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1 **Title:** Association of Serum Autotaxin Levels with Liver Fibrosis in Patients Pre- and
2 Post-Treatment with Chronic Hepatitis C

3

4 **Abstract**

5 **Background & Aim:** The evaluation of liver fibrosis in patients with chronic hepatitis
6 C virus (HCV) infection is important as it is a risk factor for hepatocellular carcinoma.
7 In the recent years, autotaxin (ATX) has been established as a new non-invasive
8 biomarker to predict liver fibrosis. However, antiviral treatment has been reported to
9 decrease serum ATX, but it is unclear whether post-treatment ATX levels reflect liver
10 fibrosis. In the present study, we analyzed the correlation between ATX and liver
11 fibrosis in pre- and post-treatment patients with HCV infection.

12 **Methods:** We used 199 samples from 136 patients with HCV infection who had
13 undergone hepatic biopsy before and/or after antiviral treatment at Osaka City
14 University Hospital. Post-treatment patients included 38 interferon (IFN)-treated
15 patients and 80 IFN-free direct-acting antiviral-treated patients; all patients achieved
16 a sustained virological response (SVR). Serum ATX levels were determined by
17 enzyme immunoassay with an AIA-2000 analyzer.

18 **Results:** Serum ATX levels were largely correlated with liver fibrosis stage in
19 patients with HCV infection before and after antiviral treatment. The measured
20 values decreased even in similar liver fibrosis stages after treatment. The area under
21 the receiver operating characteristic curve for the ability of ATX to diagnose above
22 F2 stage before treatment was 0.81 (both male and female) and that after achieving
23 SVR was 0.71 (male) and 0.72 (female).

24 **Conclusions:** Serum ATX levels were correlated with histological liver fibrosis stage
25 after achieving SVR. However, we should establish separate cutoff values before
26 and after antiviral therapy.

1

2 **Keywords:** Autotaxin, Chronic hepatitis C virus, Liver fibrosis stage, Serum liver

3 fibrosis marker, Sustained virological response

4

1 Introduction

2 Patients infected with hepatitis C virus (HCV) develop chronic hepatitis and 20-30%
3 of them progress to liver cirrhosis and/or hepatocellular carcinoma (HCC) ¹. In the
4 recent years, most patients could achieve sustained virological response (SVR) by
5 interferon (IFN)-free direct-acting antiviral agent (DAA) therapy ²⁻⁴. However, HCC
6 develops in patients with liver fibrosis regardless of achieving SVR. Therefore, the
7 evaluation of liver fibrosis stage is needed even after antiviral treatment. Hepatic
8 biopsy is the gold standard by which to evaluate liver fibrosis, but it is invasive ⁵. In
9 addition, it has problems of sampling error and inter-observer disparity. Therefore,
10 serum liver fibrosis markers are being paid more attention as non-invasive
11 biomarkers. Platelet counts are known to associate with liver fibrosis stage. The FIB-
12 4 index and aspartate aminotransferase (AST)-to-platelet ratio (APRI) are markers
13 that can be calculated by basic laboratory data ^{6,7}. In contrast, these fibrosis markers
14 are influenced by various factors, as these are calculated by age, platelet counts,
15 AST, and alanine aminotransferase (ALT).

16 Autotaxin (ATX) was discovered as an autocrine motility-stimulating protein
17 in the conditioned medium from A2058 human melanoma cell cultures ⁸. ATX has
18 lysophospholipase D activity to generate lysophosphatidic acid (LPA) from
19 lysophospholipids in the blood ⁹. LPA is involved in such physiological roles as cell
20 migration, cell proliferation, neurogenesis, and angiogenesis ¹⁰. LPA also stimulates
21 the proliferation and contractility of hepatic stellate cells. Recent studies have
22 revealed that serum ATX levels correlate with liver fibrosis stage in patients with
23 HCV ^{11,12}, chronic hepatitis B virus (HBV) ¹³, non-alcoholic fatty liver disease
24 (NAFLD) ¹⁴, and primary biliary cholangitis (PBC) ^{15,16}. Because ATX is metabolized
25 by liver sinusoidal endothelial cells, reduced metabolism by liver fibrosis and/or
26 damage leads to an increase in serum ATX ¹⁷. A sex difference in ATX has been

1 reported, where the reference value of females is higher than that of males¹⁸. In
2 Japan, ATX has been used in clinical practice since 2018, and the cutoff values for
3 predicting above the F2 stage are set at 0.91 mg/L (male) and 1.27 mg/L (female),
4 and in predicting liver cirrhosis at 1.69 mg/L (male) and 2.12 mg/L (female). On the
5 other hand, IFN-free DAA therapy has been reported to decrease serum ATX levels
6¹⁹⁻²¹. There is no data on the relationship between post-treatment histological liver
7 fibrosis stage and ATX, and it is unclear whether serum ATX levels are useful in
8 assessing liver fibrosis after achieving SVR.

9 In the present study, we revealed an association of serum ATX levels with
10 liver fibrosis stage in patients with HCV before and after antiviral treatment.

11

12 **Methods**

13 *Subjects*

14 The classification of patients in this study is shown in Figure 1. We used 199
15 samples from 136 patients with HCV infection who had undergone hepatic biopsy
16 before and/or after antiviral therapy at Osaka City University Hospital between 1999
17 and 2019. We diagnosed chronic hepatitis C based on the presence of serum HCV
18 antibody and detectable HCV RNA by real-time PCR. There were 81 samples
19 collected before antiviral treatment and 118 samples collected after achieving SVR
20 as a result of antiviral therapy. Paired biopsies before and after antiviral therapy were
21 performed in 63 patients. Serum samples were collected from these 63 patients both
22 before and after treatment. The therapy regimens of patients collected after
23 treatment were as follows: IFN-based therapy (n = 38), sofosbuvir + ledipasvir (n =
24 33), sofosbuvir + ribavirin (n = 21), asunaprevir + daclatasvir (n = 19), elbasvir +
25 grazoprevir (n = 5), and ombitasvir + paritaprevir + ritonavir (n = 2). The median
26 period from the end of treatment to sera collection was 15 months. Blood samples

1 were obtained within six months before or after hepatic biopsy. This study was
2 conducted according to the principals of the Declaration of Helsinki and was
3 approved by the Ethics Committee of the Osaka City University Graduate School of
4 Medicine (approval number: 4097).

5

6 *Measurement of serum ATX levels*

7 Serum samples were stored at -80°C until testing. Before measuring ATX, serum
8 samples were thawed at room temperature and centrifuged at 3,500 g for 5 min.

9 Serum ATX levels were measured by a two-site enzyme immunoassay with an AIA-
10 2000 analyzer (Tosoh Co.; Tokyo, Japan).

11

12 *Laboratory data*

13 Platelet counts and routine biochemical tests such as AST and ALT were analyzed
14 by standard procedures. Laboratory data were obtained by medical records. We
15 used laboratory data within seven days before and after taking the blood. The FIB-4
16 index and APRI were calculated according to the published formulae (FIB-4 index,
17 $(\text{age [years]} \times \text{AST [U/L]}) / (\text{platelet count [}10^9\text{/L]} \times \text{ALT [U/L]}^{1/2})$; APRI, $(\text{AST [U/L]} /$
18 $40 \text{ [U/L]}) \times (100 / \text{platelet count [}10^9\text{/L]})$)^{6,7}.

19

20 *Hepatic biopsy evaluation*

21 Hepatic biopsy was performed with a 16- or 18-gauge Tru-Cut needle (Merit Medical;
22 South Jordan, UT) under ultrasound guidance. We evaluated hepatic fibrosis stage
23 according to the New Inuyama Classification²². The stage of fibrosis was classified
24 from F0 to F4 as follows: F0 = no fibrosis, F1 = portal expansion of fibrosis, F2 =
25 bridging fibrosis (portal-portal or portal-central linkage), F3 = bridging fibrosis with
26 lobular distortion (disorganization), and F4 = cirrhosis.

1

2 *Statistical analysis*

3 All statistical analyses and data visualizations were performed by R software

4 (version 3.5.3; R Foundation for Statistical Computing, Vienna, Austria).

5 Comparisons of serum ATX levels for each fibrosis stage were performed by the

6 Steel–Dwass test. Comparisons of serum ATX levels before and after antiviral

7 therapy were analyzed by the Wilcoxon signed-rank test. The predictive capabilities

8 of liver fibrosis stage were analyzed by the area under the receiver operating

9 characteristic (ROC) curve (AUC). Cutoff values were identified by the Youden

10 index. A *p*-value of less than 0.05 was considered statistically significant.

11

12 **Results**

13 *Serum ATX levels and other fibrosis markers pre- and post-treatment*

14 The clinical characteristics of patients with HCV before and after antiviral therapy are

15 summarized in Table 1. First, we revealed that serum ATX levels correlated with liver

16 fibrosis stage in patients with HCV before treatment (Figure 2A). Median values of

17 ATX in patients with F1, F2, F3, and F4 stage fibrosis were 1.05, 1.23, 1.71, and

18 2.74 mg/L (male) and 1.11, 1.86, 1.44, and 2.41 mg/L (female), respectively (Figures

19 2B, C). Platelet counts, the FIB-4 index, and APRI were also correlated with liver

20 fibrosis (Supplemental Figure 1). Second, we analyzed post-treatment HCV patients.

21 We also revealed that serum ATX levels reflected liver fibrosis stage even after

22 antiviral therapy (Figure 3A). There was no significant difference between male

23 patients of each fibrosis stage, but there was a correlation (Figure 3B). Median ATX

24 values in patients with F1, F2, F3, and F4 stage fibrosis were 0.80, 0.95, 1.02, and

25 1.22 mg/L (male), and 1.21, 1.56, 1.39, and 1.83 mg/L (female), respectively

26 (Figures 3B, C). Other liver fibrosis markers also similarly correlated with liver

1 fibrosis stage (Supplemental Figure 2).

2

3 *Change in the serum ATX levels between pre- and post-treatment*

4 We selected the 63 patients who had undergone hepatic biopsy both before and
5 after antiviral treatment and analyzed the change in serum ATX levels among
6 therapy. Serum ATX levels after antiviral treatment decreased than that before
7 treatment (Figure 4A). Next, we divided patients into 3 groups according to the
8 alteration of liver fibrosis stage as follows: improved (n = 17), sustained (n = 31), and
9 exacerbated groups (n = 15). After achieving SVR, serum ATX levels decreased in
10 patients whose liver fibrosis stage improved (median value; 1.39 mg/L vs. 1.09 mg/L,
11 Figure 4B) and was sustained (1.24 mg/L vs. 1.16 mg/L, Figure 4C). Despite
12 worsening liver fibrosis, serum ATX levels tended to decrease, but without statistical
13 significance (1.37 mg/L vs. 1.31 mg/L, Figure 4D). Also, we analyzed whether IFN-
14 based therapy and IFN-free DAA therapy differed in the changes in ATX. Of the 63
15 patients who underwent paired biopsy, 36 patients received IFN-based therapy
16 (improved, 10; sustained, 16; exacerbated, 10), and 27 patients received DAA
17 therapy (improved, 7; sustained, 15; exacerbated, 5). There was no difference in the
18 rate of ATX change before and after antiviral treatment between IFN-based therapy
19 and DAA therapy (-11.99% vs. -10.42%, Supplemental Figure 3).

20

21 *Predictive capabilities of liver fibrosis markers in pre- and post-treatment*

22 We analyzed the predictive capabilities for fibrosis above F2 stage of serum ATX
23 levels, platelet counts, the FIB-4 index, and APRI. ATX had the highest AUC among
24 the other fibrosis markers in male patients before treatment (0.81, Figure 5A). The
25 AUC of female patients was equal to that of male patients (0.81, Figure 5B). In
26 contrast, the diagnostic performance of each fibrosis marker decreased after

1 achieving SVR. However, ATX had a higher AUC than platelet counts, and it was
2 equal to APRI (male, 0.71; female, 0.72; Figures 5C, D). In pre-treatment, cutoff
3 values for predicting fibrosis above F2 stage in male and female patients were 1.21
4 mg/L (sensitivity: 0.75, specificity: 0.9) and 1.11 mg/L (sensitivity: 1.0, specificity:
5 0.53), respectively. On the other hand, after achieving SVR, cutoff values for male
6 patients was 0.76 mg/L (sensitivity: 0.90, specificity: 0.47) and 1.32 mg/L (sensitivity:
7 0.76, specificity: 0.73) in female patients, respectively. We also compared AUCs
8 between serum ATX levels and liver fibrosis markers in predicting each liver fibrosis
9 stage before and after antiviral treatment (Table 2). Before treatment, ATX had the
10 highest AUC compared to platelet counts, the FIB-4 index, and APRI (\geq F2; 0.81, \geq
11 F3; 0.89, = F4; 0.89) in male patients. ATX in female patients also had a high
12 diagnostic performance (\geq F2; 0.81, \geq F3; 0.77, = F4; 0.80). However, the FIB-4
13 index and APRI had higher AUCs than ATX after achieving SVR. The diagnostic
14 performance for predicting above F3 stage (FIB-4, 0.76; APRI, 0.73) and F4 stage
15 (FIB-4, 0.76; APRI, 0.76) was superior, especially in male patients. On the other
16 hand, we also found that ATX in female patients had the highest AUC compared to
17 other fibrosis markers only for predicting F4 stage after treatment (AUC = 0.86).

18

19 Discussion

20 Previous studies revealed that ATX is a useful biomarker in predicting liver fibrosis
21 stage¹²⁻¹⁵. In the present study, we analyzed patients before treatment and after
22 achieving SVR. First, the present study showed that serum ATX levels correlated
23 with liver fibrosis stage in patients with HCV infection before treatment. Moreover,
24 the predictive potential of ATX for liver fibrosis stage was higher compared to platelet
25 counts, the FIB-4 index, and APRI in male patients. ATX in female patients also had
26 a high diagnostic capacity in predicting hepatic fibrosis. In particular, the diagnostic

1 ability of liver cirrhosis was superior and may be useful to identify high-risk patients
2 for HCC before antiviral therapy.

3 Second, we also found that serum ATX levels largely correlated with liver
4 fibrosis stage after achieving SVR. On the other hand, the measured values after
5 antiviral treatment decreased compared to before treatment. We revealed that the
6 rate of ATX reduction before and after antiviral treatment in patients whose liver
7 fibrosis stage improved tended to be higher than that in patients whose fibrosis
8 became exacerbated, but without statistical significance (median values; -26.97%
9 vs. -7.47%, $p = 0.08$, Supplemental Figure 4). We suggested that the rate of ATX
10 change predicts the improvement of the fibrosis stage, but it is necessary to evaluate
11 a larger sample size. According to a previous study, IFN-free DAA therapy has been
12 reported to decrease serum ATX levels four weeks after the start of treatment ¹⁹.
13 Because liver fibrosis does not improve in such a short period ²³, serum ATX levels
14 are expected to reflect not only liver fibrosis but also hepatitis activity and/or the
15 presence of HCV. Yamazaki et al. ¹² reported that serum ATX levels were weakly
16 correlated with serum ALT levels ($r = 0.329$). We also demonstrated a weak
17 correlation between ATX and ALT ($r = 0.231$, $p = 0.001$). In addition, we have
18 illustrated the clinical characteristics of patients whose liver fibrosis were improved or
19 sustained but serum ATX levels were not decreased (group A, B) as well as those of
20 patients whose liver fibrosis was exacerbated but serum ATX levels were decreased
21 (group C) in Supplemental Table 1. Of the 63 patients in whom we performed paired
22 biopsy, only 4 patients did not have decreased ALT levels following treatment, but 3
23 of those patients were included in group B (No. 9, 10, 13, Supplemental Table 1).
24 Also, in patients with improved or sustained fibrosis, 5 patients (of 13 patients) in the
25 ATX-elevated group and only 2 patients (of 35 patients) in the ATX-decreased group
26 had post-treatment ALT levels of > 30 U/L. We suggested that ATX reduction may

1 be inhibited in patients with residual liver inflammation. On the other hand, the
2 patients in whom liver fibrosis was exacerbated, pretreatment ALT levels in the ATX-
3 decreased group were higher than those in the ATX-elevated group (median value;
4 117 U/L vs. 23 U/L, $p = 0.03$). Patients in the ATX-decreased group may have strong
5 hepatic inflammation, increasing pretreatment ATX levels. Therefore, serum ATX
6 levels were suggested to reflect not only liver fibrosis but also liver inflammation. In
7 fact, we saw no change in serum ATX levels in patients with treatment failure (data
8 not shown). This is similar to *Wisteria floribunda agglutinin*-positive (WFA⁺) Mac-2-
9 binding protein (M2BPGi), which is a glycomarker for predicting liver fibrosis ²⁴.

10 Moreover, the present study showed that the diagnostic ability of ATX in
11 predicting liver fibrosis stage decreased after achieving SVR. From the results in
12 Table 2, ATX was expected to be superior as a liver fibrosis marker before
13 treatment. However, the FIB-4 index and APRI had better predictive performance
14 after antiviral therapy. We revealed that serum ATX levels varied even in patients
15 with similar liver fibrosis stage before and after antiviral therapy. We suspect that
16 serum ATX levels are influenced by factors other than liver fibrosis. The reference
17 values of serum ATX in females are significantly higher than in males ¹⁸. In addition,
18 ATX has been reported to be elevated in pregnant women ²⁵, and may be affected
19 by sexual hormones and female reproductive organs. A previous study has also
20 shown that serum ATX levels increased in patients with follicular lymphoma ²⁶. On
21 the other hand, it has been reported that there are few changes in ATX in kidney
22 disease, heart disease, and diabetes ²⁷. It is also unaffected by meals ²⁷. It is
23 important that the factors affecting serum ATX levels are identified to make accurate
24 clinical decisions.

25 In the recent years, IFN-free DAA therapy could achieve SVR in most
26 patients with HCV infection. However, HCC develops in patients with SVR and

1 advanced hepatic fibrosis. It is important to monitor HCC after SVR to establish
2 predictive markers for hepatic fibrosis. Our data showed that ATX is associated with
3 histological hepatic fibrosis both before and after antiviral therapy. We also showed
4 that cutoff values for predicting above the F2 stage differed before and after antiviral
5 therapy. A similar liver fibrosis marker, M2BPGi, has been suggested as a reliable
6 serum marker for liver carcinogenesis^{24,28}. ATX plays a role in converting
7 lysophosphatidylcholine to LPA, which is involved in physiological roles.
8 Interestingly, the ATX-LPA pathway has been reported to associate with the
9 development of HCC^{29,30}. ATX might be a more reliable marker for the development
10 of HCC. Further research is needed on the relationship between ATX and SVR-
11 subsequent carcinogenesis.

12 A limitation of this study is the small number of patients, especially those with
13 advanced liver fibrosis. There were only 3 male patients with F3 fibrosis after
14 treatment and 5 with F4 fibrosis. The lower ATX of F3 stage than that of F2 stage in
15 female may be due to that. The need to analyze by sex makes it more difficult to
16 collect a proper number of samples. We also could not analyze M2BPGi, type IV
17 collagen 7S, or hyaluronic acid, which are established liver fibrosis markers.
18 However, this is the first report to reveal the association of ATX with histological liver
19 fibrosis stage after achieving SVR.

20 In conclusion, we revealed that serum ATX levels were correlated with
21 histological liver fibrosis stage before and after antiviral therapy. Also, our findings
22 indicate that the diagnostic capability of ATX for predicting liver fibrosis differs before
23 and after antiviral therapy. We should establish separate cutoff values before and
24 after treatment. Although several non-invasive liver fibrosis markers have been
25 reported, we need to consider the characteristics of each marker.

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1 **Table 1. Clinical characteristics of patients with HCV infection pre- and post-**
 2 **treatment**

	Pre	Post
Male/Female	38/43	55/63
Age	61 (52-66)	68 (59-73)
Therapy regimen (A/B/C/D/E/F) [†]	—	38/33/21/19/5/2
Post-treatment period (month)	—	15 (6-37)
F0/F1/F2/F3/F4	1/26/24/11/19	4/39/47/14/14
ATX (mg/L)	1.44 (1.11-2.17)	1.17 (0.90-1.44)
PLT ($\times 10^4/\mu\text{L}$)	13.8 (10.7-17.5)	17.5 (13.2-21.3)
AST (U/L)	49 (34-82)	23 (19-32)
ALT (U/L)	49 (29-94)	19 (13-28)
FIB-4 index	3.12 (1.83-4.66)	1.97 (1.59-3.07)
APRI	0.91 (0.50-1.62)	0.36 (0.25-0.53)

Data are presented as median and interquartile range.

† A: interferon-based therapy, B: sofosbuvir + ledipasvir, C: sofosbuvir + ribavirin,
 D: asunaprevir + daclatasvir, E: elbasvir + grazoprevir, F: ombitasvir + paritaprevir + ritonavir
 HCV; hepatitis C virus, ATX; autotaxin, PLT; platelet counts,
 AST; aspartate aminotransferase, ALT; alanine aminotransferase, APRI; AST to platelet ratio

3

1 **Table 2. Comparisons of AUCs between serum ATX levels and other fibrosis**
 2 **markers in predicting fibrosis stage before and after antiviral therapy**

			≥F2	≥F3	F4
Before treatment	Male	ATX	0.81	0.89	0.89
		PLT	0.7	0.88	0.86
		FIB-4	0.72	0.84	0.84
		APRI	0.63	0.73	0.76
	Female	ATX	0.81	0.77	0.8
		PLT	0.73	0.68	0.6
		FIB-4	0.86	0.73	0.67
		APRI	0.8	0.73	0.7
After treatment	Male	ATX	0.71	0.68	0.67
		PLT	0.64	0.64	0.63
		FIB-4	0.68	0.76	0.76
		APRI	0.72	0.73	0.76
	Female	ATX	0.72	0.65	0.86
		PLT	0.6	0.63	0.67
		FIB-4	0.67	0.67	0.72
		APRI	0.73	0.76	0.78

AUC; area under the curve, ATX; autotaxin, PLT; platelet counts,
 APRI; aspartate aminotransferase to platelet ratio

3

1 Figure 1. Classification of patients

1 **Figure 2. Correlation between serum ATX levels and liver fibrosis stage before**
2 **antiviral treatment**

3 Data from all patients (A), male patients (B), and female patients (C) are shown. The
4 number of samples for each fibrosis stage is as follows: F0/F1/F2/F3/F4 =
5 1/26/24/11/19 (all), 1/9/12/6/10 (male), 0/17/12/5/9 (female). Boxes represent the
6 interquartile range of the data. The horizontal lines in the boxes indicate the median
7 values. The vertical lines connect the nearest values of 1.5 times the interquartile
8 range from the quartile point. The dots indicate outliers. ***: $p < 0.001$, **: $p < 0.01$.

1 **Figure 3. Correlation between serum ATX levels and liver fibrosis stage after**
2 **achieving SVR**

3 Data from all patients (A), male patients (B), and female patients (C) are shown. The
4 number of samples for each fibrosis stage is as follows: F0/F1/F2/F3/F4 =
5 4/39/47/14/14 (all), 1/16/30/3/5 (male), 3/23/17/11/9 (female). Boxes represent the
6 interquartile range of the data. The horizontal lines in the boxes indicate the median
7 values. The vertical lines connect the nearest values of 1.5 times the interquartile
8 range from the quartile point. The dots indicate outliers. *: $p < 0.05$.

1 **Figure 4. Change in serum ATX levels after antiviral treatment for each**
2 **alteration of liver fibrosis**

3 The figures show the change in serum ATX levels in all patients (A), the patients
4 whose liver fibrosis stage improved (n = 17) (B), sustained (n = 31) (C), and became
5 exacerbated (n = 15) (D). Boxes represent the interquartile range of the data. The
6 horizontal lines in the boxes indicate the median values. The vertical lines connect
7 the nearest values of 1.5 times the interquartile range from the quartile point. pre:
8 pre-treatment, post: post-treatment. ***: $p < 0.001$, *: $p < 0.05$, NS: not significant.

1 **Figure 5. Receiver operating characteristic curves for predicting above F2**

2 **stage**

3 The data in male patients before treatment (A), female patients before treatment (B),
4 male patients after treatment (C), and female patients after treatment (D) are shown.

5 The numbers at the bottom right are the area under the curve of each liver fibrosis
6 marker.

7

1 **Supplemental Table 1. Clinical characteristics in patients whose fibrosis and**
 2 **ATX changed opposite direction.**

	No.	Age	Sex	Therapy	pre Fibrosis	post Fibrosis	pre ATX (mg/L)	post ATX (mg/L)	pre AST (U/L)	post AST (U/L)	pre ALT (U/L)	post ALT (U/L)
improved (group A)	1	68	F	IFN	F3	F2	1.44	1.56	50	43	69	31
	2	42	F	IFN	F1	F0	0.79	0.87	15	13	17	9
sustained (group B)	3	59	F	DAA	F1	F1	0.94	0.94	34	20	31	11
	4	66	M	DAA	F2	F2	1.02	1.03	28	17	18	12
	5	46	M	DAA	F2	F2	1.23	1.28	56	47	42	40
	6	72	F	DAA	F1	F1	2.44	2.58	166	25	201	18
	7	53	F	IFN	F1	F1	1.11	1.19	16	13	12	7
	8	61	M	IFN	F1	F1	0.59	0.63	25	21	33	19
	9	41	F	IFN	F1	F1	0.72	0.87	34	96	53	125
	10	59	F	IFN	F2	F2	1.29	1.71	24	36	15	35
	11	47	F	IFN	F1	F1	0.97	1.31	19	13	18	8
	12	52	F	DAA	F1	F1	0.82	1.14	87	16	197	14
	13	38	M	IFN	F2	F2	0.53	0.76	39	53	50	50
exacerbated (group C)	14	63	M	IFN	F2	F3	1.45	1.02	103	19	151	15
	15	52	F	IFN	F2	F3	1.85	1.39	111	44	195	71
	16	68	F	IFN	F2	F4	2.35	1.84	75	35	73	30
	17	72	M	DAA	F1	F2	2.07	1.64	101	25	113	15
	18	75	F	DAA	F2	F3	1.71	1.40	30	19	27	16
	19	54	M	IFN	F1	F2	1.21	1.00	341	38	122	61
	20	53	F	IFN	F2	F3	1.13	1.00	37	26	25	15
	21	61	F	IFN	F2	F3	1.37	1.26	39	29	45	25
	22	54	M	IFN	F1	F2	1.21	1.13	341	35	122	21
	23	52	F	IFN	F2	F4	1.85	1.79	111	27	195	39

3 ATX; autotaxin, AST; aspartate aminotransferase, ALT; alanine aminotransferase, IFN; interferon, DAA; direct-acting antiviral agent

1 **Supplemental Figure 1. Correlation between serum liver fibrosis markers and**
2 **liver fibrosis stage before antiviral treatment**

3 The number of samples for each fibrosis stage is as follows: F0/F1/F2/F3/F4 =
4 1/26/24/11/19. Boxes represent the interquartile range of the data. The horizontal
5 lines in the boxes indicate the median values. The vertical lines connect the nearest
6 values of 1.5 times the interquartile range from the quartile point. The dots indicate
7 outliers. ***: $p < 0.001$, **: $p < 0.01$, *: $p < 0.05$.

1 **Supplemental Figure 2. Correlation between serum liver fibrosis markers and**
2 **liver fibrosis stage after achieving SVR.**

3 The number of samples for each fibrosis stage is as follows: F0/F1/F2/F3/F4 =
4 4/39/47/14/14. Boxes represent the interquartile range of the data. The horizontal
5 lines in the boxes indicate the median values. The vertical lines connect the nearest
6 values of 1.5 times the interquartile range from the quartile point. The dots indicate
7 outliers. **: $p < 0.01$, *: $p < 0.05$.

8

1 **Supplemental Figure 3. Comparisons of rate of ATX change between IFN-**
2 **based and DAA therapy.**

3 Boxes represent the interquartile range of the data. The horizontal lines in the boxes
4 indicate the median values. The vertical lines connect the nearest values of 1.5 times
5 the interquartile range from the quartile point. The dots indicate outliers. NS: not
6 significant.

7

1 **Supplemental Figure 4. Association between changes in liver fibrosis stage**
2 **and rate of ATX change.**

3 Boxes represent the interquartile range of the data. The horizontal lines in the boxes
4 indicate the median values. The vertical lines connect the nearest values of 1.5 times
5 the interquartile range from the quartile point. The dots indicate outliers.

6

7

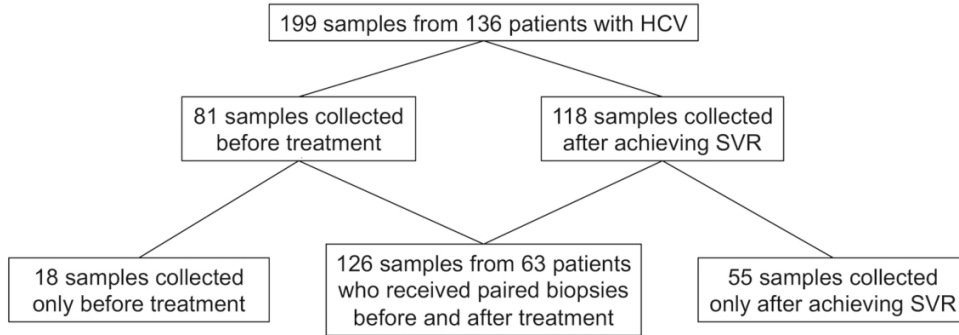


Figure 1

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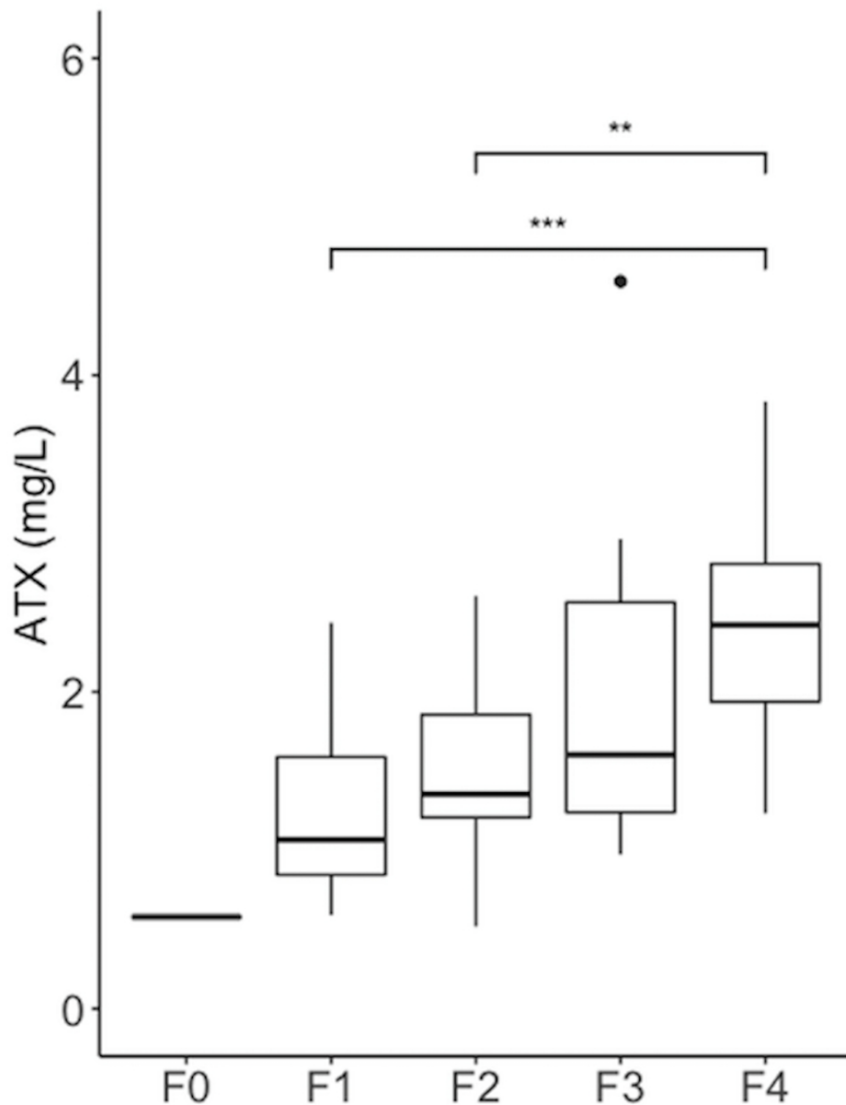


Figure 2A

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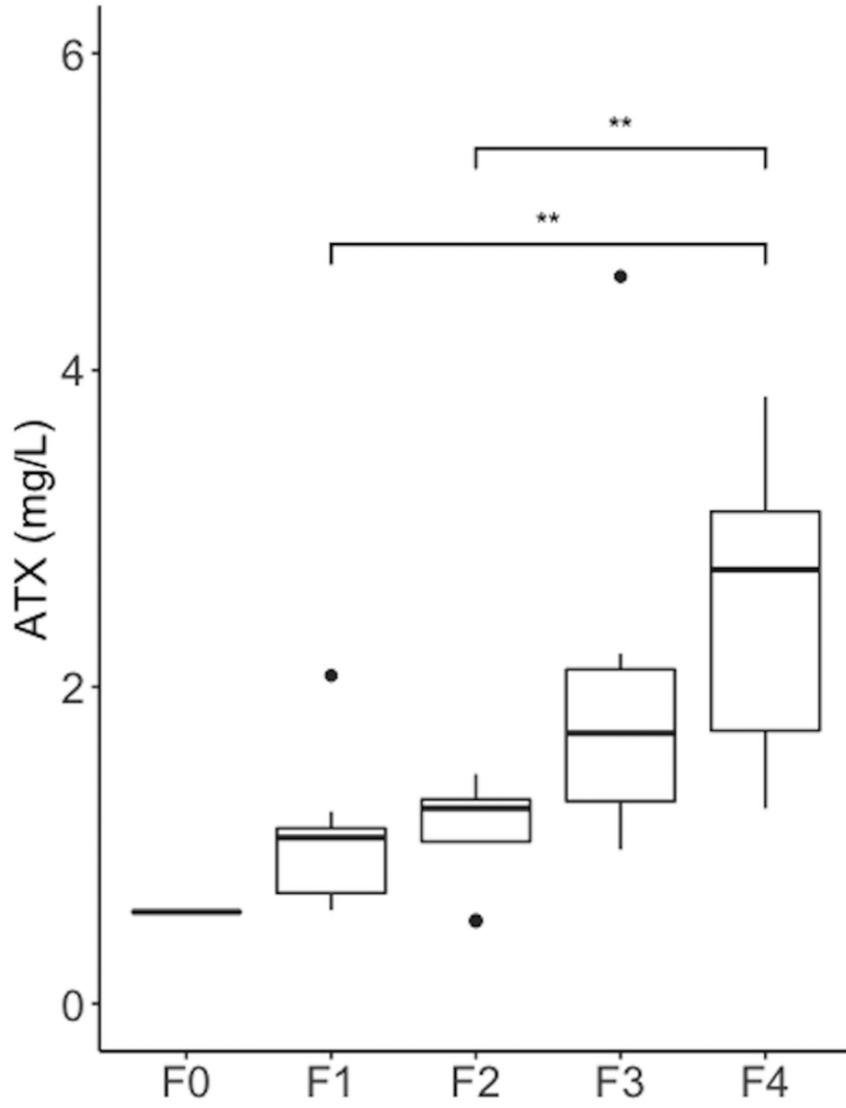


Figure 2B

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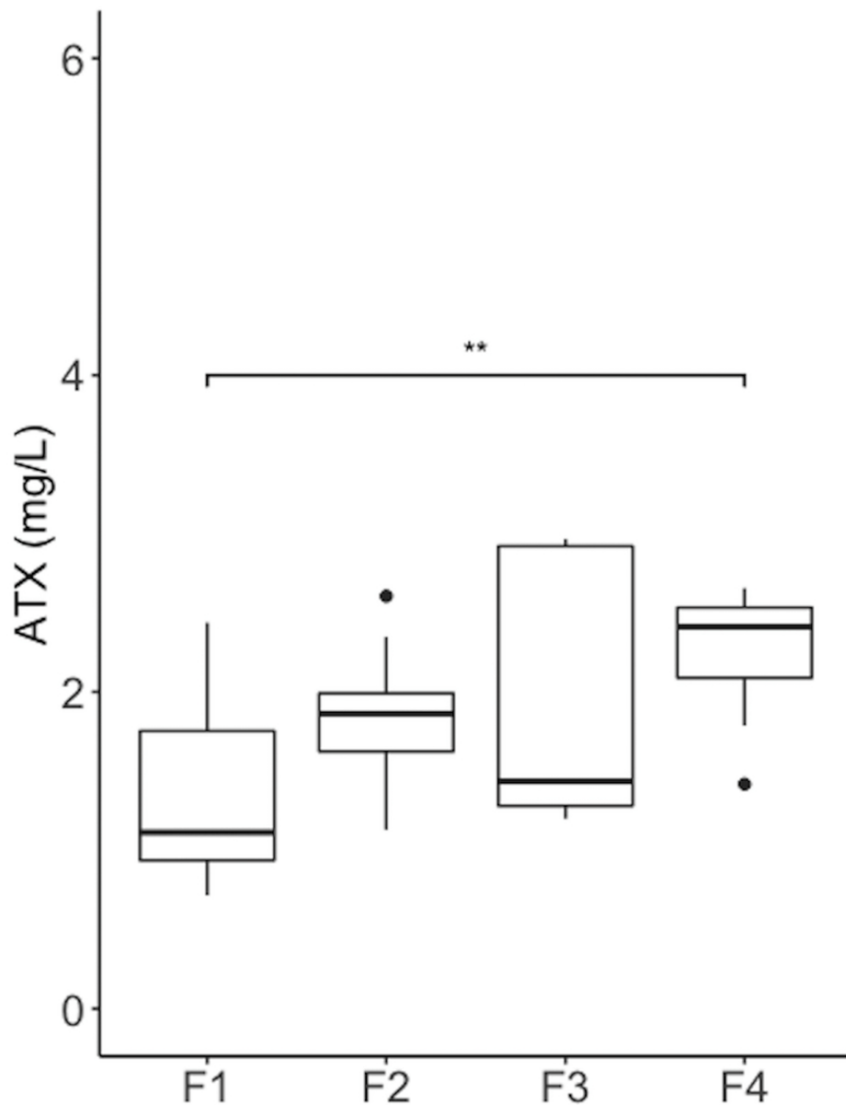


Figure 2C

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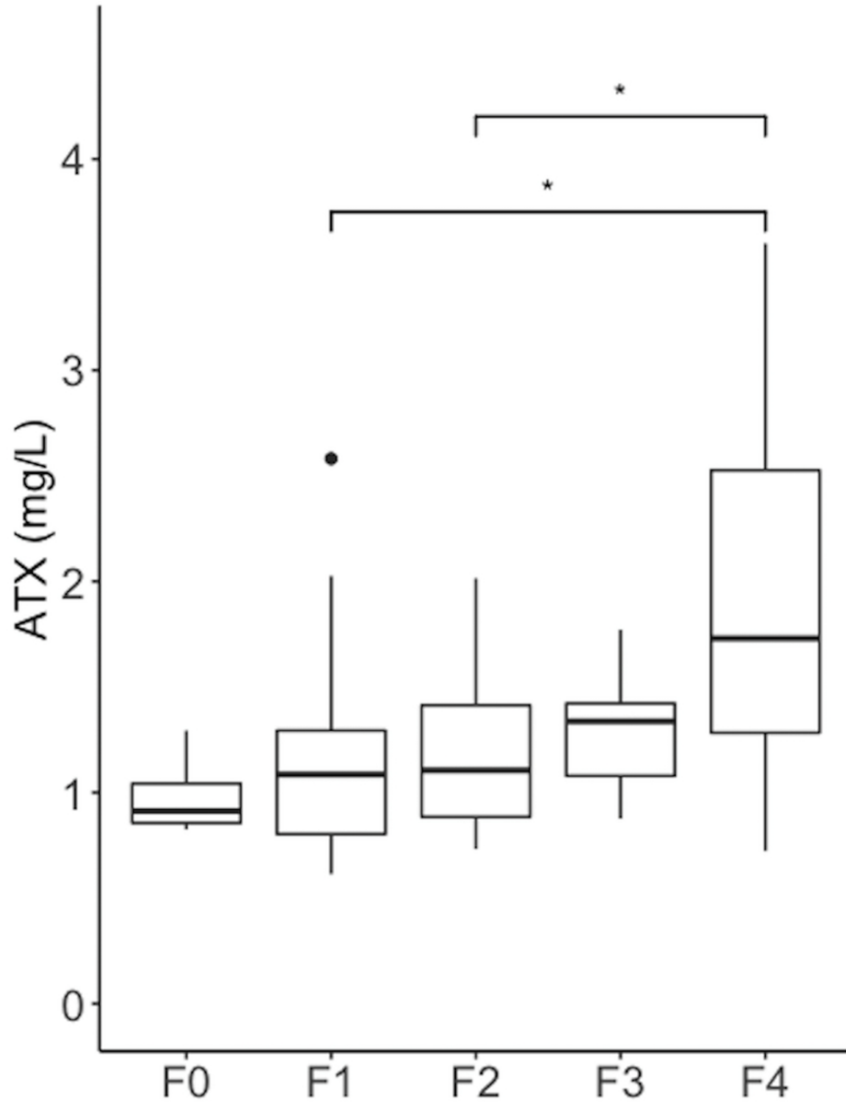


Figure 3A

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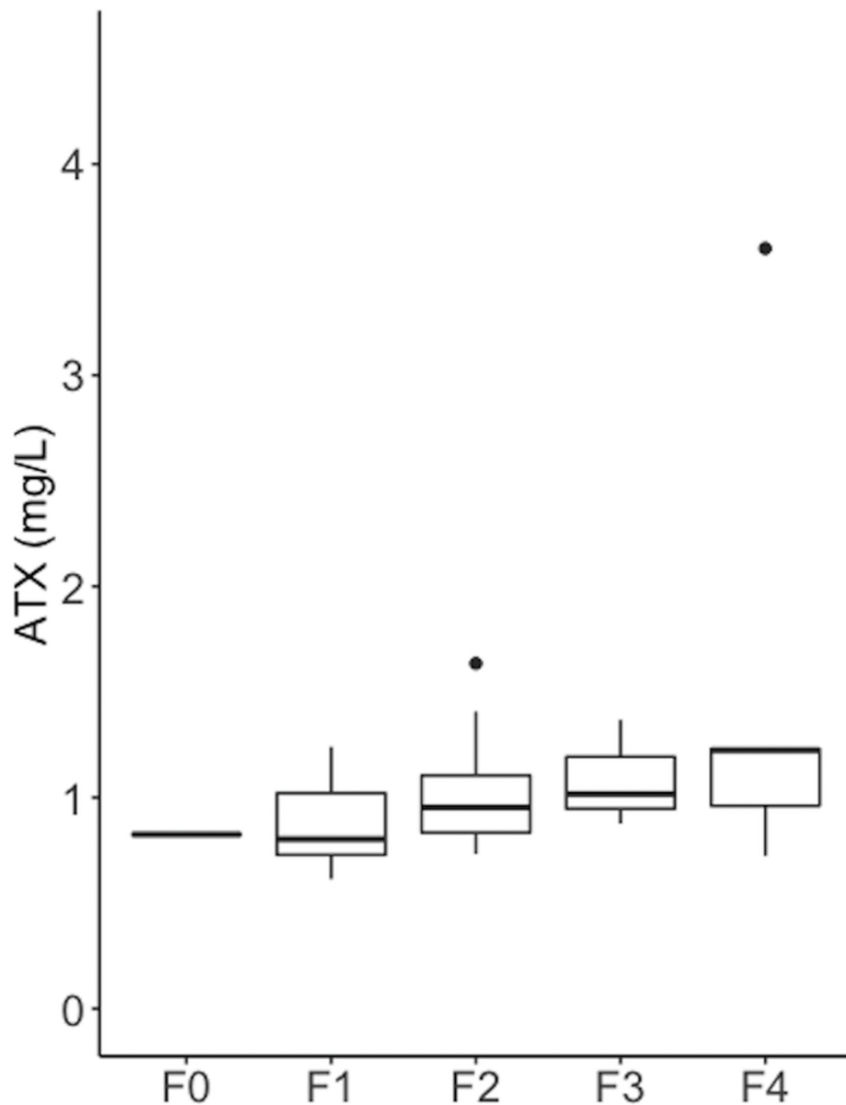


Figure 3B

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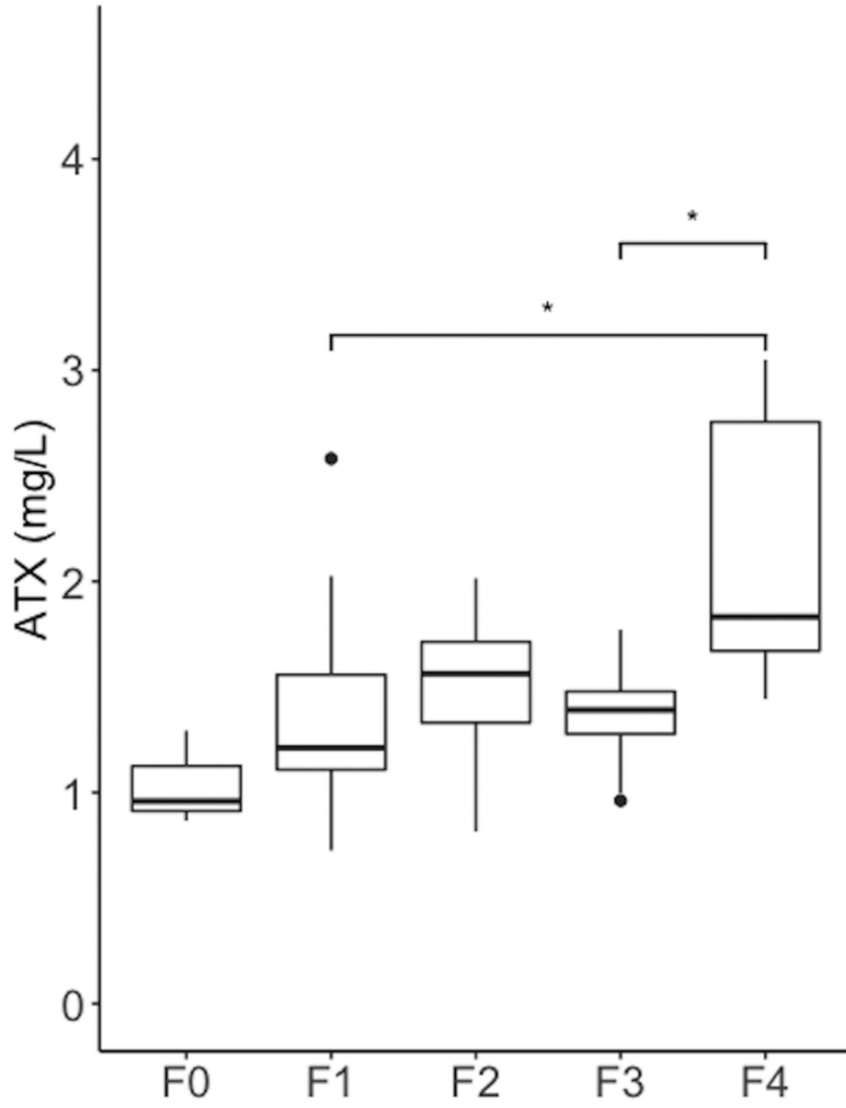


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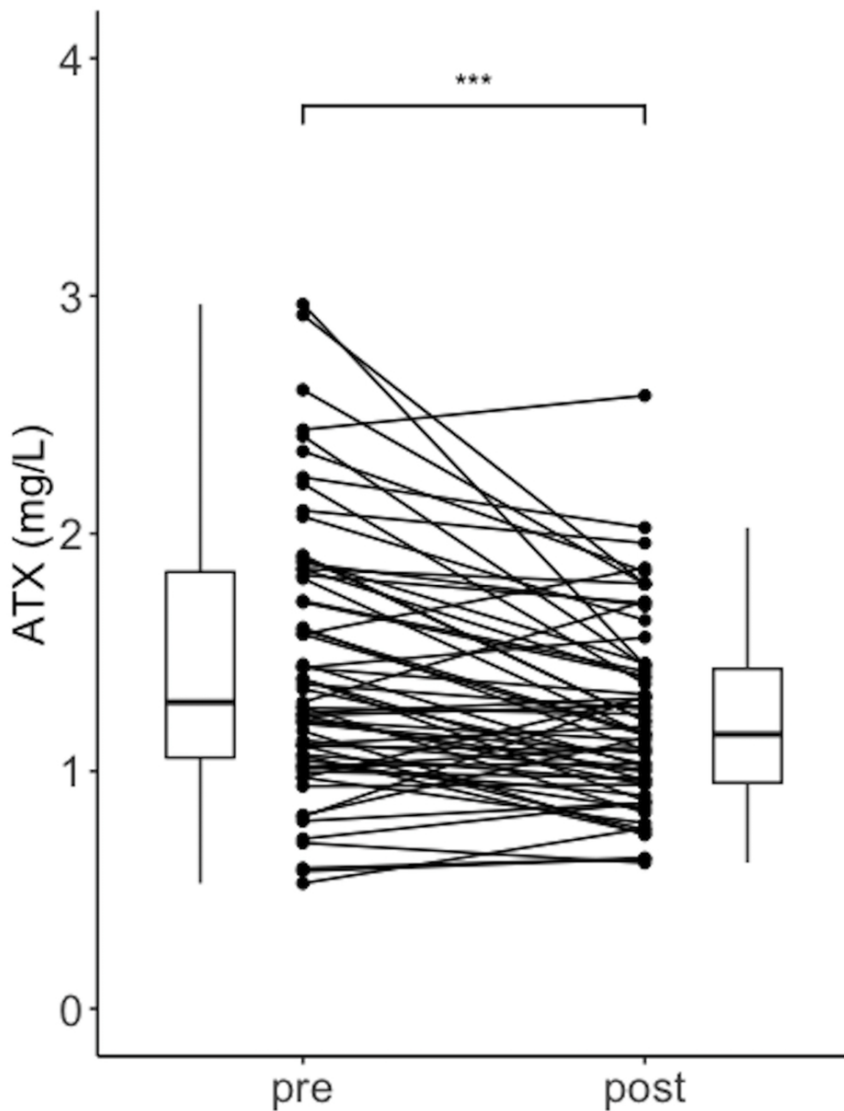


Figure 4A

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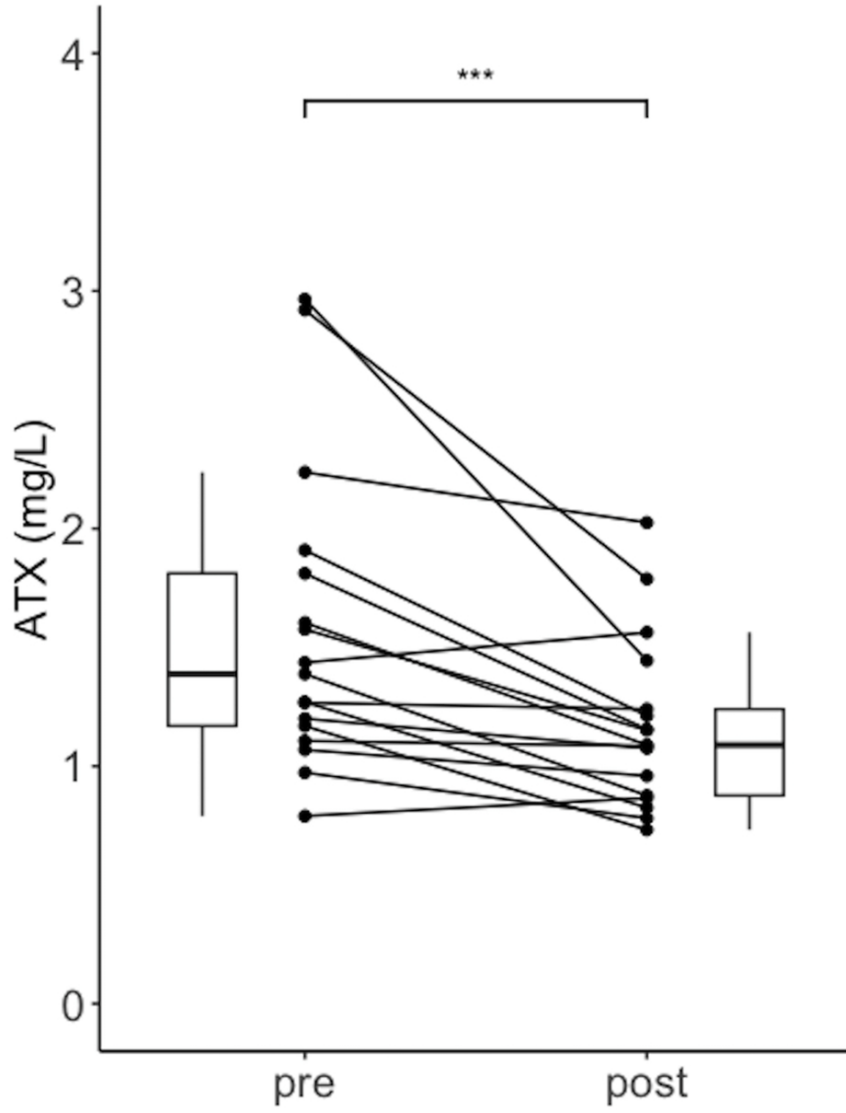


Figure 4B

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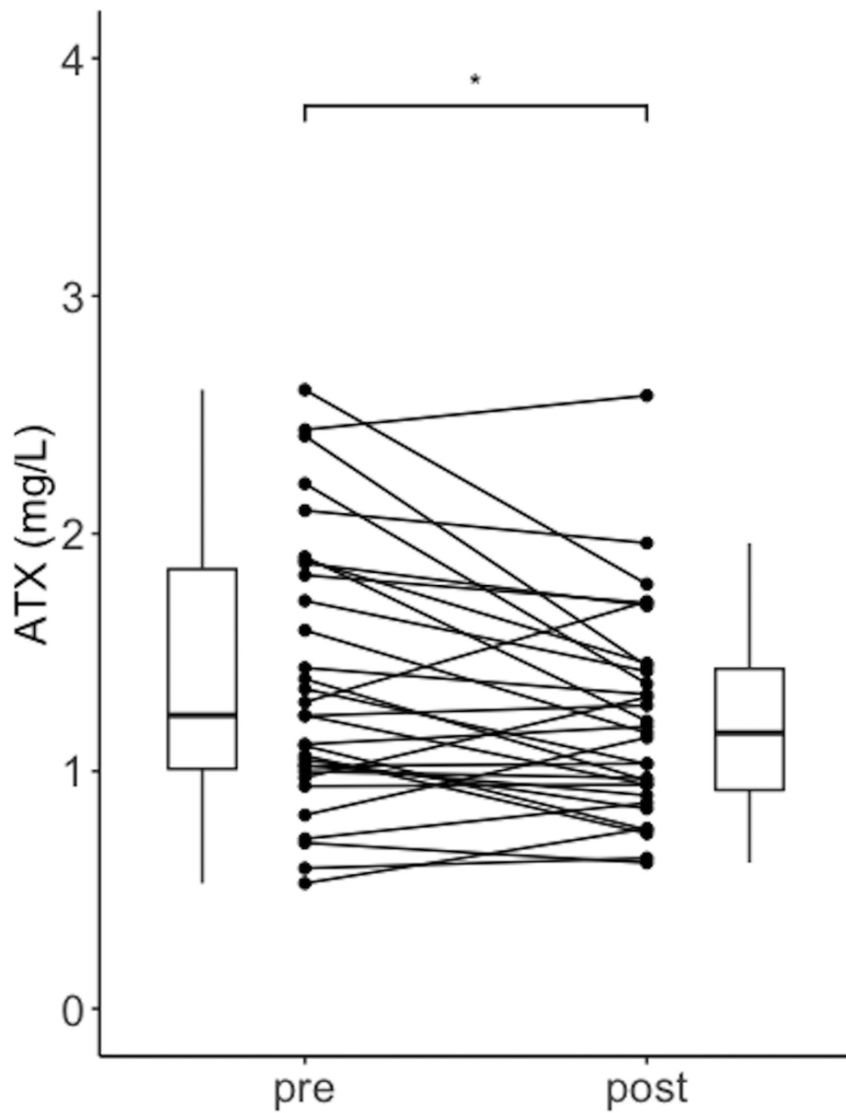


Figure 4C

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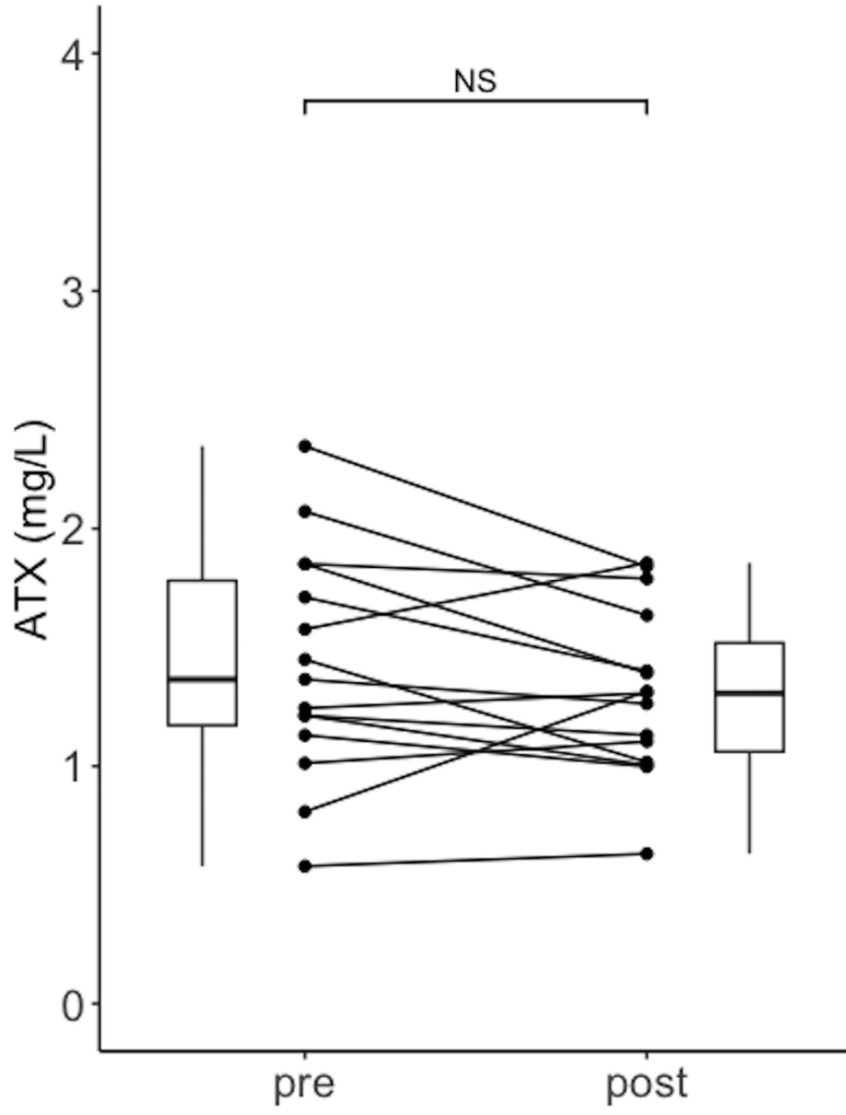


Figure 4D

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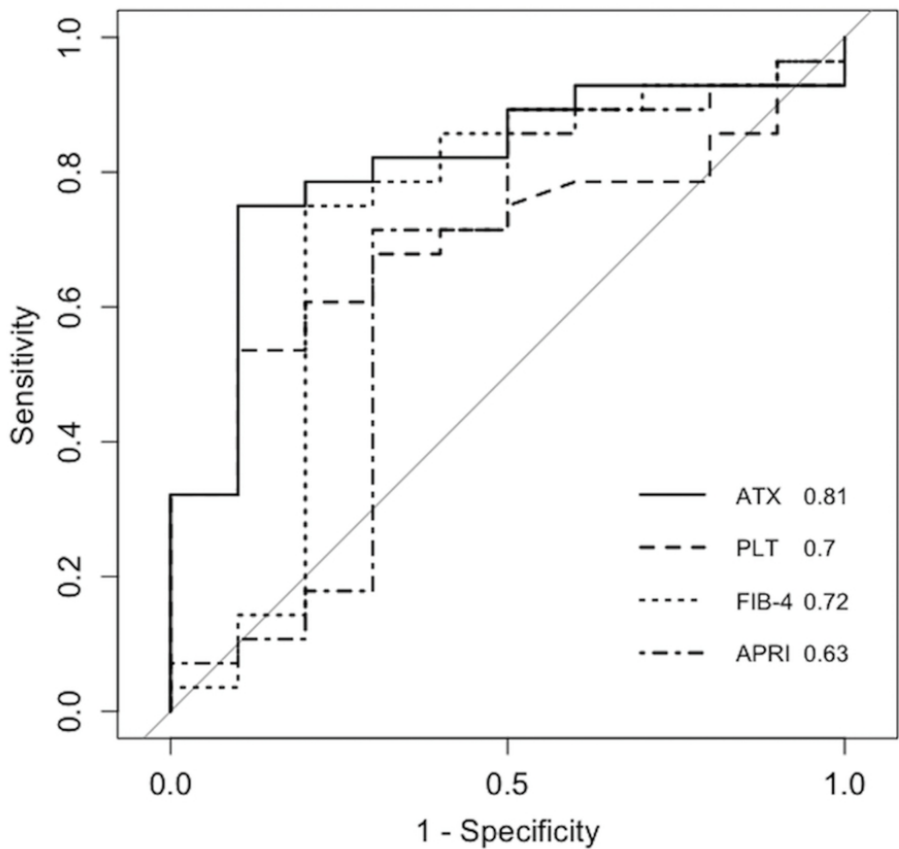


Figure 5A

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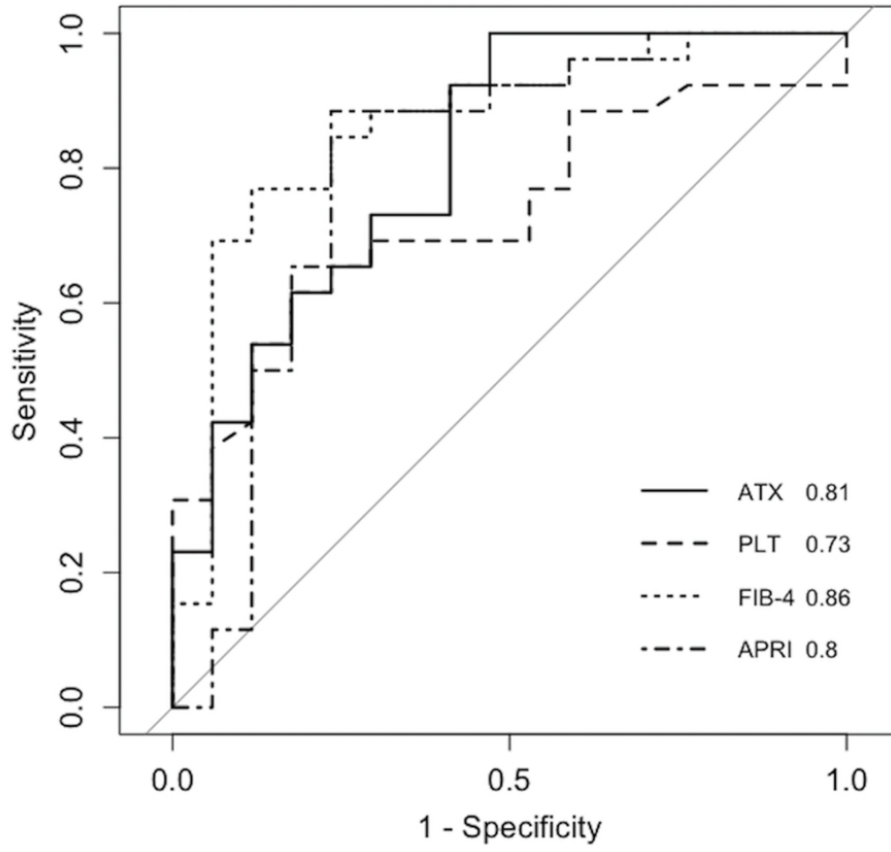


Figure 5B

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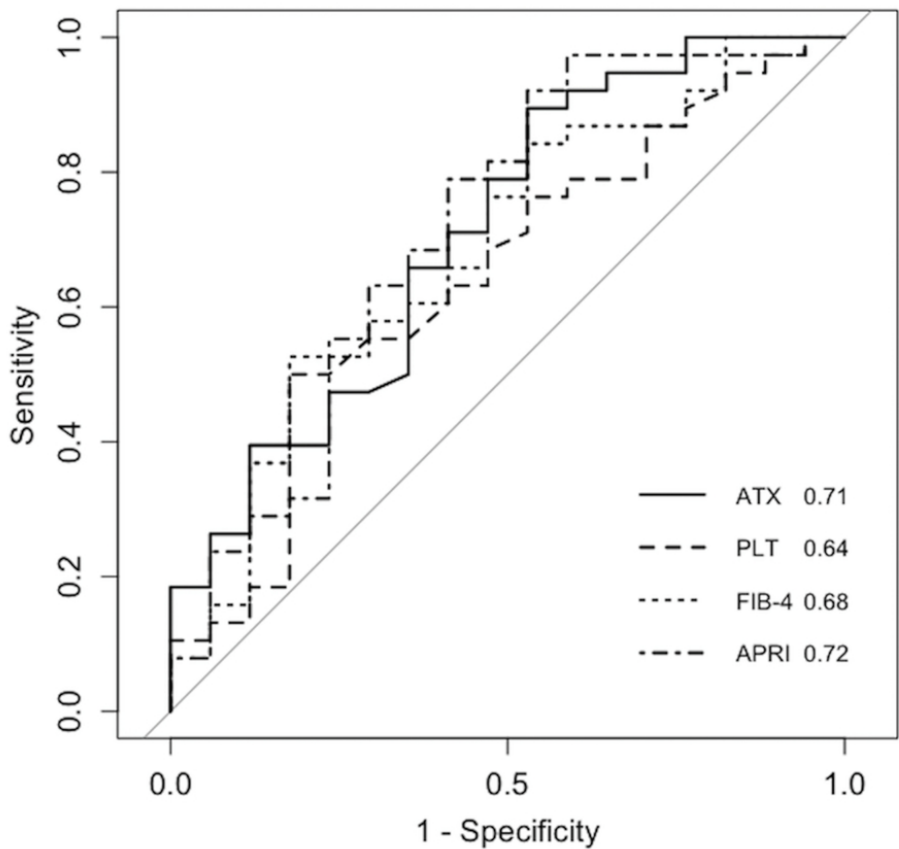


Figure 5C

79x75mm (300 x 300 DPI)

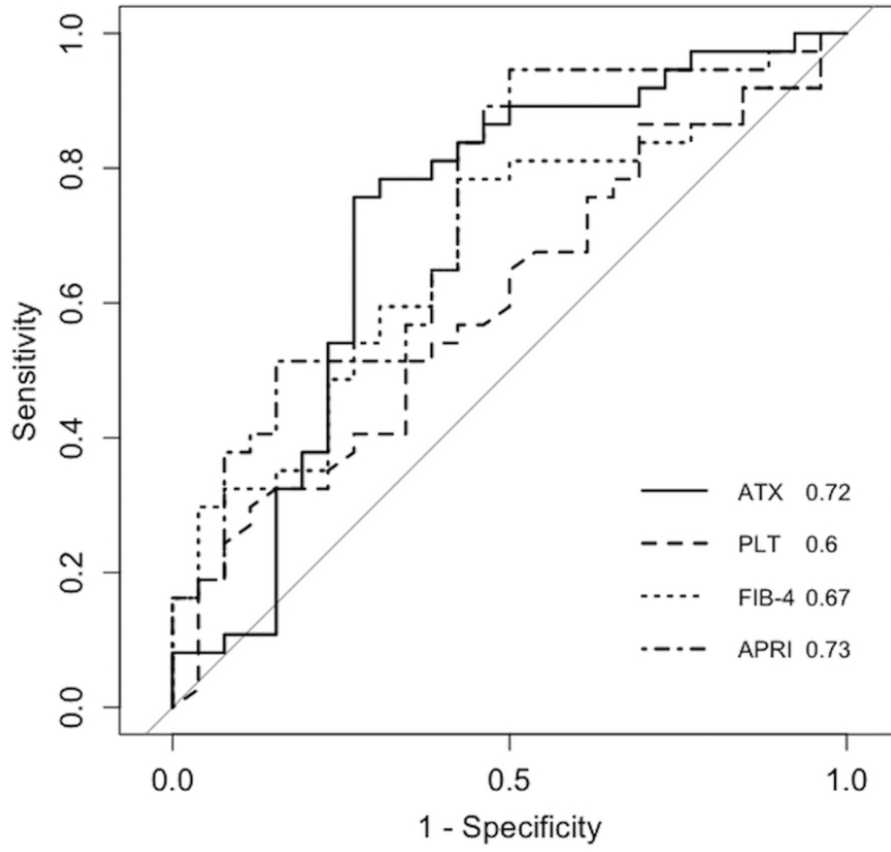
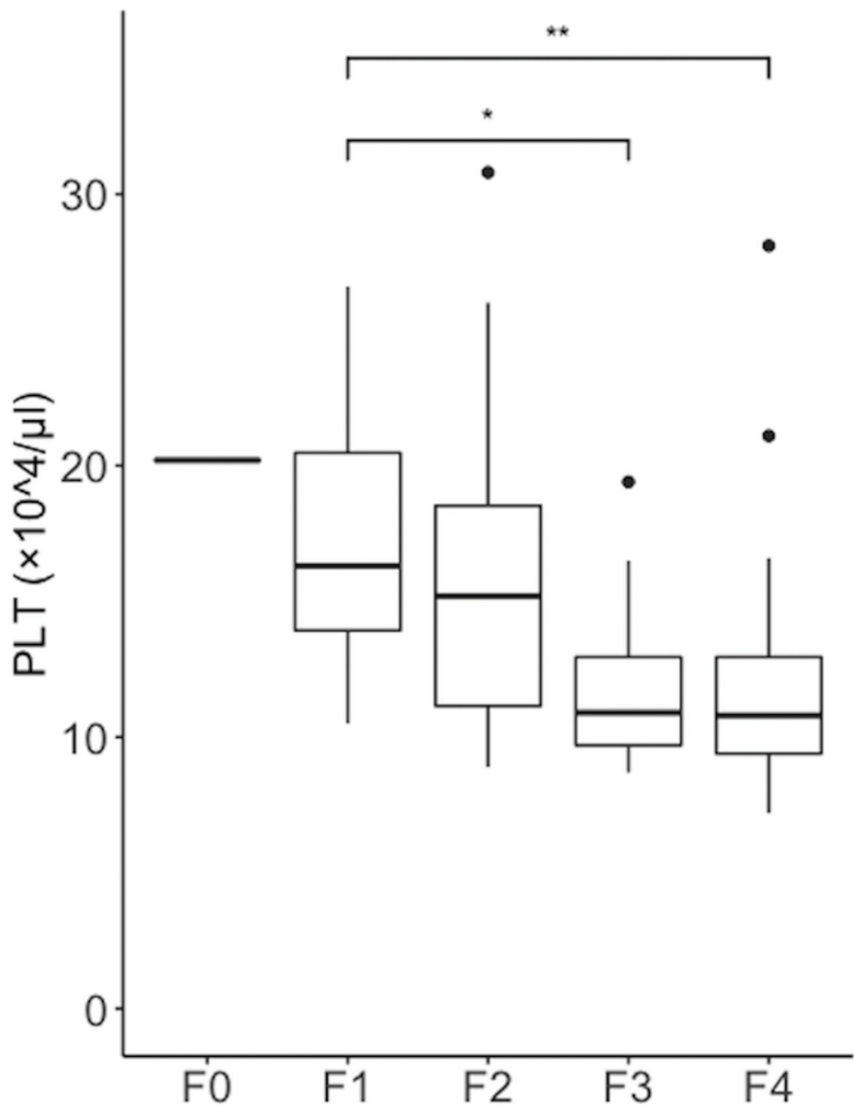


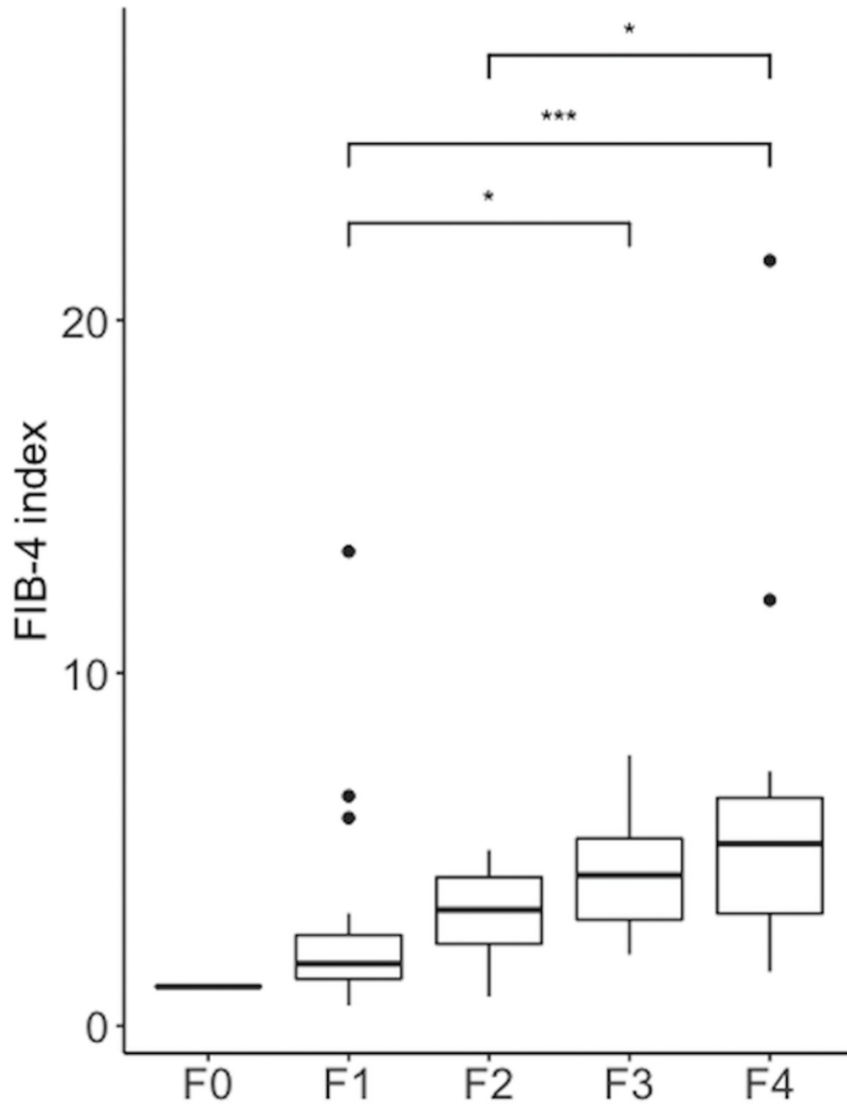
Figure 5D

79x75mm (300 x 300 DPI)



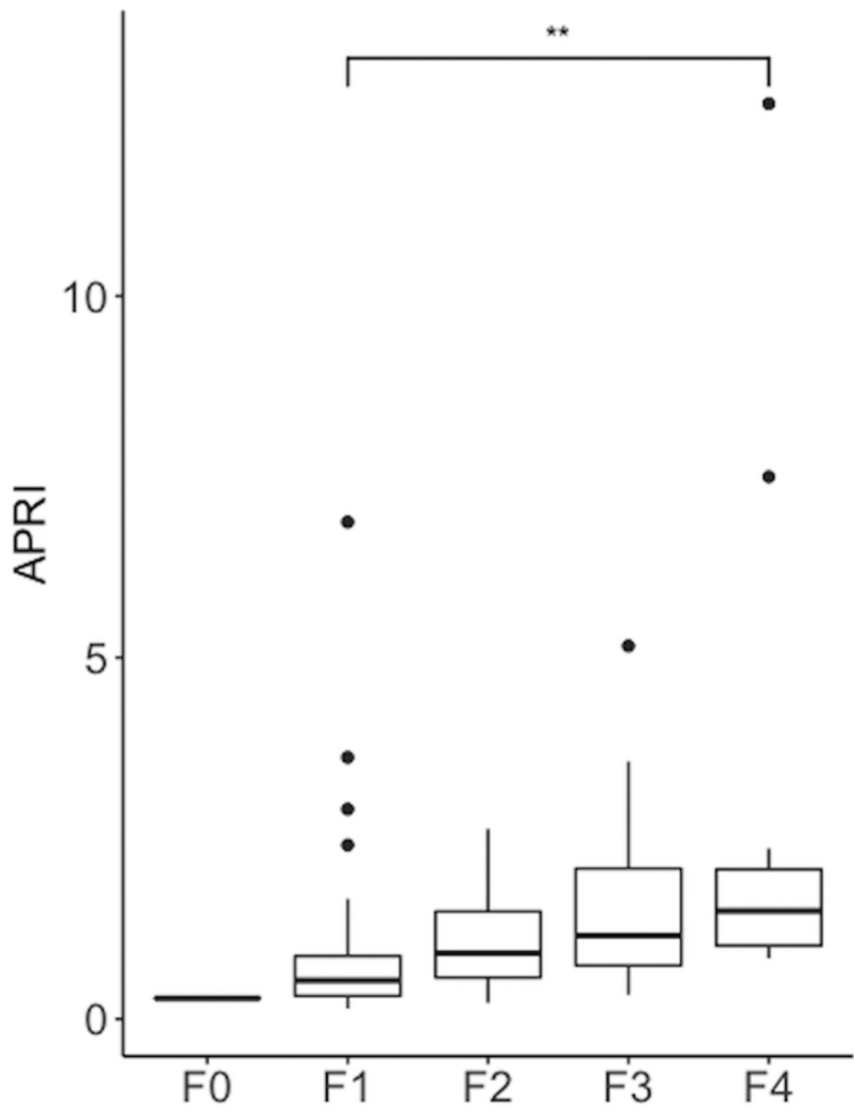
Supplemental Figure 1A

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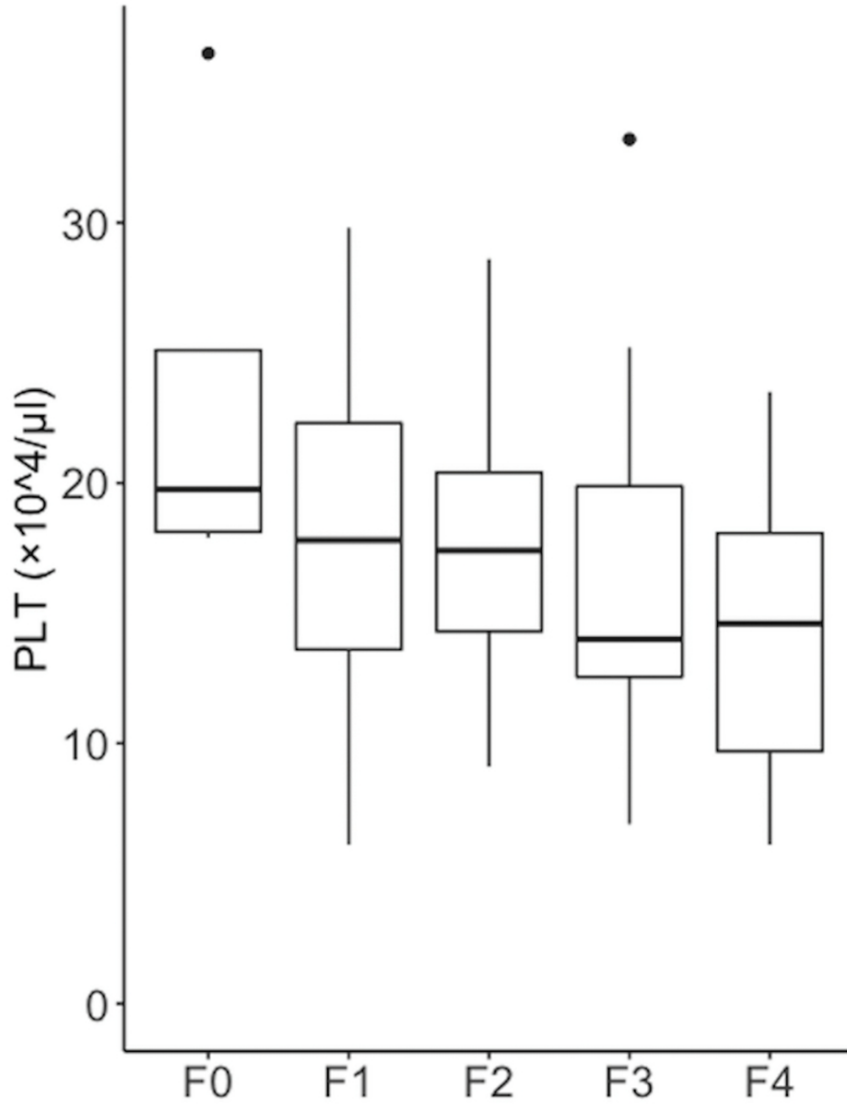
Supplemental Figure 1B

80x107mm (300 x 300 DPI)



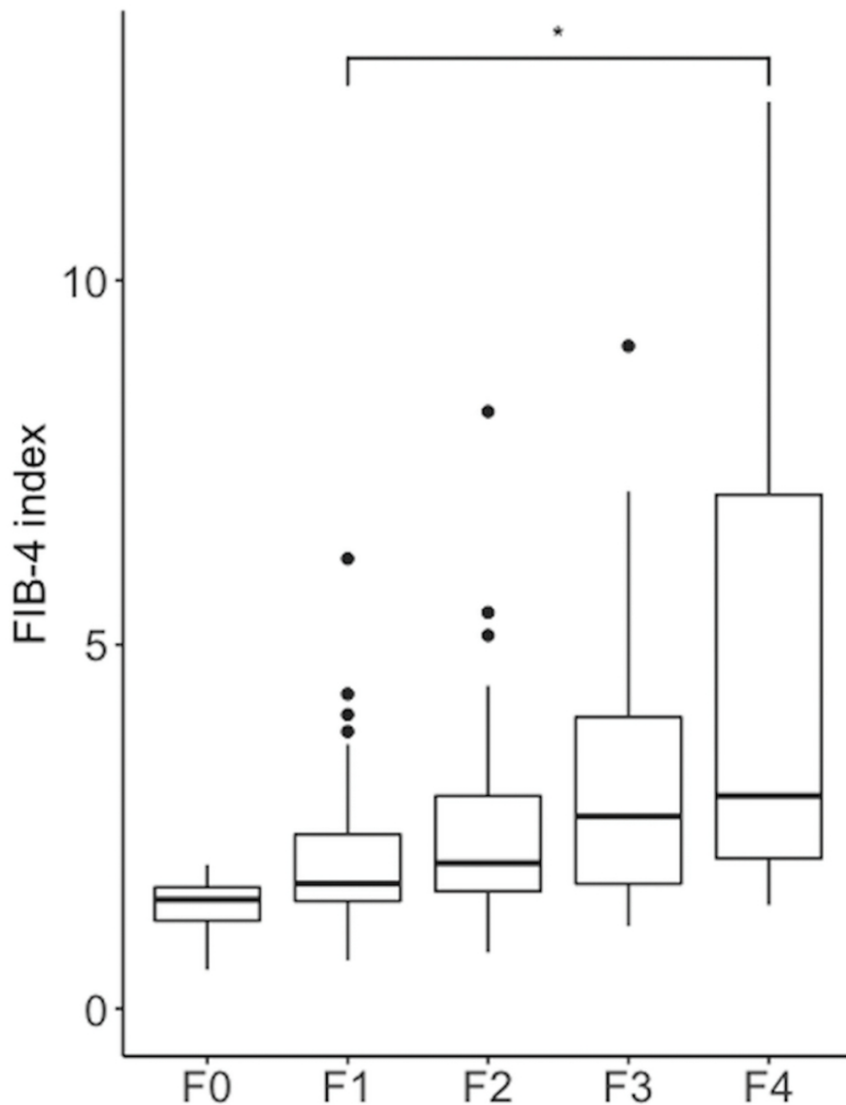
Supplemental Figure 1C

80x107mm (300 x 300 DPI)



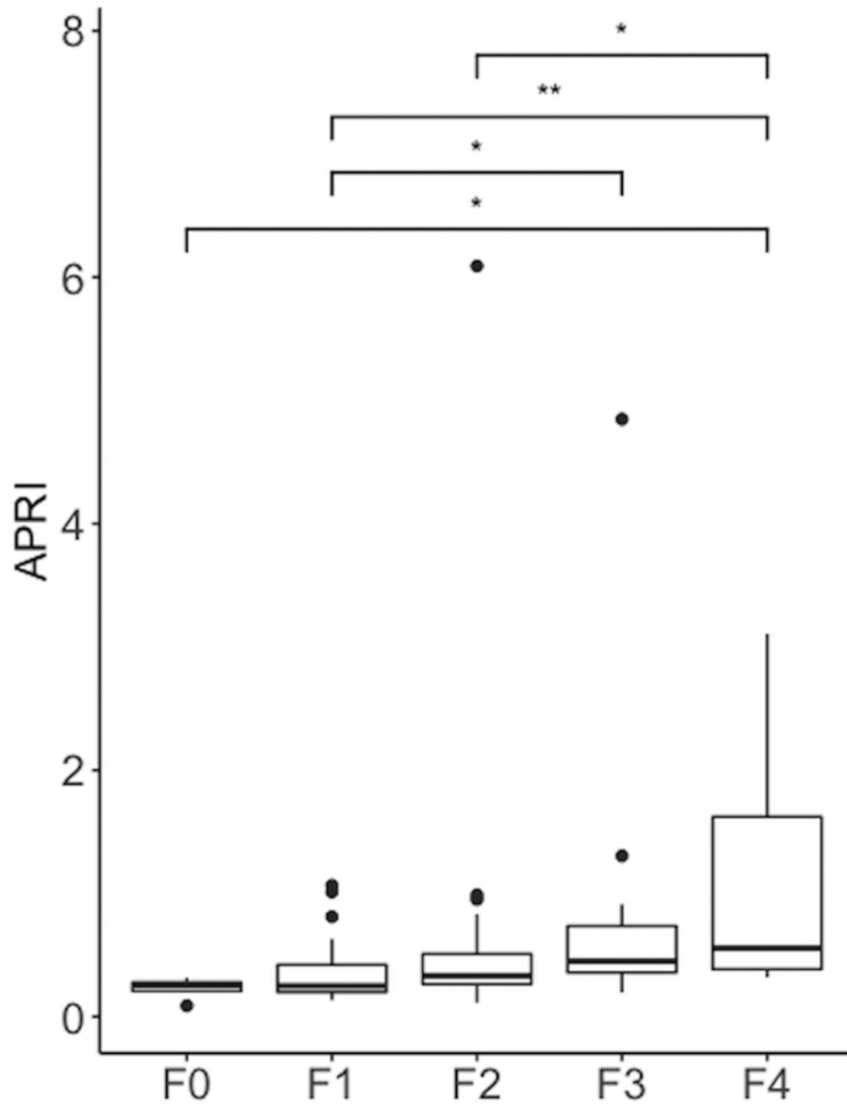
Supplemental Figure 2A

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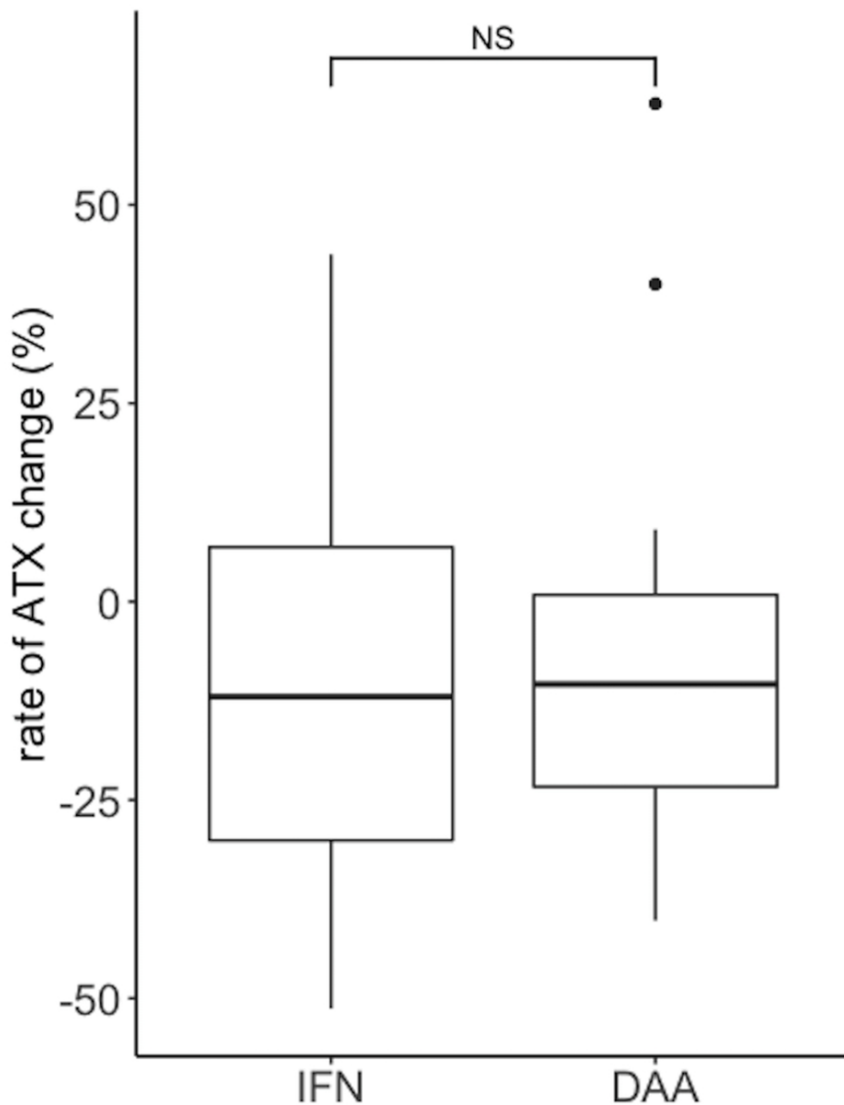
Supplemental Figure 2B

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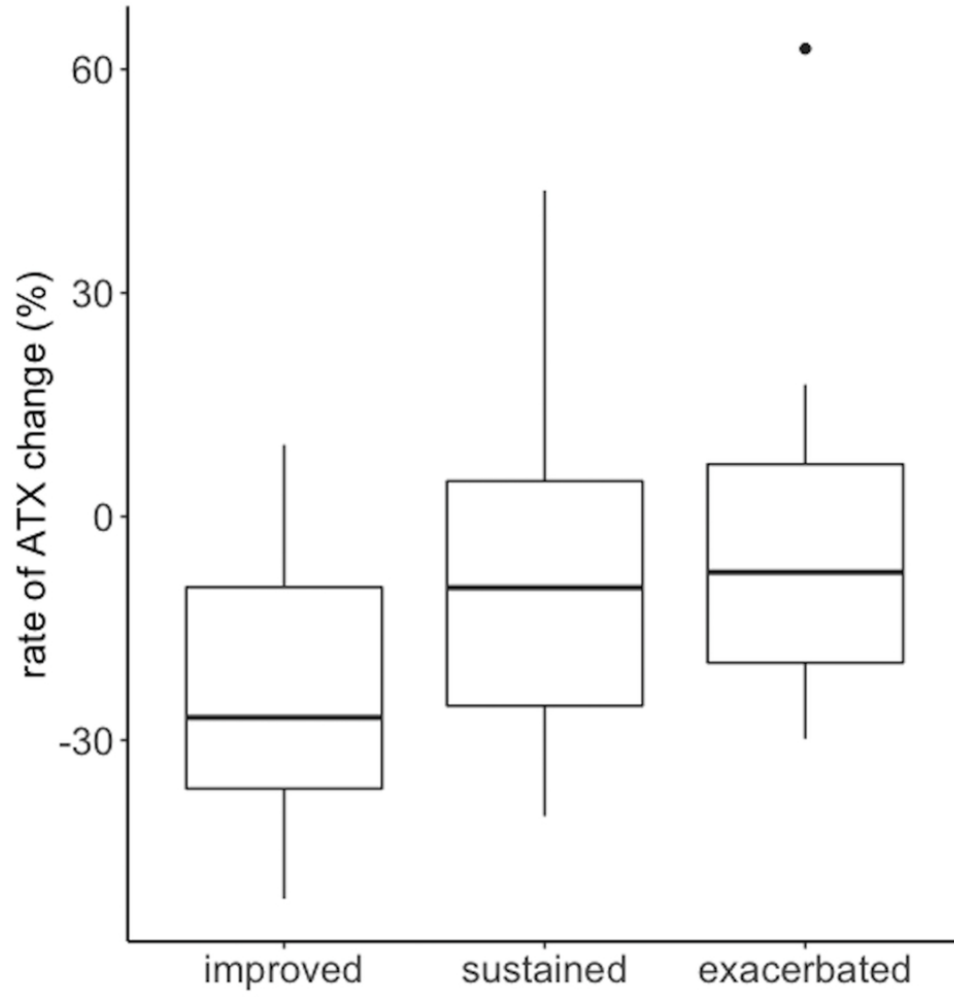
Supplemental Figure 2C

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Supplemental Figure 3

80x107mm (300 x 300 DPI)



Supplemental Figure 4

79x85mm (300 x 300 DPI)