

1 **PI-Based HCV DAAs Are Associated with Increased Risk of Aminotransferase Elevations**  
2 **but Not Hepatic Dysfunction or Decompensation**

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77

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79 **ABSTRACT**

80 Background & Aims: Cases of acute liver injury (ALI) have been reported among chronic  
81 hepatitis C virus-infected initiators of protease inhibitor (PI)-based direct-acting antiviral (DAA)  
82 regimens, predominately with decompensated cirrhosis in whom these therapies are  
83 contraindicated. No analyses have evaluated if initiators of PI versus non-PI-based DAAs have  
84 higher risk of ALI events, stratified by advanced hepatic fibrosis/cirrhosis. We compared the risk  
85 of three ALI outcomes among PI-based and non-PI-based DAA initiators, by baseline FIB-4.

86 Methods: We conducted a cohort study of 18,498 initiators of PI-based DAA therapy  
87 (paritaprevir/ritonavir/ombitasvir +/- dasabuvir, elbasvir/grazoprevir, glecaprevir/pibrentasvir)  
88 matched 1:1 on propensity score to non-PI-based DAA initiators (sofosbuvir/ledipasvir,  
89 sofosbuvir/velpatasvir) in the 1945-1965 Veterans Birth Cohort (2014-2019). During exposure to  
90 DAA therapy, we determined development of: 1) alanine aminotransferase (ALT) >200 U/L, 2)  
91 severe hepatic dysfunction (coagulopathy with hyperbilirubinemia), and 3) hepatic  
92 decompensation. Cox regression was used to determine hazard ratios (HRs) with 95% confidence  
93 intervals of each outcome, stratified by baseline advanced hepatic fibrosis/cirrhosis by FIB-4.

94 Results: Among persons with baseline FIB-4  $\leq 3.25$ , PI initiators had higher risk of ALT >200 U/L  
95 (HR, 3.98 [2.37-6.68]), but not severe hepatic dysfunction (HR, 0.67 [0.19-2.39]) or hepatic  
96 decompensation (HR, 1.01 [0.29-3.48]), compared to non-PI-initiators. For those with baseline  
97 FIB-4 >3.25, PI initiators had higher risk of ALT >200 U/L (HR, 2.15 [1.08-4.29]), but not severe  
98 hepatic dysfunction (HR, 1.23 [0.63-2.40]) or hepatic decompensation (HR, 0.87 [0.42-1.82]),  
99 compared to non-PI initiators.

100 Conclusion: While risk of incident ALT elevations was increased among PI-based DAA initiators  
101 in both FIB-4 strata, risk of severe hepatic dysfunction and hepatic decompensation did not  
102 differ between PI and non-PI-based DAA initiators in either FIB-4 stratum.

103 Highlights

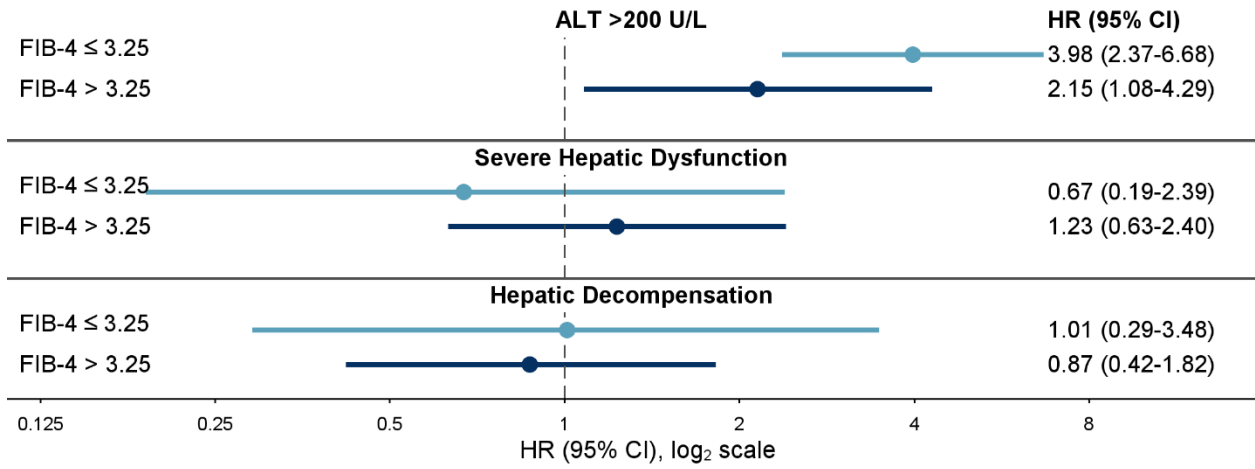
- 104 • Comparative analysis of 18,498 initiators of PI-based DAAs matched on propensity  
105 score to 18,498 initiators of non-PI-based DAAs to assess risk of 3 acute liver injury  
106 endpoints, according to advanced hepatic fibrosis/cirrhosis status by FIB-4.
- 107 • Propensity score-matched hazard ratios of ALT >200 U/L were higher for PI than non-PI-  
108 initiators in those with and without baseline advanced hepatic fibrosis/cirrhosis (i.e., FIB-  
109 4 >3.25 and FIB-4 ≤3.25, respectively).
- 110 • No differences in propensity score-matched hazard ratios of severe hepatic dysfunction  
111 or hepatic decompensation were observed between PI and non-PI-based DAA initiators,  
112 regardless of baseline advanced hepatic fibrosis/cirrhosis status by FIB-4.

113

114 Lay Summary: Cases of liver injury have been reported among patients treated with protease  
115 inhibitor-based direct-acting antivirals for hepatitis C infection, but it is not clear if risk of liver  
116 injury among people starting these drugs is increased compared to those starting non-protease  
117 inhibitor-based therapy. In this study, persons who initiated protease inhibitor-based treatment  
118 had higher risk of liver inflammation than non-protease inhibitor-based initiators, regardless of  
119 the presence of pre-treatment advanced liver fibrosis/cirrhosis. However, the risk of severe liver  
120 dysfunction and decompensation were not higher for protease inhibitor-based initiators.

121

122 **Graphical Abstract. Hazard ratios (HRs) with 95% confidence intervals (CIs) of specified**  
 123 **acute liver injury outcomes for propensity score matched cohorts of protease inhibitor**  
 124 **and non-protease inhibitor-based direct-acting antiviral therapy initiators, by baseline**  
 125 **advanced hepatic fibrosis/cirrhosis status by FIB-4.**



126

127

128 **INTRODUCTION**

129           The increased efficacy of direct-acting antivirals (DAAs) compared to interferon-based  
130 regimens has revolutionized the treatment of chronic hepatitis C virus (HCV) infection.<sup>1</sup> While DAAs  
131 also have a superior safety profile over interferon-based therapy, post-marketing surveillance has  
132 identified acute liver injury (ALI) as a potentially important DAA-related toxicity. On October 22,  
133 2015, the United States (US) Food and Drug Administration (FDA) released a Drug Safety  
134 Communication reporting cases of hepatic decompensation that developed among chronic HCV-  
135 infected patients with compensated cirrhosis during treatment with paritaprevir/ritonavir/ombitasvir  
136 (PRO), either alone or with dasabuvir (PROD).<sup>2</sup> A follow-up FDA communication on August 28,  
137 2019 reported cases of hepatic dysfunction among chronic HCV-infected patients treated with  
138 glecaprevir/pibrentasvir, elbasvir/grazoprevir, and sofosbuvir/velpatasvir/voxilaprevir.<sup>3</sup> Notably, all of  
139 the DAA regimens implicated in these FDA reports included an HCV protease inhibitor (PI). In many  
140 of the reports, ALI occurred among patients who had moderate-to-severe liver impairment (i.e.,  
141 Child-Pugh Class B and C), in whom these drugs are contraindicated.<sup>4,5</sup>

142           As a result of these reports, there have been major concerns that PI-based DAA therapy  
143 might be associated with an increased risk of ALI, particularly among persons with advanced  
144 hepatic fibrosis/cirrhosis, compared to non-PI-based treatment.<sup>6</sup> Chronic HCV-induced  
145 advanced liver fibrosis might impair cytochrome P450 activity.<sup>4</sup> This impaired activity could  
146 result in elevated serum PI concentrations during PI-based DAA treatment, which might  
147 precipitate an ALI event, particularly significant liver aminotransferase elevations, severe  
148 hepatic dysfunction (i.e., coagulopathy plus hyperbilirubinemia), or hepatic decompensation.  
149 However, no studies have examined whether PI-based DAA treatment is associated with higher  
150 risk of ALI compared to non-PI-based therapy. Moreover, it is unclear if the risk of ALI  
151 associated with PI-based DAA therapy is heightened among those with advanced hepatic  
152 fibrosis/cirrhosis. These data are needed to determine the real-world comparative hepatic safety

153 of DAAs among chronic HCV-infected patients, especially those with advanced hepatic  
154 fibrosis/cirrhosis.

155 To address these critical knowledge gaps, we evaluated the incidence and risk of ALL,  
156 defined by incident development of liver aminotransferase elevations, severe hepatic  
157 dysfunction, or hepatic decompensation, among chronic HCV-infected patients who newly  
158 initiated a PI-based compared to non-PI-based DAA regimen. Given the potential for advanced  
159 hepatic fibrosis to impair the metabolism of PI-based DAAs, we stratified our results according  
160 to baseline stage of hepatic fibrosis using the Fibrosis-4 Index for Hepatic Fibrosis (FIB-4), a  
161 non-invasive measure of advanced hepatic fibrosis/cirrhosis that has been validated compared  
162 to liver biopsy among chronic HCV-infected patients.<sup>7</sup>

163

## 164 **PATIENTS AND METHODS**

### 165 **Study Design and Data Source**

166 We conducted a retrospective cohort study among chronic HCV-infected patients who  
167 initiated DAA treatment between January 1, 2014 and June 30, 2019 within the US Department  
168 of Veterans Affairs (VA) using data from the 1945-1965 Veterans Birth Cohort (VBC).<sup>8,9</sup> The  
169 VBC consists of electronic health record data from all Veterans born between 1945 and 1965  
170 who received any VA care since October 1, 1999, encompassing >6.6 million persons aged 54-  
171 75 years. We chose to use data from the VBC since persons born between 1945 and 1965 have  
172 a 6-fold higher prevalence of HCV infection compared to all other age groups.<sup>10</sup> Available data  
173 in the VBC include demographics, inpatient and outpatient diagnoses, laboratory results, and  
174 dispensed medications. Date of death is available from the VA Vital Status file.<sup>11</sup> The study was  
175 approved by the Institutional Review Boards of the University of Pennsylvania, VA Connecticut  
176 Healthcare System, and Yale University, and was conducted under a waiver of informed  
177 consent per 45 CFR §46.117(c).

178



179 **Study Patients**

180 Chronic HCV-infected patients were eligible if they: 1) newly initiated a DAA of interest  
181 (i.e., sofosbuvir/ledipasvir, elbasvir/grazoprevir, sofosbuvir/velpatasvir, glecaprevir/pibrentasvir,  
182 or PRO/PROD) within the VA between January 1, 2014 and June 30, 2019, and 2) were in care  
183 in the VA for  $\geq 2$  years prior to DAA initiation (to permit capture of relevant baseline  
184 comorbidities, laboratory results, and medications). DAA prescriptions in the VA have been  
185 validated to accurately reflect patients receiving treatment for chronic HCV.<sup>12</sup> While PRO/PROD  
186 are no longer recommended DAA regimens, we included initiators of these drugs since they  
187 were commonly dispensed PI-based regimens during the period of interest and would provide  
188 additional evidence on the hepatotoxicity of PI-based DAAs.

189 We defined the index date as the date that the DAA of interest was initially dispensed in  
190 the VA on or after January 1, 2014. The 2 years prior to the index date represented the baseline  
191 period, during which baseline comorbidities and laboratory results were collected. Patients were  
192 excluded if during the baseline period they were: 1) diagnosed with HIV infection (since such  
193 patients may be on antiretroviral drugs that could increase the risk of ALI<sup>13</sup>); 2) dispensed  
194 warfarin or a direct-acting oral anticoagulant (i.e., apixaban, dabigatran, edoxaban,  
195 rivaroxaban), which would prevent identification of coagulopathy due to ALI; 3) identified with  
196 any prevalent ALI outcome (defined below), since we sought to ascertain incident events; 4)  
197 diagnosed with hepatocellular carcinoma (which could lead to misclassification of ALI); or 5)  
198 were missing all baseline laboratory results necessary to calculate FIB-4. We also excluded  
199 patients who were ever hepatitis B surface antigen-positive (to reduce the likelihood of detecting  
200 ALI due to hepatitis B virus reactivation<sup>14</sup>) or were dispensed a DAA within the VA at any time  
201 prior to the index date (since we wished to restrict the sample to new DAA initiators). We chose  
202 not to evaluate rates of ALI events among DAA initiators who had decompensated cirrhosis,  
203 since PI-based DAA regimens are contraindicated in this group.

204 Follow-up continued until: 1) study endpoint, 2) death, 3) switch to a different DAA, 4)

205 discontinuation of DAA (defined as no further fills within 30 days after the last prescription's  
206 supply), 5) dispensation of warfarin or a direct-acting oral anticoagulant, or 6) September 30,  
207 2019, whichever occurred first. For patients who completed or discontinued DAA therapy, we  
208 included 30 additional days of exposure time after the last days' supply to ensure capture of  
209 hepatotoxic events potentially related to DAA use.

210

### 211 **Study Outcomes**

212 To determine the full spectrum of ALI events associated with PI-based DAAs, we  
213 examined 3 incident ALI outcomes. First, we evaluated incident liver aminotransferase  
214 elevations, defined as an inpatient or outpatient alanine aminotransferase (ALT) >200 U/L  
215 (approximately 5 times the upper limit of normal of the assays used), a threshold that represents  
216 clinically important hepatic injury,<sup>15</sup> and approximately 10 times what has been considered  
217 normal liver aminotransferase levels for females (19 U/L) and males (30 U/L).<sup>16</sup> As a secondary  
218 endpoint, we evaluated development of inpatient or outpatient ALT >400 U/L, consistent with  
219 the definition of grade 4 ALT elevations employed in clinical trials.<sup>17</sup>

220 Second, we evaluated severe hepatic dysfunction, defined by an inpatient or outpatient  
221 international normalized ratio (INR)  $\geq 1.5$  and total bilirubin >2 times the upper limit of normal  
222 within up to 30 days of each other. This definition of severe hepatic dysfunction has been used  
223 by the US FDA's Sentinel System to assess serious and clinically significant drug-induced ALI in  
224 the post-marketing period.<sup>18</sup> While Hy's Law has also been used by Sentinel to identify ALI, we  
225 selected the definition that identifies liver injury at an advanced stage, such that serum liver  
226 aminotransferases might not be sufficiently elevated to meet Hy's Law.<sup>18</sup> If the laboratory  
227 abnormalities presented on different dates, the event was considered to have occurred on the  
228 date that the latter abnormality occurred.

229 Third, we determined incident hepatic decompensation, defined by 1 hospital discharge  
230 diagnosis (principal or contributory) or 2 or more outpatient diagnoses (recorded within 1 year)

231 of ascites, spontaneous bacterial peritonitis, esophageal variceal hemorrhage, or hepatic  
232 encephalopathy (**Supplementary Table 1**).<sup>19</sup> The decompensation date was defined as the  
233 hospital discharge date (if event was identified by hospital diagnosis) or initial outpatient  
234 diagnosis date (if identified by outpatient diagnoses).

235

## 236 **Data Collection**

237 Baseline clinical data included age, sex, race/ethnicity, body mass index, diabetes  
238 mellitus (defined by random glucose  $\geq 200$  mg/dL, hemoglobin A1c  $\geq 6.5\%$ , and/or anti-diabetic  
239 medication use),<sup>20</sup> previously validated diagnoses of alcohol dependence/abuse,<sup>21</sup> and use of  
240 ribavirin as part of the DAA regimen. Baseline laboratory data included HCV RNA, HCV  
241 genotype, ALT, aspartate aminotransferase (AST), INR, total bilirubin, hemoglobin, platelets,  
242 and serum creatinine. Baseline FIB-4 was calculated by: (age [years] x AST [U/L]) / (platelet  
243 count [ $10^9$ /L] x (ALT [U/L])<sup>1/2</sup>).<sup>7</sup> FIB-4  $> 3.25$  identifies advanced hepatic fibrosis/cirrhosis  
244 (METAVIR stages F3 or F4) with a high degree of accuracy versus liver biopsy in chronic HCV  
245 infection.<sup>7</sup> When multiple laboratory results were assessed during the baseline period, we  
246 collected the result closest, but prior, to the index date. The estimated glomerular filtration rate  
247 (mL/min/1.73 m<sup>2</sup>) was calculated using the Modification of Diet in Renal Disease equation:  $175 \times$   
248 (serum creatinine)<sup>-1.154</sup> x (age)<sup>-0.203</sup> x (0.742, if female) x (1.212, if Black).<sup>22</sup> The Model for End-  
249 Stage Liver Disease (MELD) score was calculated by:  $3.78 \times \ln[\text{total bilirubin (mg/dL)}] +$   
250  $11.2 \times \ln[\text{INR}] + 9.57 \times \ln[\text{creatinine (mg/dL)}] + 6.43$ .<sup>23</sup>

251 Data collected during follow-up included: outpatient and inpatient ALT, INR, and total  
252 bilirubin results; diagnoses of hepatic decompensation; and sustained virologic response (SVR12;  
253 defined by undetectable HCV RNA on the first test  $\geq 12$  weeks after DAA treatment end date).

254

255           **Statistical Analysis**

256           To reduce the potential for selection bias in examining rates of ALI events between PI-  
257 based DAA therapy (glecaprevir/pibrentasvir, elbasvir/grazoprevir, or PRO/PROD) and non-PI-  
258 based regimens (sofosbuvir/ledipasvir or sofosbuvir/velpatasvir), we developed propensity  
259 scores, which determine each patient’s probability of being assigned to a particular treatment  
260 given their observed set of baselined covariates.<sup>24</sup> Propensity score methods allow for the  
261 reduction of bias when estimating treatment effects by accounting for the differential probability  
262 of receiving PI-based or non-PI-based DAA therapy. The propensity score model was developed  
263 using logistic regression, with potential determinants of PI-based DAA therapy as independent  
264 variables and PI-based DAA treatment exposure as the dependent variable.<sup>25</sup> Variables selected  
265 for the propensity score were those that might affect clinicians’ decision to prescribe a PI or non-  
266 PI DAA regimen and influence risk of acute liver injury and included: age, sex, race/ethnicity,  
267 body mass index, diabetes, diagnosis of alcohol dependence/abuse, HCV genotype,  
268 hemoglobin, platelet count, ALT, AST, INR, total bilirubin, eGFR, MELD score, ribavirin use, and  
269 date of DAA initiation. Within FIB-4 strata, each PI initiator was matched on propensity score  
270 (nearest-neighbor matching within 0.02 of the propensity score) to one non-PI initiator. This  
271 matching allows us to create a comparison group of non-PI-treated patients whose baseline  
272 characteristics resemble those of PI-treated patients.<sup>25</sup> We compared the baseline characteristics  
273 between PI and non-PI initiators prior to and after propensity score matching using standardized  
274 differences, of which a value exceeding 0.1 is generally considered meaningful.<sup>26</sup>

275           For propensity score-matched PI and non-PI initiator cohorts, we determined incidence  
276 rates (events per 1,000 person-years) of each ALI outcome (as independent events) with 95%  
277 confidence intervals (CIs), stratified by baseline advanced hepatic fibrosis/cirrhosis status by  
278 FIB-4 ( $\leq 3.25$  versus  $> 3.25$ ). Additionally, among PI-based and non-PI-based DAA initiators who  
279 had an ALI event defined by ALT  $> 200$  U/L, we evaluated the median ALT level within each 4-  
280 week period of treatment over 32 weeks of follow-up, by FIB-4. For individuals with multiple ALT

281 results within a given 4-week period, we analyzed the highest assessed ALT. We then  
282 determined the proportion whose ALT decreased to  $\leq 100$  U/L. To assess if development of ALT  
283  $> 200$  U/L compromised likelihood of achieving HCV cure, we compared the proportions that  
284 achieved SVR12 for persons who did and did not develop ALT  $> 200$  U/L, by FIB-4 status, among  
285 those tested for SVR12.

286 Cox regression was then used to determine the hazard ratio (HR) of each ALI outcome  
287 associated with PI-based DAA therapy compared to use of non-PI-based regimens.<sup>27</sup> We  
288 confirmed the adequacy of the propensity score as a continuous variable by: 1) observing  
289 overlap in the distribution between PI-based and non-PI-based DAA users (**Supplementary**  
290 **Figs. 1 and 2**), and 2) confirming linearity of propensity score categories within outcome models.  
291 Results were stratified by baseline FIB-4 ( $\leq 3.25$ ;  $> 3.25$ ). Proportionality of hazards was assessed  
292 by log-log plots and Schoenfeld residuals.<sup>27</sup> Data were analyzed using SAS 9.4 (SAS Institute  
293 Inc., Cary, NC).

294

## 295 **RESULTS**

### 296 **Patient Characteristics**

297 We identified 96,720 chronic-HCV-infected persons who were dispensed one of the  
298 DAAs of interest between January 1, 2014 and June 30, 2019. After exclusions, 20,169 new  
299 initiators of a PI-based DAA regimen (5,994 PRO/PROD; 8,301 elbasvir/grazoprevir; 5,874  
300 glecaprevir/pibrentasvir) and 51,222 non-PI-based initiators (43,813 sofosbuvir/ledipasvir; 7,409  
301 sofosbuvir/velpatasvir) remained (**Fig. 1**).

302 Prior to propensity score matching, initiators of PI-based DAA regimens more commonly  
303 were Black, infected with HCV genotype 1, and had diabetes mellitus, severe anemia  
304 (hemoglobin  $< 10$  g/dL), and renal insufficiency (estimated glomerular filtration rate  $< 30$   
305 mL/min/1.73 m<sup>2</sup>) (**Tables 1 and 2**). Initiators of non-PI-based DAAs more commonly had alcohol  
306 dependence/abuse history and MELD score  $< 10$ . Among those with baseline FIB-4  $\leq 3.25$ ,

307 14,985 initiators of PI-based DAAs were propensity score-matched to 14,985 initiators of non-  
308 PI-based DAAs. Among those with baseline FIB-4 >3.25, 3,513 initiators of PI-based DAAs  
309 were propensity score-matched to 3,513 initiators of non-PI-based DAAs. Propensity score-  
310 matching generally balanced the frequencies of characteristics between the cohorts.

311

### 312 **Incidence Rates of ALI Events, by Baseline FIB-4**

313 **Table 3** reports the absolute risk and unadjusted incidence rates of ALT >200 U/L,  
314 severe hepatic dysfunction, and hepatic decompensation for the propensity score-matched PI-  
315 based and non-PI-based initiator cohorts by baseline FIB-4. **Supplementary Table 2** reports  
316 the frequencies of specific decompensation diagnoses. **Supplementary Table 3** reports the  
317 absolute risk and unadjusted rates of ALI outcomes in the overall study sample prior to  
318 propensity score matching, by baseline FIB-4. Regardless of baseline FIB-4 score, the absolute  
319 risk of each ALI outcome was rare (<2%).

320 Among persons with baseline FIB-4  $\leq$ 3.25, incidence rates of ALT >200 U/L (**Table 3**)  
321 and ALT >400 U/L (**Supplementary Table 4**) were higher in magnitude for PI than non-PI  
322 initiators. Among PI and non-PI initiators who developed ALT >200 U/L, median ALT levels  
323 peaked at week 4 of treatment for PI-based initiators and at week 8 for non-PI-based initiators  
324 and then declined over follow-up (**Fig. 2A**). Similar proportions of PI-based and non-PI-based  
325 DAA initiators who developed ALT >200 U/L experienced subsequent decrease in ALT to  $\leq$ 100  
326 U/L (90.1% versus 88.9%, respectively;  $p=0.88$ ). The proportion of DAA initiators achieving  
327 SVR12 did not differ between those who did and did not develop ALT >200 U/L (**Supplementary**  
328 **Table 5**). Incidence rates of severe hepatic dysfunction and hepatic decompensation were  
329 similar between PI- and non-PI-based DAA initiators for those with baseline FIB-4  $\leq$ 3.25.

330 For patients with baseline FIB-4 >3.25 (advanced hepatic fibrosis/cirrhosis), rates of both  
331 ALT >200 U/L (**Table 3**) and ALT >400 U/L (**Supplementary Table 4**) were higher in magnitude  
332 for PI-based initiators. Among patients who had ALT >200 U/L, median ALT levels peaked at

333 week 4 for PI-based and non-PI initiators, and then declined over follow-up (**Fig. 2B**). Similar  
334 proportions of PI-based and non-PI-based DAA initiators who developed ALT >200 U/L  
335 experienced subsequent decrease in ALT to  $\leq 100$  U/L (92.0% vs 100%, respectively;  $p=0.31$ ).  
336 There were no significant differences in achievement of SVR12 between those who did versus  
337 did not develop ALT >200 U/L (**Supplementary Table 5**). In contrast to findings among those  
338 with FIB-4  $\leq 3.25$ , incidence rates of severe hepatic dysfunction and hepatic decompensation  
339 were higher in magnitude for initiators of non-PI regimens.

340

#### 341 **Risk of ALI Events with PI-Based Versus Non-PI-Based DAAs, by Baseline FIB-4**

342 Among persons with baseline FIB-4  $\leq 3.25$ , initiators of PI-based DAA regimens had  
343 higher relative hazards of ALT >200 U/L (HR, 3.98 [95% CI, 2.37-6.68]; **Fig. 3**) and ALT >400  
344 U/L (HR, 3.02 [95% CI, 1.10-8.30]) than those who received a non-PI-based regimen. However,  
345 among persons in this FIB-4 stratum, PI initiators did not have significantly higher relative  
346 hazards of either severe hepatic dysfunction (HR, 0.67 [95% CI, 0.19-2.39]) or hepatic  
347 decompensation (HR, 1.01 [95% CI, 0.29-3.48]) than non-PI initiators.

348 For those with baseline FIB-4 >3.25, initiators of PI-based DAAs had significantly higher  
349 relative hazards of ALT >200 U/L (HR, 2.15 [95% CI, 1.08-4.29]), but not of ALT >400 U/L (HR,  
350 1.52 [95% CI, 0.25-9.11]), severe hepatic dysfunction (HR, 1.23 [95% CI, 0.63-2.40]), or hepatic  
351 decompensation (HR, 0.87 [95% CI, 0.42-1.82]) compared to non-PI initiators (**Fig. 3**).

352

## 353 **DISCUSSION**

354 In this national sample of treatment-naïve chronic HCV-infected patients within the VA  
355 system, we identified PI-based DAA initiators (glecaprevir/pibrentasvir, elbasvir/grazoprevir, or  
356 PRO/PROD) and matched them 1:1 on propensity scores to non-PI-based initiators  
357 (sofosbuvir/ledipasvir or sofosbuvir/velpatasvir) to ensure that the cohorts were similar with  
358 regards to the frequencies of important baseline demographic and clinical characteristics. We

359 observed that the absolute risk of the three ALI outcomes of interest was low (<2%) among  
360 initiators of both PI-based and non-PI-based DAAs. Regardless of baseline advanced hepatic  
361 fibrosis/cirrhosis status by FIB-4, incidence rates and relative hazards of ALT >200 U/L were  
362 higher for PI- than non-PI-based initiators. However, relative hazards of severe hepatic  
363 dysfunction and hepatic decompensation were not significantly increased among users of PI-  
364 based DAA regimens, regardless of baseline FIB-4.

365 We observed that PI-based DAA therapy was associated with higher risk of liver  
366 aminotransferase elevations, but not severe hepatic dysfunction or hepatic decompensation,  
367 compared to non-PI therapy. Transient elevations in liver aminotransferases have been reported  
368 following initiation of PI-based DAA regimens, particularly PRO/PROD and  
369 elbasvir/grazaprevir.<sup>28-30</sup> The biologic mechanism remains unclear but may be due to either  
370 immune-mediated hepatocyte injury in the setting of viral clearance or idiosyncratic drug-  
371 induced ALI.<sup>31,32</sup> These reports have suggested that ALT elevations during PI-based DAA  
372 therapy are largely asymptomatic and that levels normalize by week 8.<sup>28,29</sup> Consistent with those  
373 findings, we observed that among PI-based and non-PI-based DAA initiators who developed an  
374 ALT >200 U/L, median ALT levels generally decreased after 4 weeks following the initial event.  
375 These findings suggest that the majority of ALT elevations during DAA therapy are transient.  
376 Furthermore, our findings suggest that the ALT elevations did not decrease the likelihood of  
377 achieving SVR12.

378 In our primary analysis, we found no association between PI therapy and severe hepatic  
379 dysfunction or hepatic decompensation, regardless of baseline advanced hepatic  
380 fibrosis/cirrhosis status by FIB-4. This is a valuable observation particularly for patients with  
381 baseline advanced hepatic fibrosis/cirrhosis, since compensated cirrhosis may predispose to  
382 increased PI exposure and subsequently an increased risk of hepatotoxicity.<sup>33-35</sup> Our findings  
383 suggest that the risk of serious ALI events is not increased among chronic HCV-infected persons  
384 without decompensated cirrhosis who initiated PI-based versus non-PI-based DAA therapy.



385 Prior studies evaluating the real-world safety of DAAs estimated that the absolute risk of  
386 hepatic decompensation following DAA initiation was 0.2-1.1%, similar to our findings.<sup>34,36-38</sup>  
387 One multicenter observational study of 33,808 initiators of DAAs from 2012-2017 reported the  
388 incidence of hepatic decompensation to be 23.8 events/1,000 person-years.<sup>37</sup> However, rates  
389 were not stratified by cirrhosis status, as in this study. A previous study among US Veterans  
390 observed more than 10-fold higher incidence rates of hepatic decompensation among initiators  
391 of PRO/PROD and sofosbuvir/ledipasvir with cirrhosis compared to those without cirrhosis.<sup>34</sup> In  
392 this study, we found that rates of each ALI outcome were substantially higher among persons  
393 with FIB-4 >3.25, highlighting how cirrhosis might modify the risk of ALI associated with DAA  
394 therapy.

395 Our study had several limitations. First, we undertook these analyses from an  
396 epidemiological standpoint, comparing the relative incidences and risk of ALI events according to  
397 PI-based DAA status in order to identify hepatotoxicity signals. However, we were unable to  
398 ascertain the etiology of each ALI event given the challenges in confirming a drug-induced  
399 etiology of ALI in clinical practice. Second, there is potential for confounding by indication,<sup>39</sup> since  
400 patients were assigned to PI-based DAA treatment by clinician choice. Our implementation of  
401 propensity score matching attempted to account for this in our risk models by matching patients  
402 on the probability of receipt of PI-based versus non-PI-based DAA therapy to create comparable  
403 cohorts for analysis. Third, we were unable to capture alcohol use that might have begun during  
404 DAA therapy as well as concomitant use of hepatotoxic medications during treatment, and these  
405 factors might have contributed to ALI events. Fourth, since ALT was assessed as part of routine  
406 clinical care and not per standardized protocol, our secondary analysis examining median ALT  
407 over 4-week periods of DAA treatment may not have accurately assessed the course of ALT  
408 increase over time. Finally, our study sample was predominantly comprised of male US Veterans  
409 and may not be generalizable to women. Our study is also not generalizable to patients with  
410 decompensated cirrhosis prior to DAA treatment.

411 Our study had a number of strengths. We included a large, national cohort of patients  
412 who initiated different DAA therapies. We evaluated a spectrum of clinically relevant ALI  
413 outcomes and stratified results by baseline FIB-4 to assess the risk of these events by  
414 advanced hepatic fibrosis/cirrhosis status. Finally, we used propensity scores to account for  
415 important variables that might influence prescription of PI versus non-PI-based DAA therapy  
416 and which might be associated with ALI.

417 In conclusion, our study found that PI-based DAA therapy was associated with higher  
418 risk of liver aminotransferase elevations, but not severe hepatic dysfunction or hepatic  
419 decompensation events, compared to non-PI therapy. Liver aminotransferase elevations during  
420 PI-based DAA therapy might be due to immune-mediated inflammation accompanying viral  
421 eradication or transient drug-induced ALI; however, clinically apparent acute severe hepatic  
422 dysfunction or hepatic decompensation were not more common among PI-based DAA initiators.  
423 These findings demonstrate the comparable hepatic safety of PI-based and non-PI-based DAA  
424 therapies among chronic HCV-infected persons without decompensated cirrhosis.

425  
426 **Abbreviations:** ALI, acute liver injury; ALT, alanine aminotransferase; CI, confidence interval;  
427 DAA, direct-acting; antivirals; FDA, Food and Drug Administration; FIB-4, Fibrosis-4 Index for  
428 Hepatic Fibrosis; HCV, hepatitis C virus; HIV, human immunodeficiency virus; HR, hazard ratio;  
429 INR, international normalized ratio; PI, protease inhibitor; PRO, paritaprevir/ritonavir/ombitasvir;  
430 PROD, paritaprevir/ritonavir/ombitasvir/dasabuvir; RNA, ribonucleic acid; US, United States; VA,  
431 Veterans Administration; VBC, Veterans Birth Cohort

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537

**Table 1. Baseline characteristics of patients with FIB-4  $\leq$ 3.25 prescribed protease inhibitor-based and non-protease inhibitor (PI)-based direct-acting antiviral regimens of interest for chronic hepatitis C virus infection prior to and after propensity score matching.**

Characteristics	Prior to Propensity Score Matching			After Propensity Score Matching		
	PI-Based DAA Regimen* (n=16,353)	Non-PI-Based DAA Regimen† (n=40,639)	SDP	PI-Based DAA Regimen* (n=14,985)	Non-PI-Based DAA Regimen† (n=14,985)	SDP
<b>Age</b>			0.17			0.02
<55 years	611 (3.7%)	2,290 (5.6%)		584 (3.9%)	552 (3.7%)	
55-59 years	3,430 (21.0%)	9,994 (24.6%)		3,233 (21.6%)	3,255 (21.7%)	
60-64 years	6,259 (38.3%)	15,592 (38.4%)		5,751 (38.4%)	5,834 (38.9%)	
65-69 years	5,114 (31.3%)	11,428 (28.1%)		4,610 (30.8%)	4,585 (30.6%)	
$\geq$ 70 years	939 (5.7%)	1,335 (3.3%)		807 (5.4%)	759 (5.1%)	
<b>Male sex</b>	15,848 (96.9%)	39,245 (96.6%)	0.02	14,509 (96.8%)	14,487 (96.7%)	<0.01
<b>Race/ethnicity</b>			0.17			0.05
Black	8,140 (49.8%)	17,153 (42.2%)		7,187 (48.0%)	7,096 (47.4%)	
White	6,665 (40.8%)	19,733 (48.6%)		6,359 (42.4%)	6,587 (44.0%)	
Hispanic	831 (5.1%)	1,822 (4.5%)		769 (5.1%)	617 (4.1%)	
Other/Unknown	717 (4.4%)	1,931 (4.8%)		670 (4.5%)	685 (4.6%)	
<b>Body mass index</b>			0.27			0.06
Underweight (<18.50 kg/m <sup>2</sup> )	291 (1.8%)	680 (1.7%)		267 (1.8%)	267 (1.8%)	
Normal (18.50-24.99 kg/m <sup>2</sup> )	4,682 (28.6%)	11,872 (29.2%)		4,263 (28.4%)	4,343 (29.0%)	
Overweight (25.00-29.99 kg/m <sup>2</sup> )	5,679 (34.7%)	14,985 (36.9%)		5,221 (34.8%)	5,303 (35.4%)	
Obesity (30.00-34.99 kg/m <sup>2</sup> )	2,917 (17.8%)	8,096 (19.9%)		2,716 (18.1%)	2,757 (18.4%)	
Morbid obesity ( $\geq$ 35.00 kg/m <sup>2</sup> )	1,215 (7.4%)	3,684 (9.1%)		1,118 (7.5%)	1,146 (7.6%)	
Unknown	1,569 (9.6%)	1,322 (3.3%)		1,400 (9.3%)	1,169 (7.8%)	
<b>Diabetes mellitus</b>	5,259 (32.2%)	11,402 (28.1%)	0.09	4,439 (29.6%)	4,425 (29.5%)	<0.01
<b>Alcohol dependence/abuse diagnosis</b>	8,430 (51.6%)	21,501 (52.9%)	0.03	7,795 (52.0%)	7,822 (52.2%)	<0.01
<b>HCV RNA &gt;800,000 IU/mL</b>	11,220 (68.6%)	28,285 (69.6%)	0.08	10,473 (69.9%)	10,014 (66.8%)	0.09
<b>HCV genotype</b>			0.49			0.06
Genotype 1a	7,553 (46.2%)	25,057 (61.7%)		7,006 (46.8%)	6,781 (45.3%)	
Genotype 1b	6,123 (37.4%)	7,053 (17.4%)		5,467 (36.5%)	5,759 (38.4%)	
Genotype 1, subtype unknown	261 (1.6%)	980 (2.4%)		253 (1.7%)	231 (1.5%)	
Genotype 2	702 (4.3%)	2,714 (6.7%)		664 (4.4%)	618 (4.1%)	
Genotype 3	330 (2.0%)	1,701 (4.2%)		318 (2.1%)	376 (2.5%)	
Other genotype	906 (5.5%)	2,424 (6.0%)		839 (5.6%)	862 (5.8%)	
<b>Hemoglobin &lt;10 g/dL</b>	368 (2.3%)	261 (0.6%)	0.14	170 (1.1%)	138 (0.9%)	0.02
<b>Alanine aminotransferase</b>			0.15			0.03
<30 U/L	4,287 (26.2%)	8,503 (20.9%)		3,622 (24.2%)	3,457 (23.1%)	
30-60 U/L	7,689 (47.0%)	19,078 (46.9%)		7,159 (47.8%)	7,196 (48.0%)	
>60 U/L	3,734 (22.8%)	10,931 (26.9%)		3,603 (24.0%)	3,728 (24.9%)	
<b>Aspartate aminotransferase</b>			0.11			0.02
<30 U/L	4,785 (29.3%)	10,461 (25.7%)		4,127 (27.5%)	4,013 (26.8%)	
30-60 U/L	8,850 (54.1%)	21,926 (54.0%)		8,265 (55.2%)	8,305 (55.4%)	
>60 U/L	2,055 (12.6%)	5,953 (14.6%)		1,972 (13.2%)	2,049 (13.7%)	
<b>Platelet count</b>			0.04			<0.01
$\geq$ 150,000/ $\mu$ L	14,923 (91.3%)	37,388 (92.0%)		13,757 (91.8%)	13,725 (91.6%)	
<150,000/ $\mu$ L	1,388 (8.5%)	3,104 (7.6%)		1,186 (7.9%)	1,218 (8.1%)	
<b>Total bilirubin</b>			<0.01			<0.01
$\leq$ 2 mg/dL	16,323 (99.8%)	40,563 (99.8%)		14,956 (99.8%)	14,961 (99.8%)	
>2 mg/dL	30 (0.2%)	76 (0.2%)		29 (0.2%)	24 (0.2%)	
<b>International normalized ratio</b>			0.03			<0.01



<1.5	13,132 (80.3%)	32,172 (79.2%)		11,904 (79.4%)	11,869 (79.2%)	
≥1.5	3,221 (19.7%)	8,467 (20.8%)		3,081 (20.6%)	3,116 (20.8%)	
<b>Model of End-Stage Liver Disease (MELD) score<sup>‡</sup></b>			0.39			0.09
<10	10,721 (65.6%)	29,849 (73.4%)		10,500 (70.1%)	10,709 (71.5%)	
10-14	1,199 (7.3%)	2,218 (5.5%)		1,126 (7.5%)	1,060 (7.1%)	
≥15	1,233 (7.5%)	161 (0.4%)		290 (1.9%)	140 (0.9%)	
<b>Estimated glomerular filtration rate &lt;30 mL/min/1.73m<sup>2</sup></b>	1,320 (8.1%)	116 (0.3%)	0.40	298 (2.0%)	115 (0.8%)	0.11
<b>Use of ribavirin as part of DAA regimen</b>	3,075 (18.8%)	3,441 (8.5%)	0.30	2,899 (19.3%)	3,003 (20.0%)	0.02
<b>Year of DAA initiation</b>			0.58			0.38
2014-2015	3,701 (22.6%)	9,167 (22.6%)		3,595 (24.0%)	1,990 (13.3%)	
2016-2017	7,557 (46.2%)	27,579 (67.9%)		6,813 (45.5%)	9,520 (63.5%)	
2018-2019	5,095 (31.2%)	3,893 (9.6%)		4,577 (30.5%)	3,475 (23.2%)	

DAA, direct-acting antiviral; FIB-4, Fibrosis-4 index for liver fibrosis; HCV, hepatitis C virus; IQR, interquartile range; PI,

protease inhibitor; RNA, ribonucleic acid; SDP, standardized difference in proportion

\* Includes glecaprevir/pibrentasvir, elbasvir/grazoprevir, or paritaprevir/ritonavir/ombitasvir with or without dasabuvir

† Includes sofosbuvir/ledipasvir or sofosbuvir/velpatasvir

‡ MELD score calculation:  $3.78 \times \ln[\text{total bilirubin (mg/dL)}] + 11.2 \times \ln[\text{INR}] + 9.57 \times \ln[\text{creatinine (mg/dL)}] + 6.43$

**Table 2. Baseline characteristics of patients with FIB-4 >3.25 prescribed protease inhibitor-based and non-protease inhibitor (PI)-based direct-acting antiviral regimens of interest for chronic hepatitis C virus infection prior to and after propensity score matching.**

Characteristics	Prior to Propensity Score Matching			After Propensity Score Matching		
	PI-Based DAA Regimen* (n=3,816)	Non-PI-Based DAA Regimen† (n=10,583)	SDP	PI-Based DAA Regimen* (n=3,513)	Non-PI-Based DAA Regimen† (n=3,513)	SDP
<b>Age</b>			0.12			0.04
<55 years	80 (2.1%)	324 (3.1%)		75 (2.1%)	62 (1.8%)	
55-59 years	619 (16.2%)	1,971 (18.6%)		585 (16.7%)	610 (17.4%)	
60-64 years	1,465 (38.4%)	4,124 (39.0%)		1,366 (38.9%)	1,347 (38.3%)	
65-69 years	1,400 (36.7%)	3,644 (34.4%)		1,260 (35.9%)	1,276 (36.3%)	
≥70 years	252 (6.6%)	520 (4.9%)		227 (6.5%)	218 (6.2%)	
<b>Male sex</b>	3,738 (98.0%)	10,310 (97.4%)	0.04	3,442 (98.0%)	3,426 (97.5%)	0.03
<b>Race/ethnicity</b>			0.16			0.05
Black	1,771 (46.4%)	4,138 (39.1%)		1,552 (44.2%)	1,491 (42.4%)	
White	1,611 (42.2%)	5,296 (50.0%)		1,546 (44.0%)	1,637 (46.6%)	
Hispanic	248 (6.5%)	641 (6.1%)		237 (6.7%)	220 (6.3%)	
Other/Unknown	186 (4.9%)	508 (4.8%)		178 (5.1%)	165 (4.7%)	
<b>Body mass index</b>			0.11			0.03
Underweight (<18.50 kg/m <sup>2</sup> )	87 (2.3%)	229 (2.2%)		80 (2.3%)	77 (2.2%)	
Normal (18.50-24.99 kg/m <sup>2</sup> )	1,201 (31.5%)	3,112 (29.4%)		1,088 (31.0%)	1,065 (30.3%)	
Overweight (25.00-29.99 kg/m <sup>2</sup> )	1,303 (34.1%)	3,758 (35.5%)		1,196 (34.0%)	1,211 (34.5%)	
Obesity (30.00-34.99 kg/m <sup>2</sup> )	671 (17.6%)	2,025 (19.1%)		624 (17.8%)	611 (17.4%)	
Morbid obesity (≥35.00 kg/m <sup>2</sup> )	321 (8.4%)	1,010 (9.5%)		305 (8.7%)	323 (9.2%)	
Unknown	233 (6.1%)	449 (4.2%)		220 (6.3%)	226 (6.4%)	
<b>Diabetes mellitus</b>	1,230 (32.2%)	3,197 (30.2%)	0.04	1,056 (30.1%)	1,095 (31.2%)	0.02
<b>Alcohol dependence/abuse diagnosis</b>	2,136 (56.0%)	6,161 (58.2%)	0.05	1,997 (56.8%)	2,002 (57.0%)	<0.01
<b>HCV RNA &gt;800,000 IU/mL</b>	2,522 (66.1%)	6,946 (65.6%)	0.04	2,338 (66.6%)	2,306 (65.6%)	0.05
<b>HCV genotype</b>			0.54			0.10
Genotype 1a	1,841 (48.2%)	6,635 (62.7%)		1,715 (48.8%)	1,835 (52.2%)	
Genotype 1b	1,458 (38.2%)	1,704 (16.1%)		1,302 (37.1%)	1,162 (33.1%)	
Genotype 1, subtype unknown	60 (1.6%)	280 (2.6%)		59 (1.7%)	51 (1.5%)	
Genotype 2	74 (1.9%)	495 (4.7%)		71 (2.0%)	55 (1.6%)	
Genotype 3	109 (2.9%)	664 (6.3%)		108 (3.1%)	108 (3.1%)	
Other genotype	206 (5.4%)	575 (5.4%)		192 (5.5%)	225 (6.4%)	
<b>Hemoglobin &lt;10 g/dL</b>	113 (3.0%)	157 (1.5%)	0.10	64 (1.8%)	49 (1.4%)	0.03
<b>Alanine aminotransferase</b>			0.06			<0.01
<30 U/L	326 (8.5%)	748 (7.1%)		237 (6.7%)	235 (6.7%)	
30-60 U/L	1,109 (29.1%)	3,051 (28.8%)		1,007 (28.7%)	1,008 (28.7%)	
>60 U/L	2,145 (56.2%)	6,108 (57.7%)		2,054 (58.5%)	2,061 (58.7%)	
<b>Aspartate aminotransferase</b>			0.10			0.02
<30 U/L	94 (2.5%)	156 (1.5%)		62 (1.8%)	57 (1.6%)	
30-60 U/L	951 (24.9%)	2,365 (22.3%)		822 (23.4%)	847 (24.1%)	
>60 U/L	2,530 (66.3%)	7,325 (69.2%)		2,408 (68.5%)	2,398 (68.3%)	
<b>Platelet count</b>			0.09			0.03
≥150,000/μL	1,182 (31.0%)	2,859 (27.0%)		1,103 (31.4%)	1,055 (30.0%)	
<150,000/μL	2,619 (68.6%)	7,690 (72.7%)		2,397 (68.2%)	2,445 (69.6%)	
<b>Total bilirubin</b>			0.17			<0.01
≤2 mg/dL	3,765 (98.7%)	10,152 (95.9%)		3,462 (98.5%)	3,463 (98.6%)	
>2 mg/dL	51 (1.3%)	431 (4.1%)		51 (1.5%)	50 (1.4%)	

<b>International normalized ratio</b>			0.04			0.02
<1.5	3,310 (86.7%)	9,036 (85.4%)		3,033 (86.3%)	3,010 (85.7%)	
≥1.5	506 (13.3%)	1,547 (14.6%)		480 (13.7%)	503 (14.3%)	
<b>Model of End-Stage Liver Disease (MELD) score<sup>‡</sup></b>			0.32			0.06
<10	2,635 (69.1%)	7,696 (72.7%)		2,569 (73.1%)	2,588 (73.7%)	
10-14	426 (11.2%)	1,310 (12.4%)		411 (11.7%)	395 (11.2%)	
≥15	282 (7.4%)	114 (1.1%)		80 (2.3%)	53 (1.5%)	
<b>Estimated glomerular filtration rate &lt;30 mL/min/1.73m<sup>2</sup></b>	281 (7.4%)	42 (0.4%)	0.37	62 (1.8%)	41 (1.2%)	0.05
<b>Use of ribavirin as part of DAA regimen</b>	1,320 (34.6%)	2,434 (23.0%)	0.26	1,236 (35.2%)	1,344 (38.3%)	0.06
<b>Year of DAA initiation</b>			0.25			0.18
2014-2015	1,337 (35.0%)	3,515 (33.2%)		1,260 (35.9%)	1,001 (28.5%)	
2016-2017	1,722 (45.1%)	5,828 (55.1%)		1,545 (44.0%)	1,844 (52.5%)	
2018-2019	757 (19.8%)	1,240 (11.7%)		708 (20.2%)	668 (19.0%)	

DAA, direct-acting antiviral; FIB-4, Fibrosis-4 index for liver fibrosis; HCV, hepatitis C virus; IQR, interquartile range; PI,

protease inhibitor; RNA, ribonucleic acid; SDP, standardized difference in proportion

\* Includes glecaprevir/pibrentasvir, elbasvir/grazoprevir, or paritaprevir/ritonavir/ombitasvir with or without dasabuvir

† Includes sofosbuvir/ledipasvir or sofosbuvir/velpatasvir

‡ MELD score calculation:  $3.78 \times \ln[\text{total bilirubin (mg/dL)}] + 11.2 \times \ln[\text{INR}] + 9.57 \times \ln[\text{creatinine (mg/dL)}] + 6.43$

**Table 3. Absolute risk and unadjusted incidence rates of specified acute liver injury outcomes among propensity score-matched protease inhibitor (PI) and non-PI initiators, by baseline advanced hepatic fibrosis/cirrhosis status by FIB-4. Incidence rates are reported as events per 1,000 person-years.**

Regimen	No. Exposed	ALT >200 U/L			Severe Hepatic Dysfunction*			Hepatic Decompensation†		
		No. Events	Absolute Risk	Incidence Rates (95% CI)	No. Events	Absolute Risk	Incidence Rates (95% CI)	No. Events	Absolute Risk	Incidence Rates (95% CI)
<b>FIB-4 ≤3.25</b>										
<b>PI-based</b>	14,985	71	0.47%	16.96 (13.44-21.40)	4	0.03%	0.95 (0.36-2.54)	5	0.03%	1.19 (0.50-2.86)
Gle/Pib	4,739	5	0.11%	4.42 (1.84-10.61)	1	0.02%	0.88 (0.12-6.27)	0	0.00%	-
Elb/Gra	5,938	17	0.29%	9.57 (5.95-15.39)	2	0.03%	1.12 (0.28-4.50)	1	0.02%	0.56 (0.08-3.99)
PRO/PROD	4,308	49	1.1%	38.35 (28.98-50.74)	1	0.02%	0.78 (0.11-5.52)	4	0.09%	3.11 (1.17-8.29)
<b>Non-PI-based</b>	14,985	18	0.12%	4.23 (2.66-6.71)	6	0.04%	1.41 (0.63-3.13)	5	0.03%	1.17 (0.49-2.82)
Sof/Led	12,711	14	0.11%	3.91 (2.31-6.60)	4	0.03%	1.12 (0.42-2.97)	5	0.04%	1.39 (0.58-3.35)
Sof/Vel	2,274	4	0.18%	5.92 (2.22-15.77)	2	0.09%	2.96 (0.74-11.82)	0	0.00%	-
<b>FIB-4 &gt;3.25</b>										
<b>PI-based</b>	3,513	25	0.71%	24.35 (16.45-36.04)	19	0.54%	18.48 (11.79-28.97)	13	0.37%	12.63 (7.33-21.75)
Gle/Pib	691	2	0.29%	10.92 (2.73-43.65)	1	0.14%	5.45 (0.77-38.70)	0	0.00%	-
Elb/Gra	1,356	5	0.37%	12.40 (5.16-29.79)	3	0.22%	7.43 (2.40-23.04)	3	0.22%	7.43 (2.40-23.03)
PRO/PROD	1,466	18	1.2%	40.88 (25.76-64.89)	15	1.0%	34.03 (20.51-56.44)	10	0.68%	22.63 (12.17-42.05)
<b>Non-PI-based</b>	3,513	12	0.34%	11.17 (6.35-19.67)	16	0.46%	14.92 (9.14-24.35)	16	0.46%	14.91 (9.13-24.34)
Sof/Led	2,943	10	0.34%	11.13 (5.99-20.69)	13	0.44%	14.49 (8.41-24.95)	15	0.51%	16.71 (10.07-27.72)
Sof/Vel	570	2	0.35%	11.40 (2.85-45.57)	3	0.53%	17.12 (5.52-53.09)	1	0.18%	5.70 (0.80-40.48)

ALT, alanine aminotransferase; CI, confidence interval; Elb, elbasvir; FIB-4, Fibrosis-4 Index for Hepatic Fibrosis; Gle, glecaprevir; Gra, grazoprevir; Led, ledipasvir; PI, protease inhibitor; Pib, pibrentasvir; PRO, paritaprevir/ritonavir/ombitasvir; PROD, paritaprevir/ritonavir/ombitasvir with dasabuvir; Sof, sofosbuvir; Vel, velpatasvir

\* Severe hepatic dysfunction defined by 1 inpatient or outpatient international normalized ratio ≥1.5 and total bilirubin >2 times the upper limit of normal within 30 days of each other.

† Hepatic decompensation defined by 1 hospital discharge diagnosis or 2 or more outpatient diagnoses of ascites, spontaneous bacterial peritonitis, esophageal variceal hemorrhage, or hepatic encephalopathy. The decompensation date was defined as the hospital discharge date (if event was identified by hospital diagnosis) or initial outpatient diagnosis date (if identified by outpatient diagnoses).

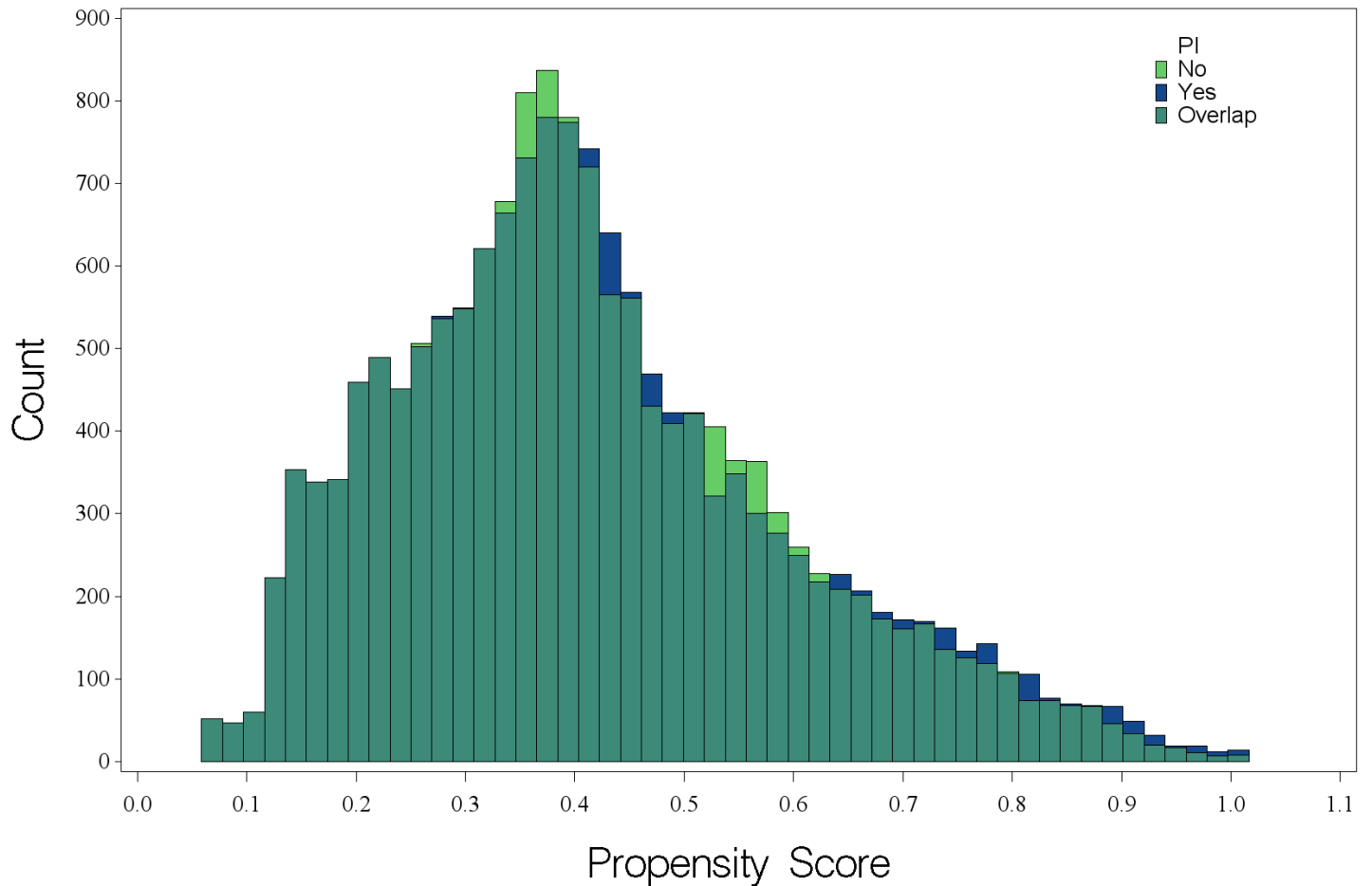
# **PI-Based HCV DAAs Are Associated with Increased Risk of Aminotransferase Elevations but Not Hepatic Dysfunction or Decompensation**

Jessie Torgersen, Craig W. Newcomb, Dena M. Carbonari, Christopher T. Rentsch, Lesley S. Park, Alyssa Mezochow, Rajni L. Mehta, Lynn Buchwalder, Janet P. Tate, Norbert Bräu, Debika Bhattacharya, Joseph K. Lim, Tamar H. Taddei, Amy C. Justice, Vincent Lo Re III

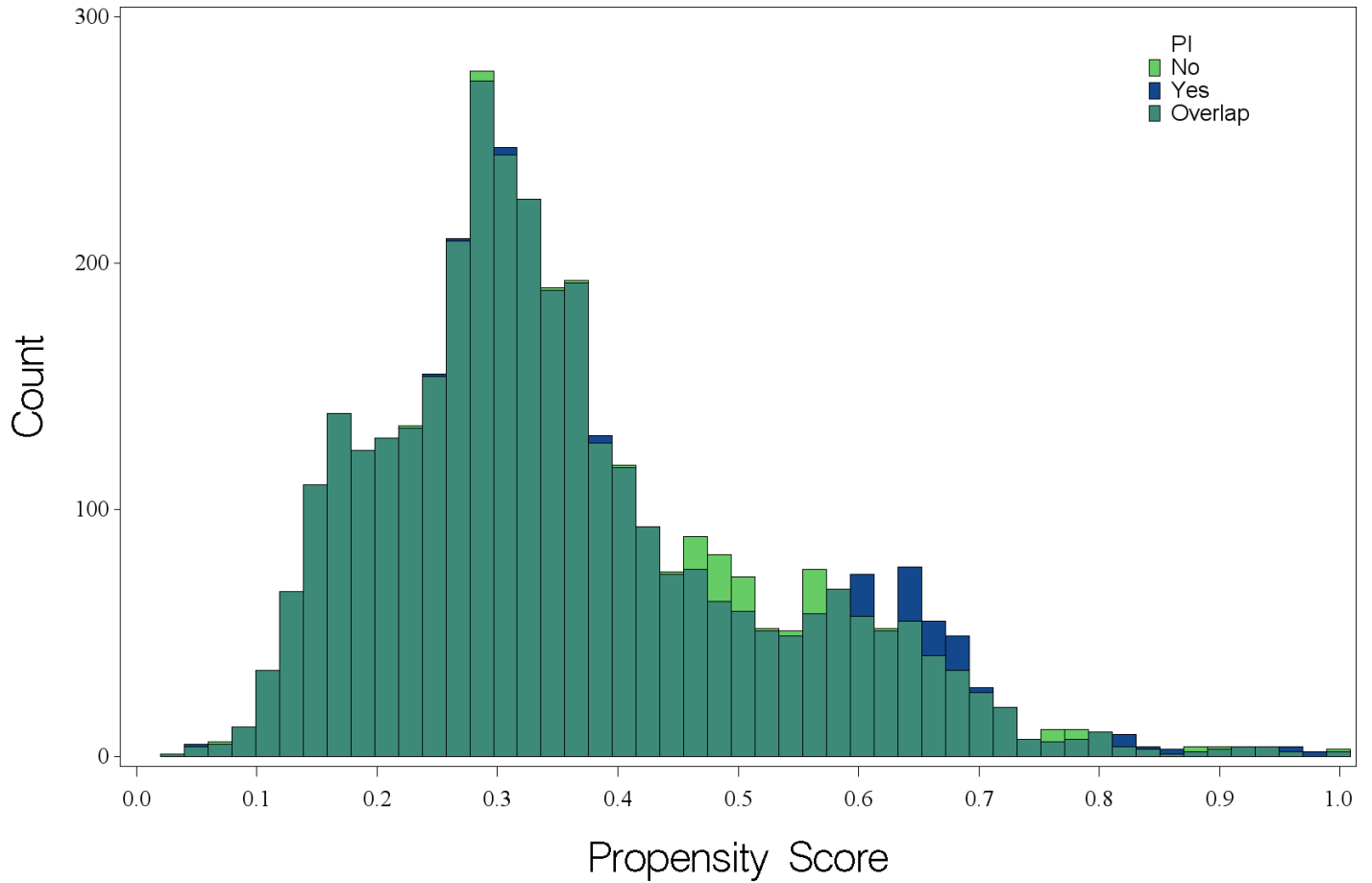
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**Supplementary Figure 1. Propensity score histogram by protease inhibitor (PI)-based and non-PI-based direct-acting antiviral (DAA) therapy for patients with FIB-4  $\leq 3.25$ .** Distribution illustrated as percent of patients receiving PI-based (light green) and non-PI-based (blue) DAA therapy by propensity score and overlap (dark green) of these two groups.



**Supplementary Figure 2. Propensity score histogram by protease inhibitor (PI)-based and non-PI-based direct-acting antiviral (DAA) therapy for patients with FIB-4 >3.25.** Distribution illustrated as percent of patients receiving PI-based (light green) and non-PI-based (blue) DAA therapy by propensity score and overlap (dark green) of these two groups.



**Supplementary Table 1. International Classification of Diseases, Ninth Revision (ICD-9) and International Classification of Diseases, Tenth Revision (ICD-10) diagnoses used to identify hepatic decompensation events among protease inhibitor and non-protease inhibitor-based direct-acting antiviral initiator cohorts.**

ICD-9 Code(s)	ICD-10 Code(s)	Description
456.0; 456.2	I85.01; I85.11	Esophageal Varices With Bleeding
567.23	K65.2	Spontaneous Bacterial Peritonitis
572.2	K70.41; K72.11; K72.91	Hepatic Coma; Hepatic Failure With Coma
789.5; 789.59	K70.11; K70.31; K71.51; R18.8	Ascites
572.4	K76.7	Hepatorenal Syndrome
--	K76.81	Hepatopulmonary Syndrome



**Supplementary Table 2. Frequencies of specific hepatic decompensation diagnoses among propensity score-matched protease inhibitor (PI) and non-PI initiators, by baseline advanced hepatic fibrosis/cirrhosis status by FIB-4.**

<b>Regimen</b>	<b>Bleeding Esophageal Varices</b>	<b>Spontaneous Bacterial Peritonitis</b>	<b>Hepatic Coma</b>	<b>Ascites</b>	<b>Hepatorenal Syndrome</b>	<b>Hepatopulmonary Syndrome</b>
<b>FIB-4 ≤3.25</b>						
PI-based	0	1	0	3	1	0
Non-PI-based	1	0	0	4	0	0
<b>FIB-4 &gt;3.25</b>						
PI-based	3	0	3	7	0	0
Non-PI-based	4	1	1	9	0	1

FIB-4, Fibrosis-4 index for liver fibrosis; PI, protease inhibitor