

Non-estrogenic Xanthohumol Derivatives Mitigate Insulin Resistance and Cognitive Impairment in High-Fat Diet-induced Obese Mice

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Supplementary Information: Effect of xanthohumol (XN), α,β -dihydroxanthohumol (DXN), or tetrahydroxanthohumol (TXN) on the viability of C2C12 cells (Figure S1a) and of HepG2 cells (Figure S1b); Structural characterization of DXN by ¹H-NMR (Figure S2) and ¹³C-NMR spectroscopy (Figure S3) and by electrospray mass spectrometry (Figure S4); Structural characterization of TXN by ¹H-NMR (Figure S5) and ¹³C-NMR spectroscopy (Figure S6) and by electrospray mass spectrometry (Figure S7); The MS/MS spectra provided in Figures S4 and S7 were recorded on a Sciex 4000 QTrap mass spectrometer (see Methods). Supplementary Figure S8 contains full-length Western blots of liver and muscle biopsies from untreated mice (control, lanes 1-2) and treated with XN (lanes 3-4), DXN (lanes 5-6) and TXN (lanes 7-8). Tissue extracts were incubated with specific antibodies against pAMPK, AMPK and β -actin. These original images were used to generate Figure 7. No exposure modifications were recorded.

Figure S1a. Exposure of C2C12 cells to XN, DXN, or TXN for one hour at concentrations 1-50 μ M did not affect viability ($p > 0.05$ by ANOVA). Cell viability of treated cells is displayed as a percentage of vehicle treated cells (controls). Data are represented as mean $\pm$ SE of 4 replicate wells.

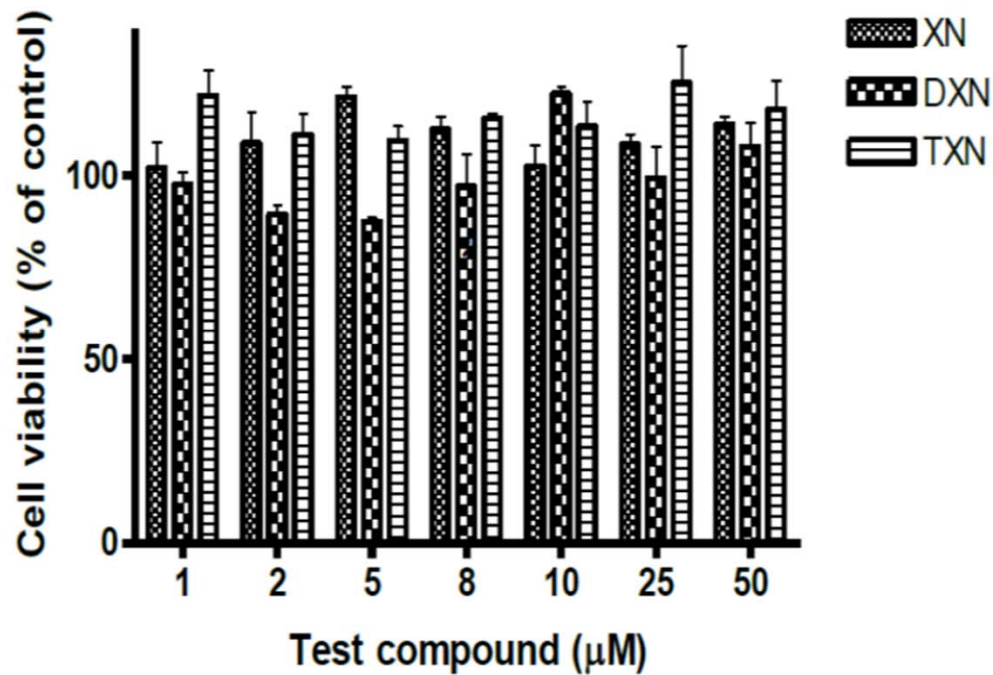


Figure S1b. Exposure of HepG2 cells to XN, DXN, or TXN for 24 h did not affect viability at concentrations up to 25 μM ($p > 0.05$ by ANOVA). However, at 50 μM , all the test compounds were cytotoxic. Cell viability of treated cells is displayed as a percentage of vehicle treated cells (controls). Data are represented as mean $\pm$ SE of 4 replicate wells.

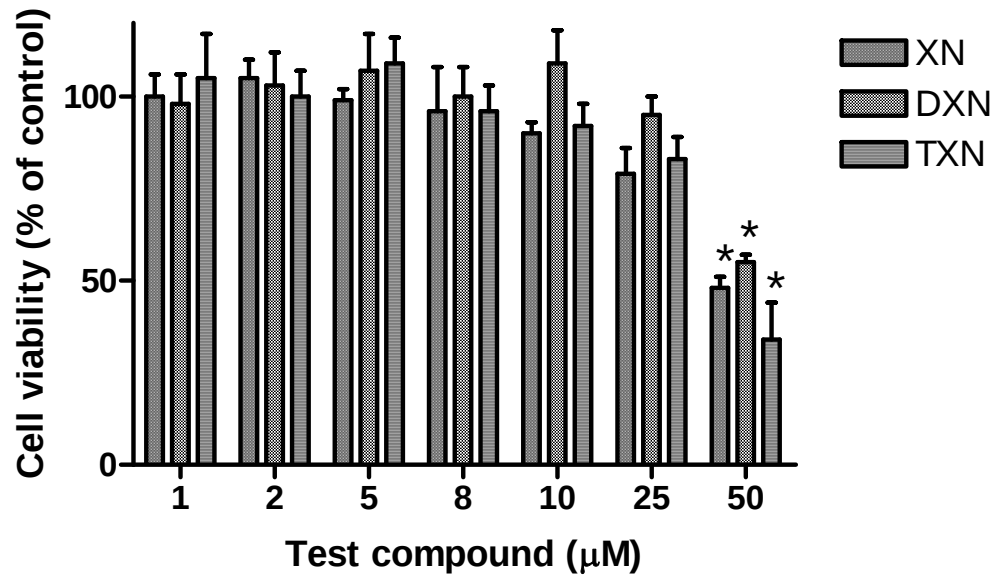


Figure S2. ^1H -NMR (700 MHz) spectrum of α,β -dihydroxanthohumol (DXN) recorded in methanol- d_4 .

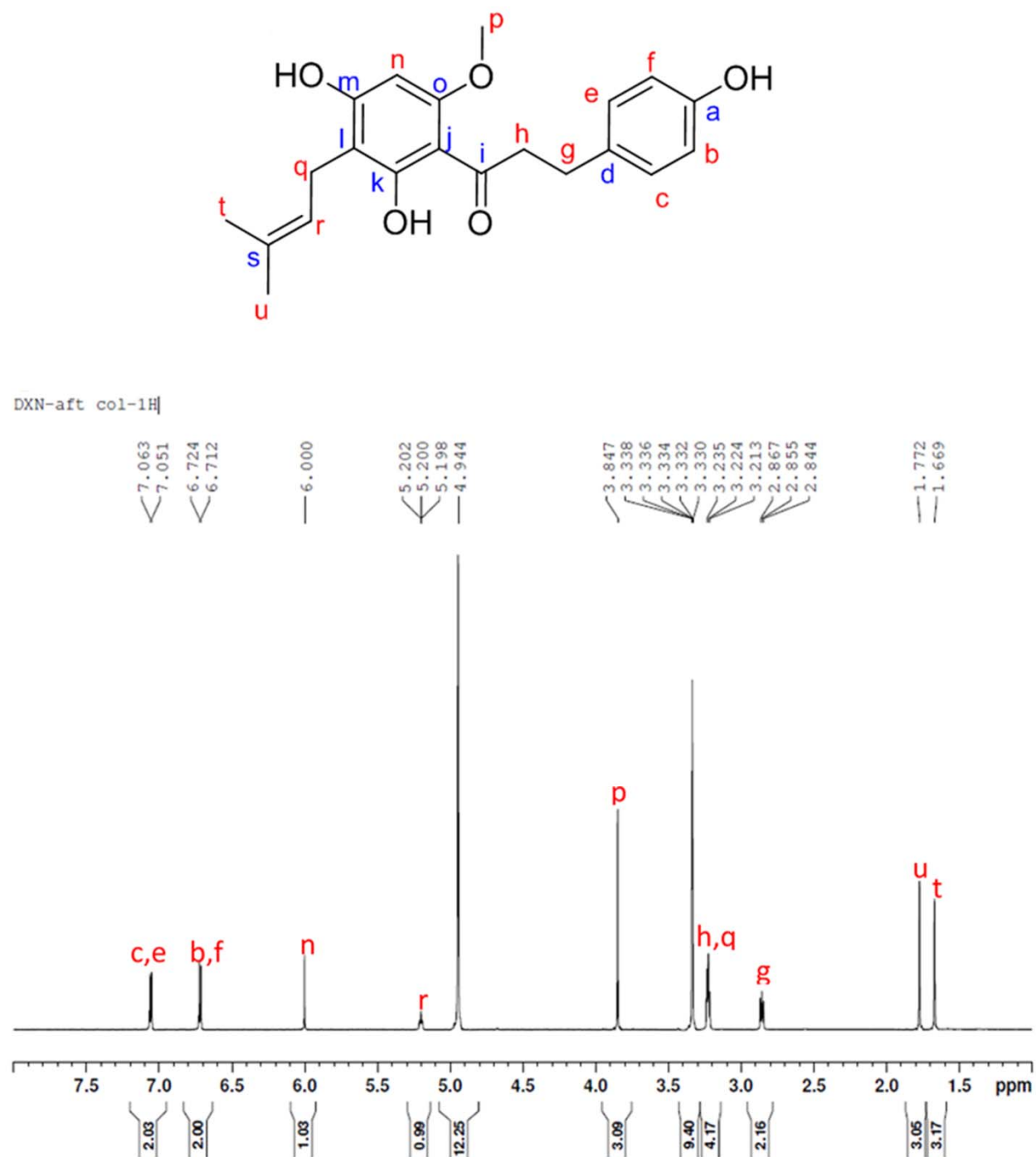


Figure S3. ^{13}C -NMR (175 MHz) spectrum of α,β -dihydroxanthohumol (DXN) recorded in **methanol- d_4** . For atom labels, see Figure S2. ^{13}C signals were assigned on the basis of HSQC and HMBC experiments.

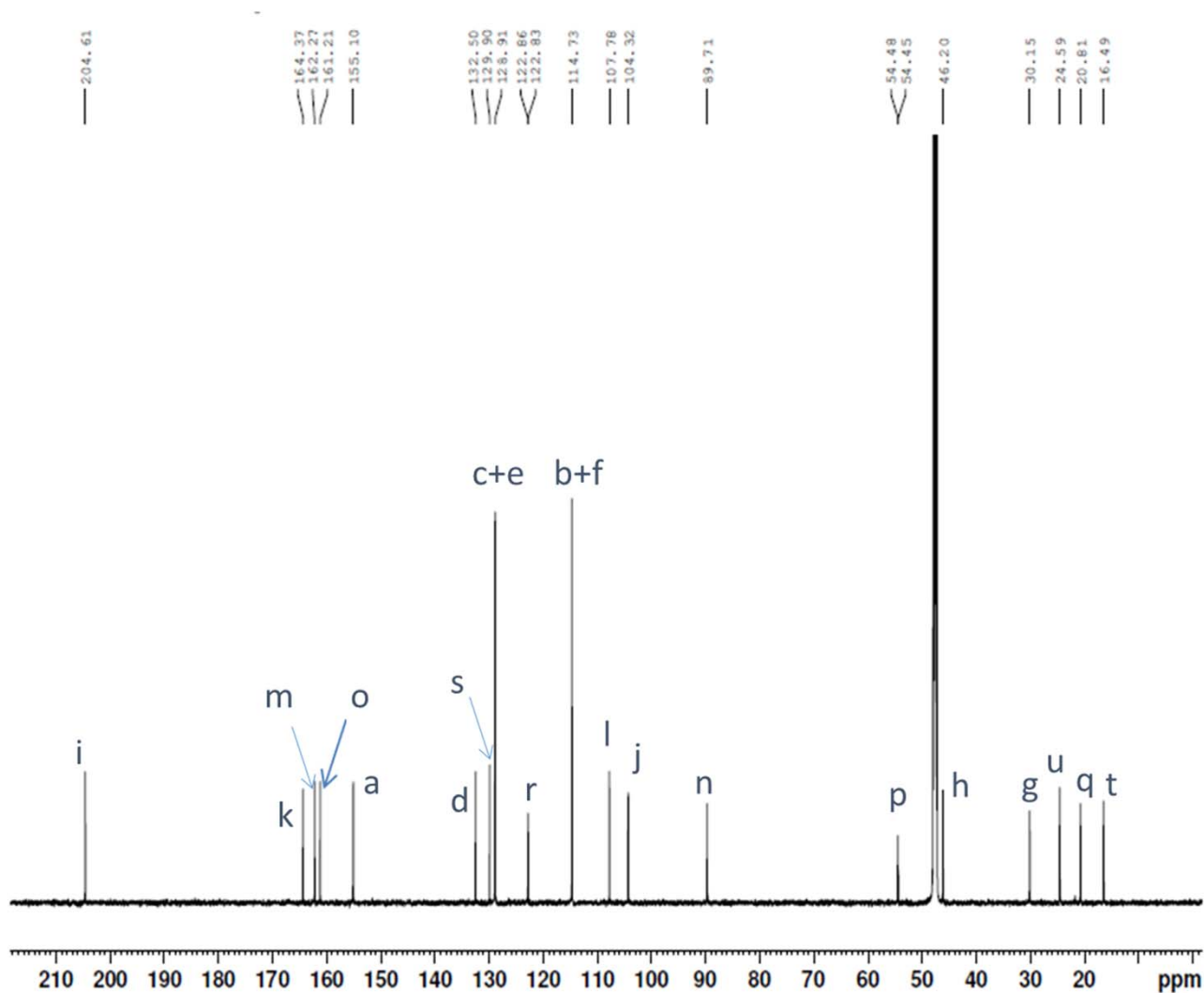
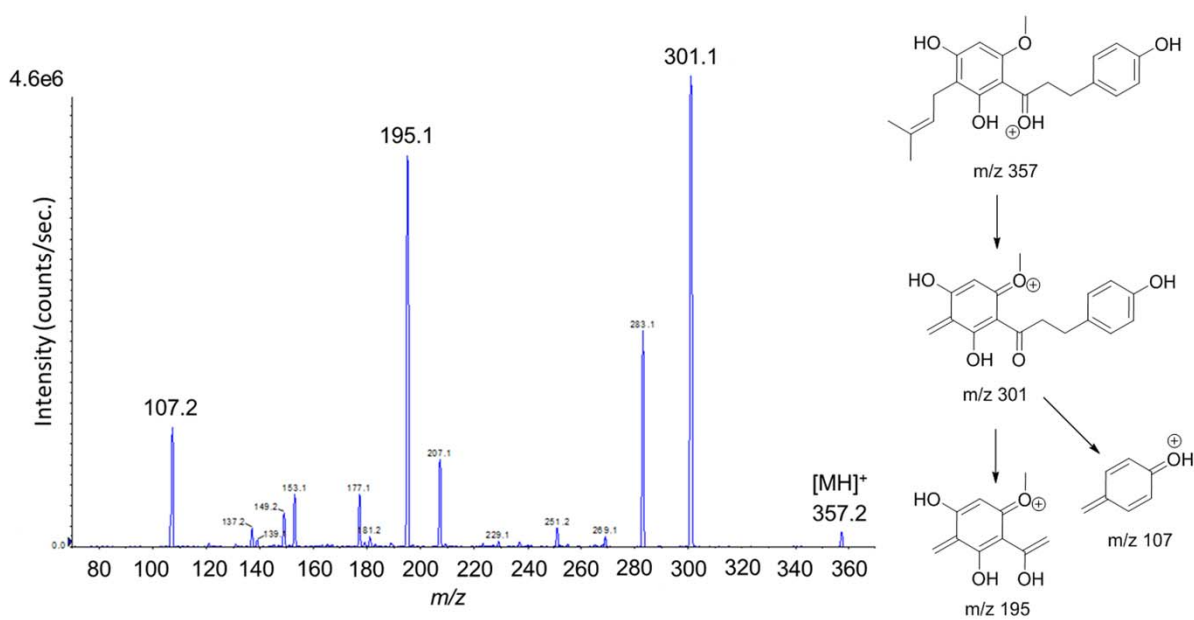


Figure S4. Electrospray MS/MS spectrum of α,β -dihydroxanthohumol (DXN) and its proposed fragmentation pathway.



The chemical structure of compound 10 is shown at the top, with protons labeled with letters: a, b, c, d, e, f, g, h, i, j, k, l, m, n, o, p, q, r, s, t, u. The ¹H NMR spectrum (400 MHz, CDCl₃) is displayed below, with peaks assigned to these labels. The x-axis represents the chemical shift in ppm, ranging from 0.0 to 8.0. Integration values are provided below the baseline for each group of peaks.

Chemical structure of compound 10: CC(C)C[C@H](O)[C@@H](OC(=O)C[C@H](O)c1ccc(O)cc1)c2cc(O)c(OC)c(O)c2

¹H NMR spectrum (400 MHz, CDCl₃) showing peaks assigned to protons in compound 10:

- Peak assignments: c+e (~7.0 ppm), b+f (~6.7 ppm), n (~6.0 ppm), unlabeled (~4.8 ppm), p (~3.8 ppm), h (~3.3 ppm), g (~3.1 ppm), q (~2.6 ppm), s (~1.5 ppm), r (~1.3 ppm), t+u (~0.9 ppm).
- Integration values (from left to right): 1.94, 1.92, 0.98, 2.94, 1.95, 1.95, 1.95, 0.99, 1.97, 0.22, 5.88.

Figure S6. ^{13}C -NMR (175 MHz) spectrum of tetrahydroxanthohumol (TXN) recorded in **methanol- d_4** . For atom labels, see Figure S5. ^{13}C signals were assigned on the basis of HSQC and HMBC experiments.

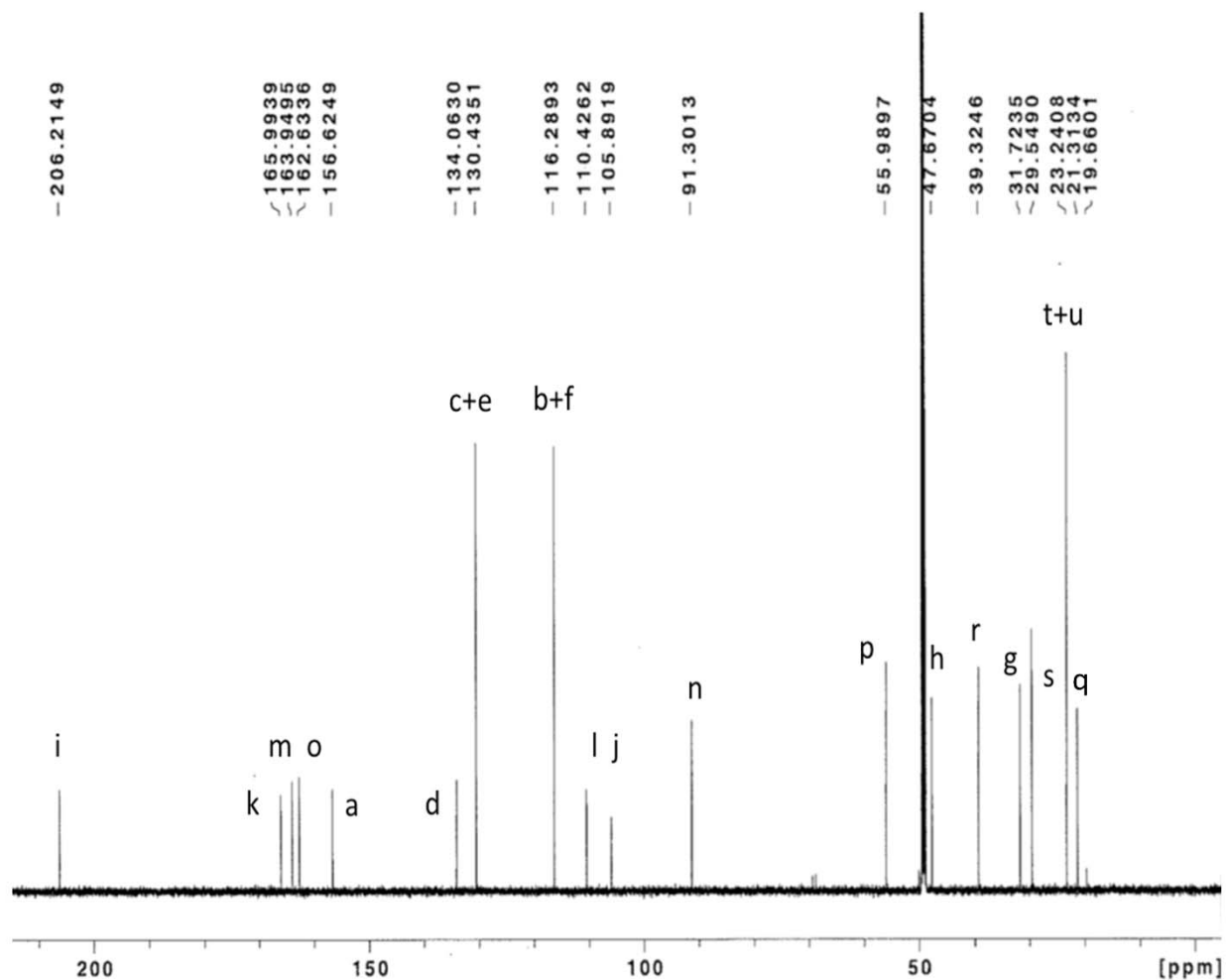


Figure S7. Electrospray MS/MS spectrum of tetrahydroxanthohumol (TXN) and its proposed fragmentation pathway.

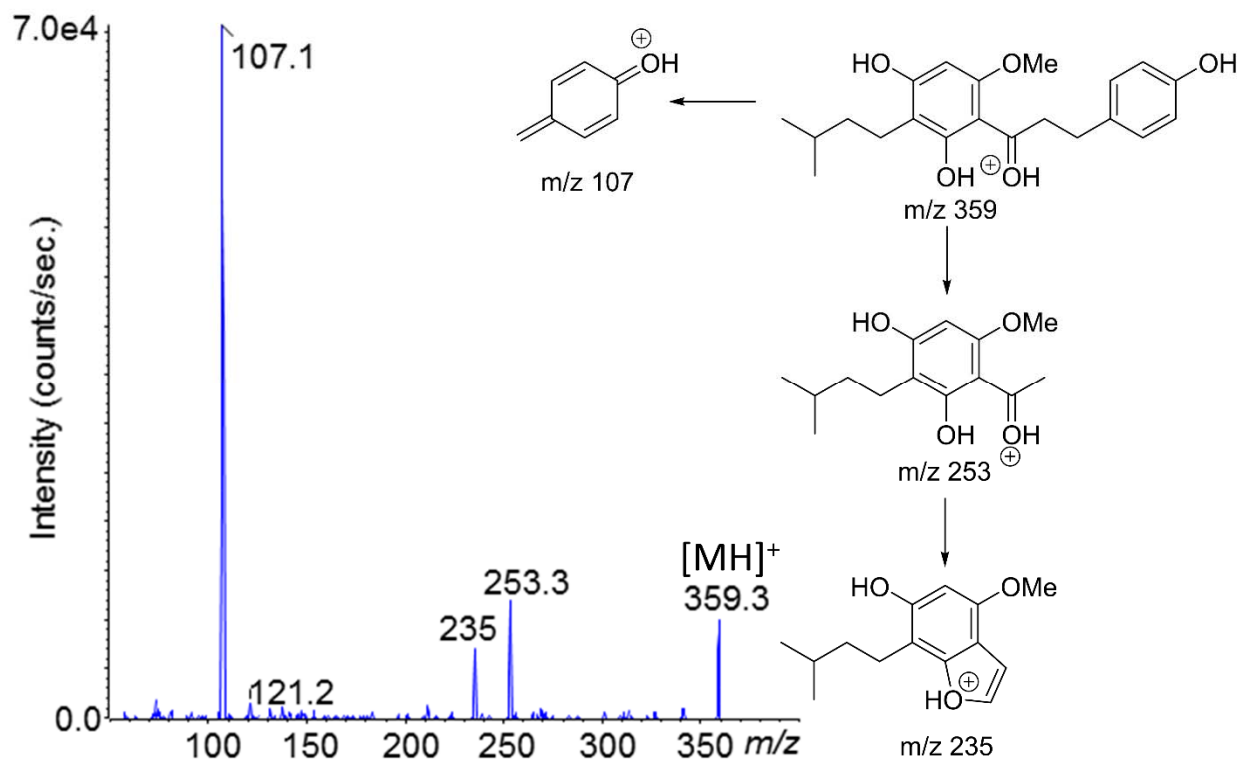


Figure S8. Full-length Western blots of liver and muscle biopsies from untreated mice (control, lanes 1-2) and treated with XN (lanes 3-4), DXN (lanes 5-6) and TXN (lanes 7-8). Tissue extracts were incubated with specific antibodies against pAMPK, AMPK and β -actin. These original images were used to generate Figure 7. No exposure modifications were recorded.

