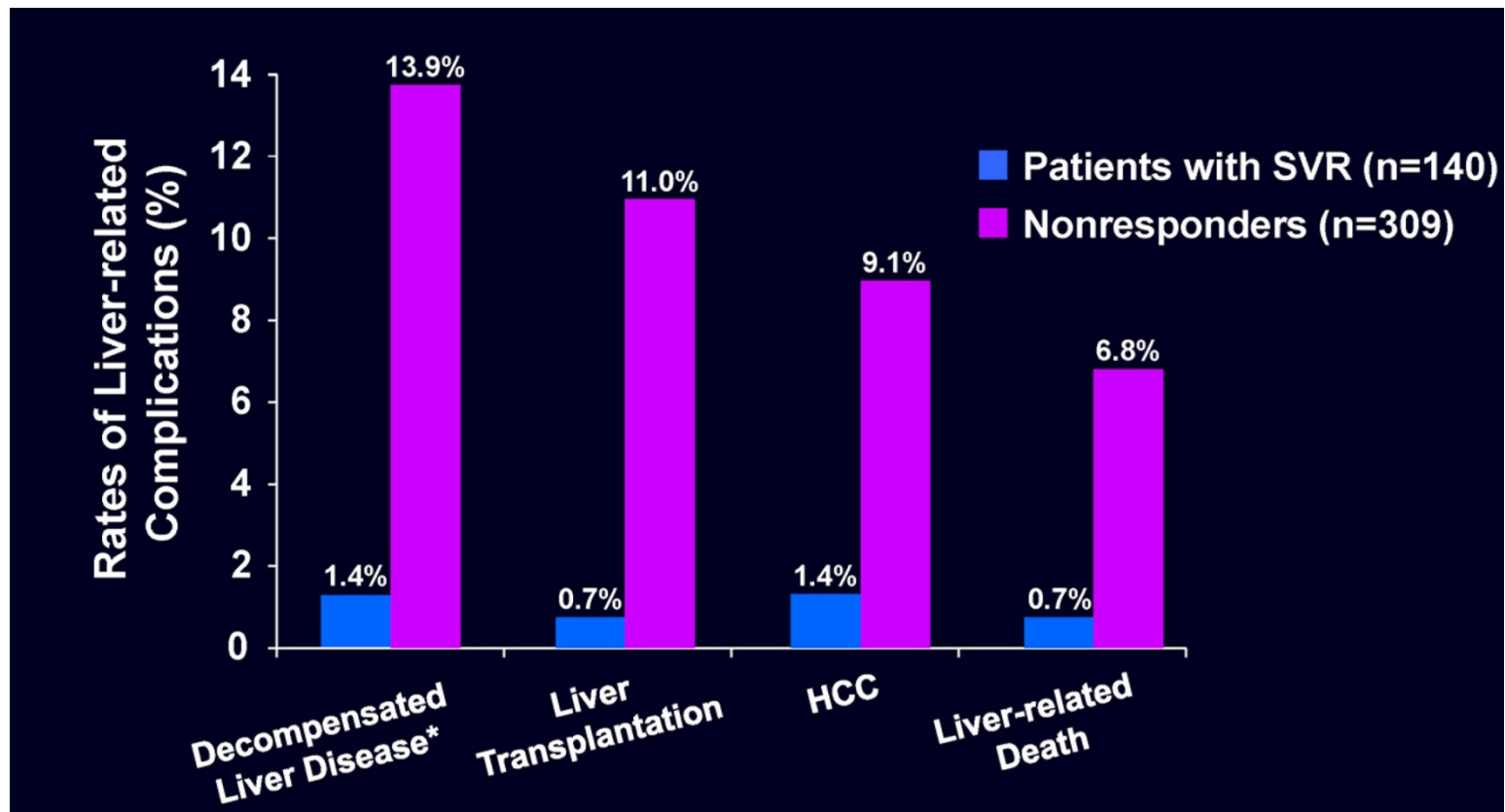


- Improved outcomes in patients who achieve SVR
 - All-cause mortality
 - Liver-related mortality
 - Liver cancer
 - Liver failure or need for liver transplantation
 - Reduce insulin resistance and diabetes mellitus

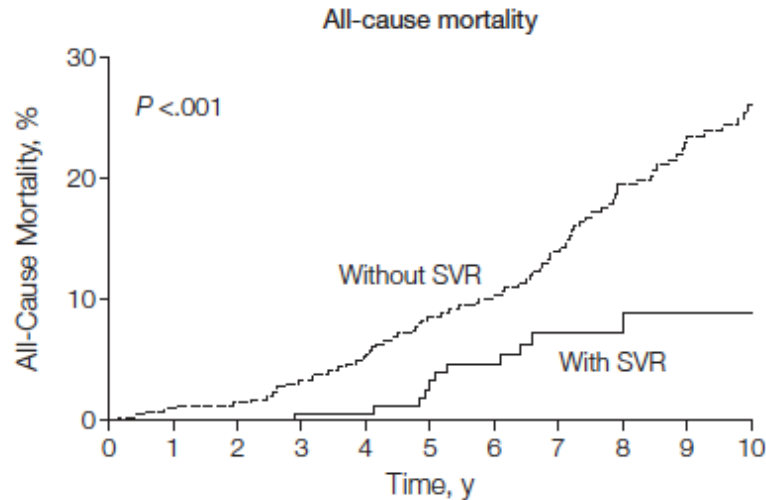
SVR Reduces Liver-Related Complications



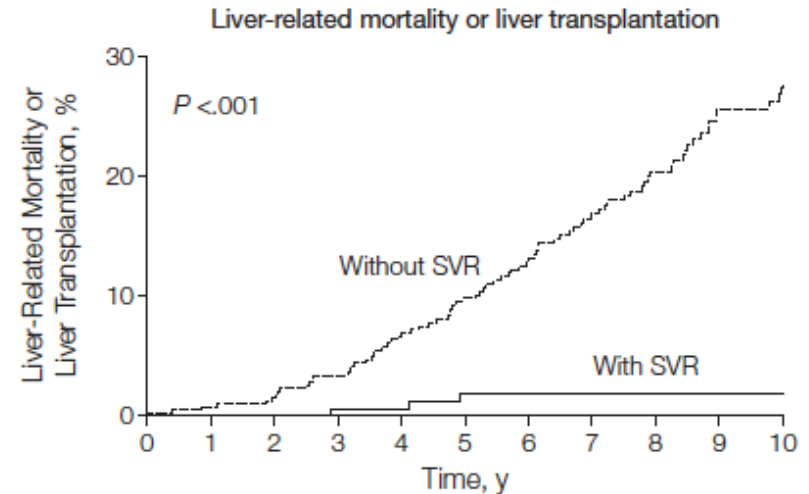
- HALT-C Cohort (patients with bridging fibrosis/cirrhosis)
- Median follow-up 85.8 months (SVR) and 78.4 months (non-SVR)



SVR Reduces All-Cause and Liver-Related Mortality



| No. at risk | |
|-------------|---|
| Without SVR | 405 393 382 363 344 317 295 250 207 164 135 |
| With SVR | 192 181 168 162 155 144 125 88 56 40 28 |



| No. at risk | |
|-------------|---|
| Without SVR | 405 392 380 358 334 305 277 229 187 146 119 |
| With SVR | 192 181 168 162 155 144 125 88 56 40 28 |

| Outcomes | With SVR | | | Without SVR | | | P Value ^b |
|--|-------------|----------------------------------|------------------------------------|-------------|----------------------------------|------------------------------------|----------------------|
| | Events, No. | Observation Period, Person-Years | Rate per 100 Person-Years (95% CI) | Events, No. | Observation Period, Person-Years | Rate per 100 Person-Years (95% CI) | |
| Any event ^a | 18 | 1260 | 1.43 (0.77-2.09) | 169 | 2921 | 5.79 (4.91-6.66) | <.001 |
| All-cause mortality | 13 | 1283 | 1.01 (0.46-1.56) | 100 | 3410 | 2.93 (2.36-3.51) | <.001 |
| Liver-related mortality or liver transplantation | 3 | 1283 | 0.23 (<0.01-0.50) | 103 | 3120 | 3.20 (2.58-3.82) | <.001 |

Baseline Factors Predictive of SVR With an HCV PI Plus PegIFN and RBV therapy



Viral factors

- HCV RNA level
- HCV genotype (GT; 1 vs 2,3)
- GT 1 subtype (1a vs 1b)

Host factors

- Race and ethnicity
- Gender
- Interleukin 28B genotype (CC vs CT or TT)
- Body mass index (BMI)

Combined

- Liver histology

Baseline Predictors of SVR: SPRINT-2



| Effect | Odds Ratio (95% CI) | P value |
|---|---------------------|---------|
| Baseline HCV RNA: ≤ 400,000 vs > 400,000 | 11.6 (1.5, 87.8) | .02 |
| <i>IL28B</i> rs12979860: CC vs TT | 2.6 (1.3, 5.1) | .006 |
| <i>IL28B</i> rs12979860: CC vs CT | 2.1 (1.2, 3.7) | .01 |
| <i>IL28B</i> rs12979860: CT vs TT | 1.2 (0.7, 2.2) | .48 |
| Cirrhosis: no vs yes | 4.3 (1.6, 11.9) | .004 |
| Genotype: 1b vs 1a | 2.0 (1.2, 3.4) | .005 |
| Race: non-black vs black | 2.0 (1.1, 3.7) | .03 |
| BMI: ≤ 30 vs > 30 | 1.6 (1.0, 2.5) | .07 |

CI = confidence interval

Viral Factors that Predict Response to Therapy (HCV Genotype, HCV RNA level)



Viral Factors



- **Genotype**
 - Current therapies (PegIFN/RBV/PI) centered on GT 1
 - Standard of care for GT 2,3: PegIFN and RBV
- **Subtype**
 - SVR rates higher for GT 1b vs 1a
 - 1a - lower barrier to emergence of resistant variants for HCV PIs
- **HCV RNA level**
 - Higher HCV RNA associated with slightly diminished SVR
 - Effect strong

Host Factors that Predict Response to Therapy (Race, Age, Sex, *IL28B* Genotype)

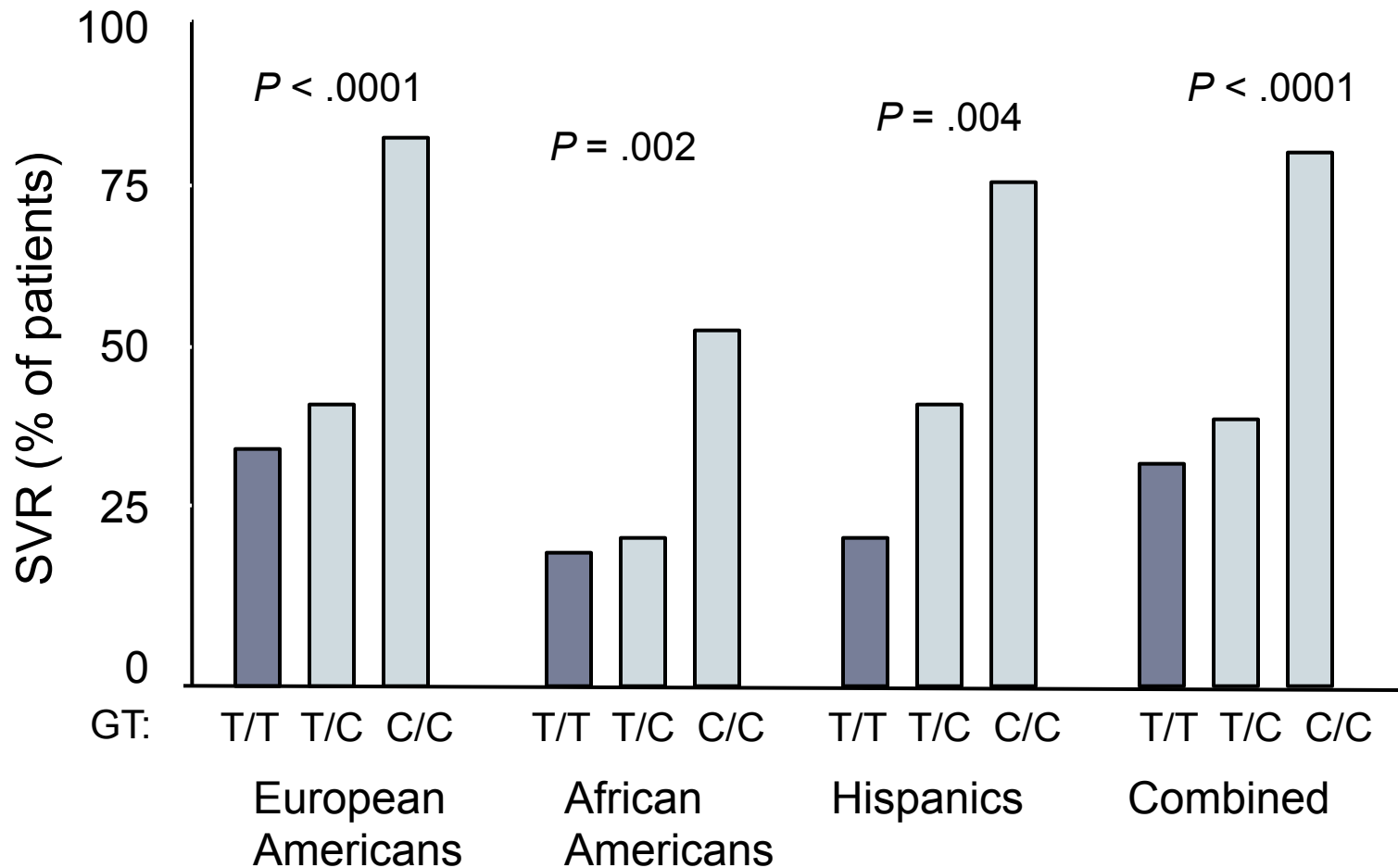


Host Factors

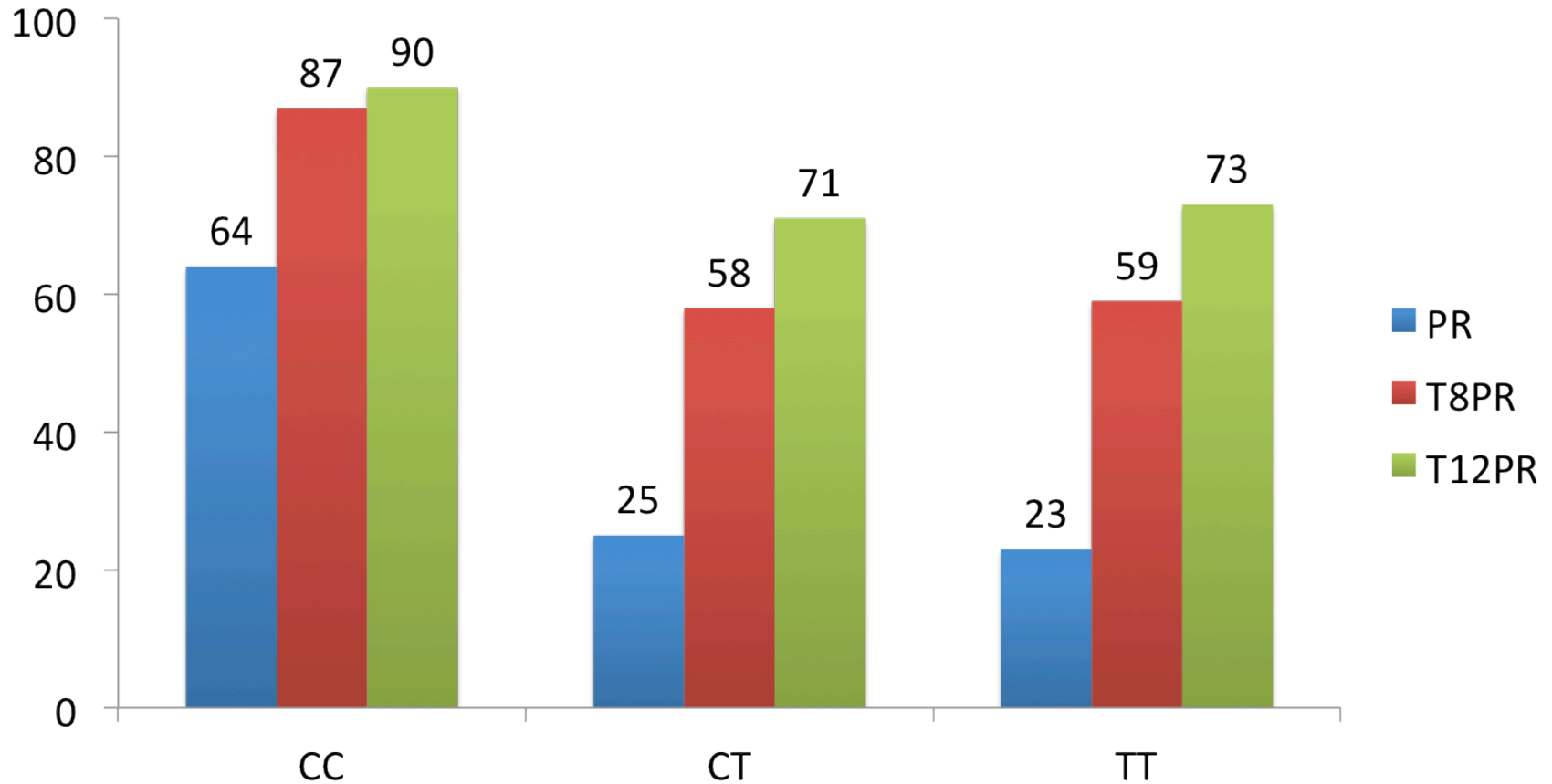


- Race (non-black > black)
- Age (younger > older)
- Sex (F > M)
- *IL28B* GT (CC > CT, TT for the rs12979860 single nucleotide polymorphism [SNP])
 - Strongest predictor of PegIFN and RBV response in GT 1, but effect less stark with PegIFN and RBV plus a PI

SVR to PegIFN and RBV Profoundly Influenced by *IL28B* Genotype



SVR According to *IL28B*: PegIFN, RBV, and Telaprevir



Degree of Fibrosis and Predicting Treatment Response



Combined Factors



- **Extent of fibrosis**
 - Patients without cirrhosis more responsive than patients with cirrhosis
- **Mechanism unclear**
 - Pharmacokinetic and pharmacodynamic considerations with distorted architecture
 - Cirrhosis associated with innate and adaptive immune deficits
- **Decision to treat patients with cirrhosis must be weighed against projected response rate and potential for adverse effects**
- **Similarly, deferral of treatment may be considered in patients without cirrhosis with the impending approval of direct-acting antivirals (DAAs)**

Summary



- Cure of HCV achievable
- SVR associated with significant clinical benefit
 - Slows or reverses disease progression
 - Reduces hepatic and all-cause mortality
 - Improves extrahepatic manifestations
- Host, viral, and mixed factors all influence success of treatment with PegIFN, RBV, and a PI
- Discussion of these factors should take place in preparation for treatment
- Availability of potent DAA combination regimens will dilute the effect of these predictors

End



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