

***Urtica dioica* L. inhibits proliferation and enhances cisplatin cytotoxicity in NSCLC cells via Endoplasmic Reticulum-stress mediated apoptosis**

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Figure S1. Western Blot analyses of the proteins involved in apoptosis of NSCLC (H1299 and A549) cell lines, normal bronchial epithelial (Beas2b) cells and human fibroblasts (Wi38)

Figure S2. The expression levels of ER-stress related proteins

Figure S3. CIGAR experiment of *Urtica dioica* crude extract. R= rutin. OXY=oxylipins

Figure S4. HSQC-TOCSY experiment of *Urtica dioica* crude extract

Figure S5. DQCOSY experiment of *Urtica dioica* crude extract. R= rutin. OXY=oxylipins

Figure S6. HMBC experiment of rutin

Figure S7. ^1H -NMR spectrum of oxylipins' enriched fraction

Figure S8. H2BC experiment of oxylipins' enriched fraction

Figure S9. ESI Q-TOF MS spectrum of main oxylipin acquired in negative (A) and positive ion mode (B)

Figure S10. HSQC experiment of the main oxylipin

Table S1. NMR data of oxylipin recorded in CD_3OD . ^1H and ^{13}C are measured in ppm

Figure S1.

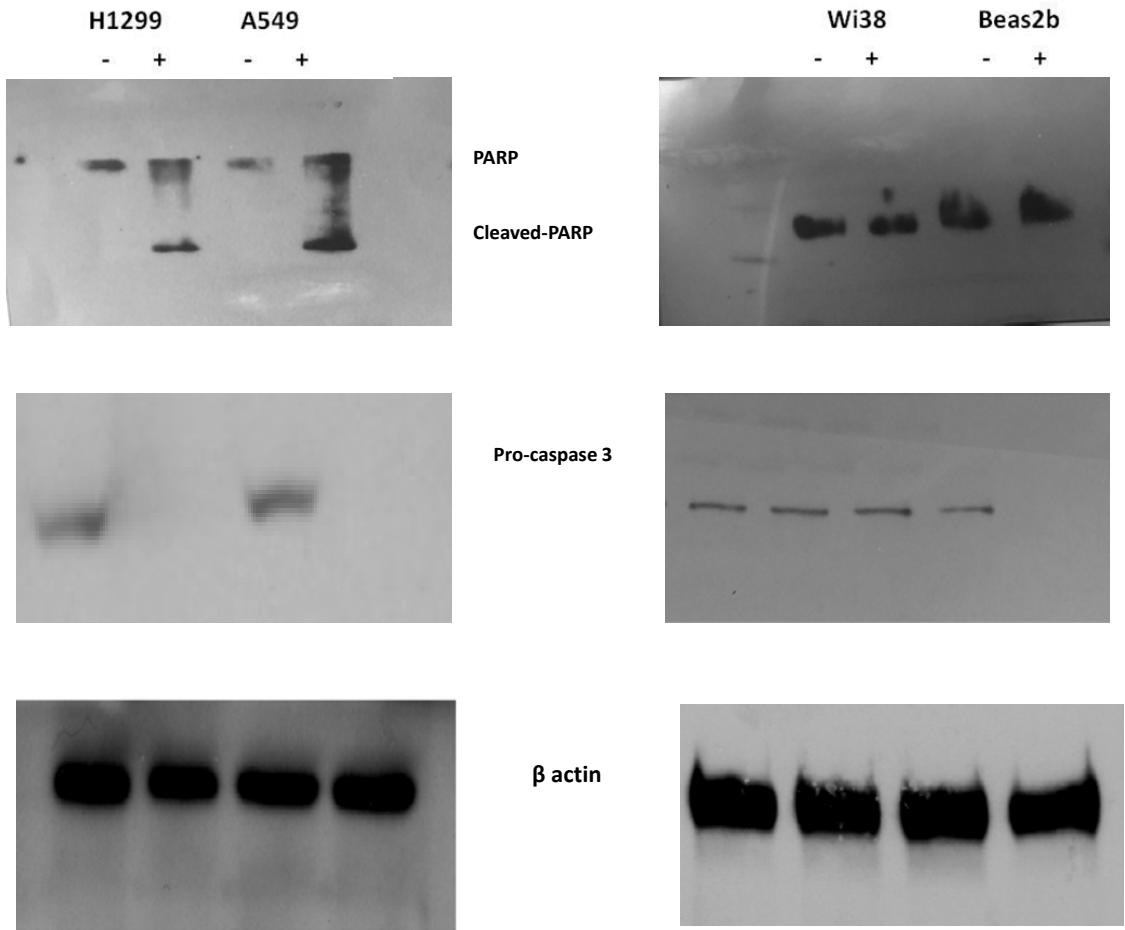


Figure S1. Western Blot analyses of the proteins involved in apoptosis of NSCLC (H1299 and A549) cell lines, normal bronchial epithelial (Beas2b) cells and human fibroblasts (Wi38) . The Western Blot analyses for the indicated proteins were performed on the protein lysates from cells, which were incubated with (+) and without (-) *Urtica dioica* extract (60 µg/mL) for 72 h. The blots have been cut out according to the molecular weight of the proteins to allow multiple hybridizations with specific antibodies including anti-actin antibody that was used to normalise the obtained signals.

After hybridization, the cropped images were acquired and the blots were re-hybridized where applicable.

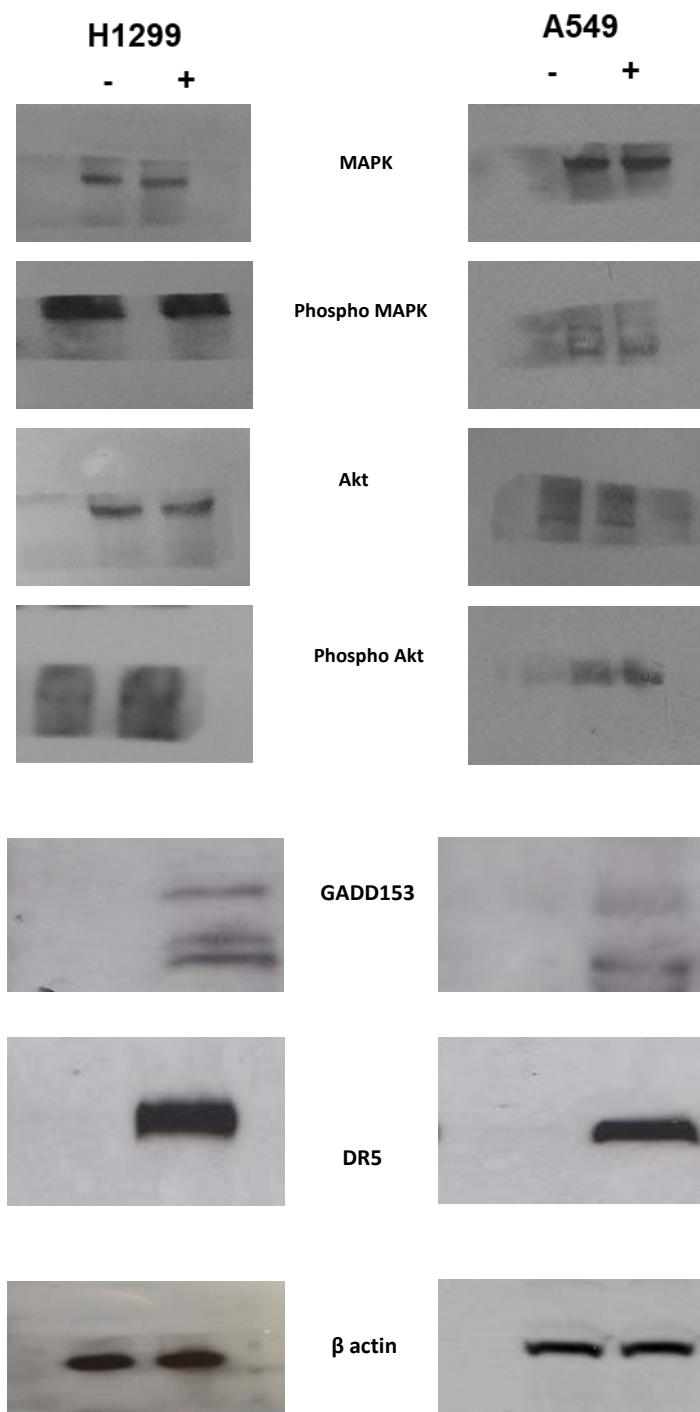
Figure S2.

Figure S2. The expression levels of ER-stress related proteins. The expression levels of ER-stress related proteins were investigated on H1299 and A549 cancer cells following treatment with (+) and without (-) *Urtica dioica* extract (60 µg/mL) for 72h. The blots have been cut out according to the molecular weigh of the proteins to allow multiple hybridizations with specific antibodies, including anti-actin antibody that was used to normalize the obtained signals. After hybridization, the cropped images were acquired and the blots were re-hybridized where applicable.

Figure S3A. CIGAR experiment of *Urtica dioica* crude extract. R= rutin. OXY=oxylipins

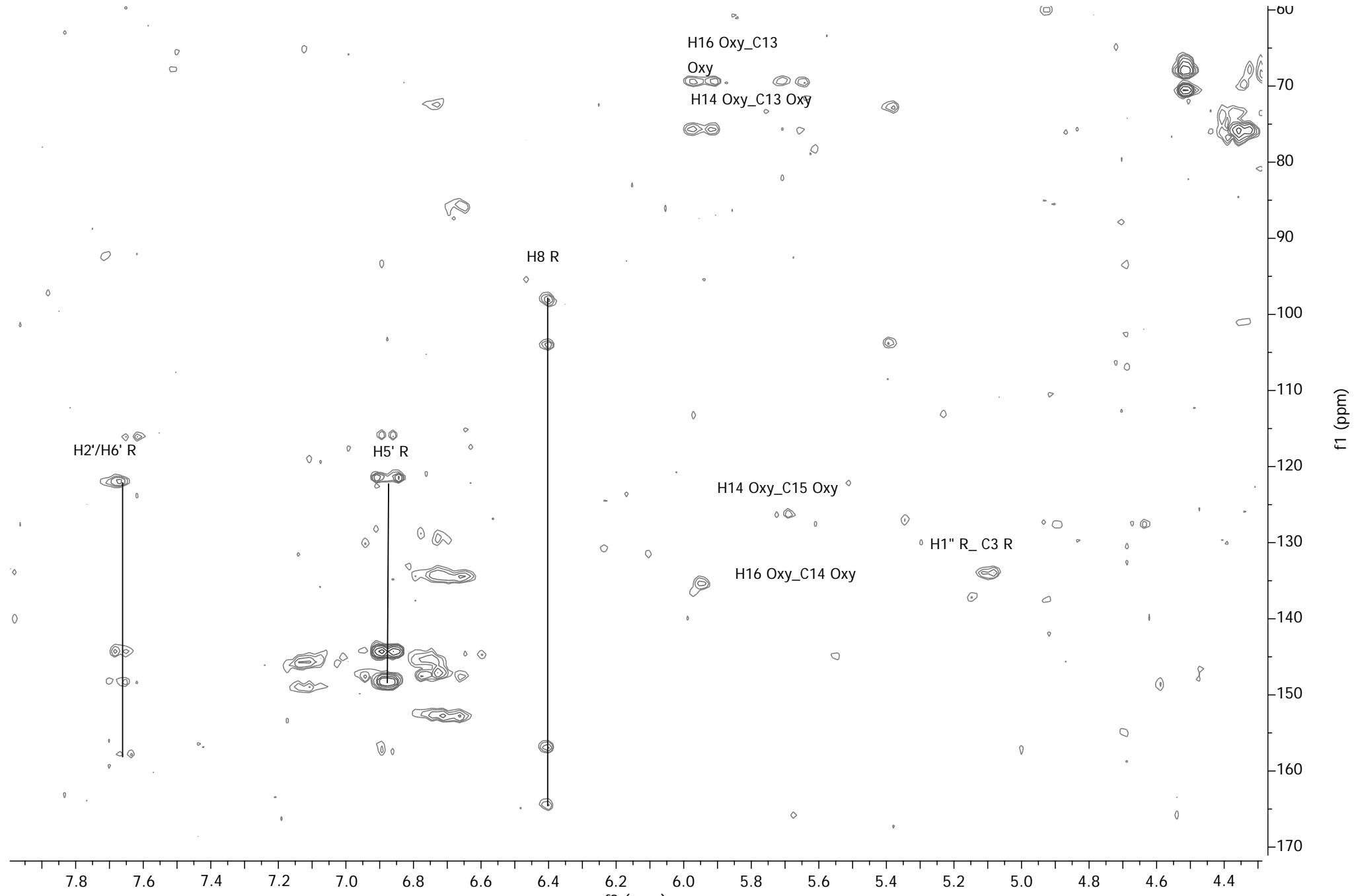


Figure S3B. Aliphatic region of CIGAR experiment of *Urtica dioica* crude extract, R= rutin. OXY=oxylipins

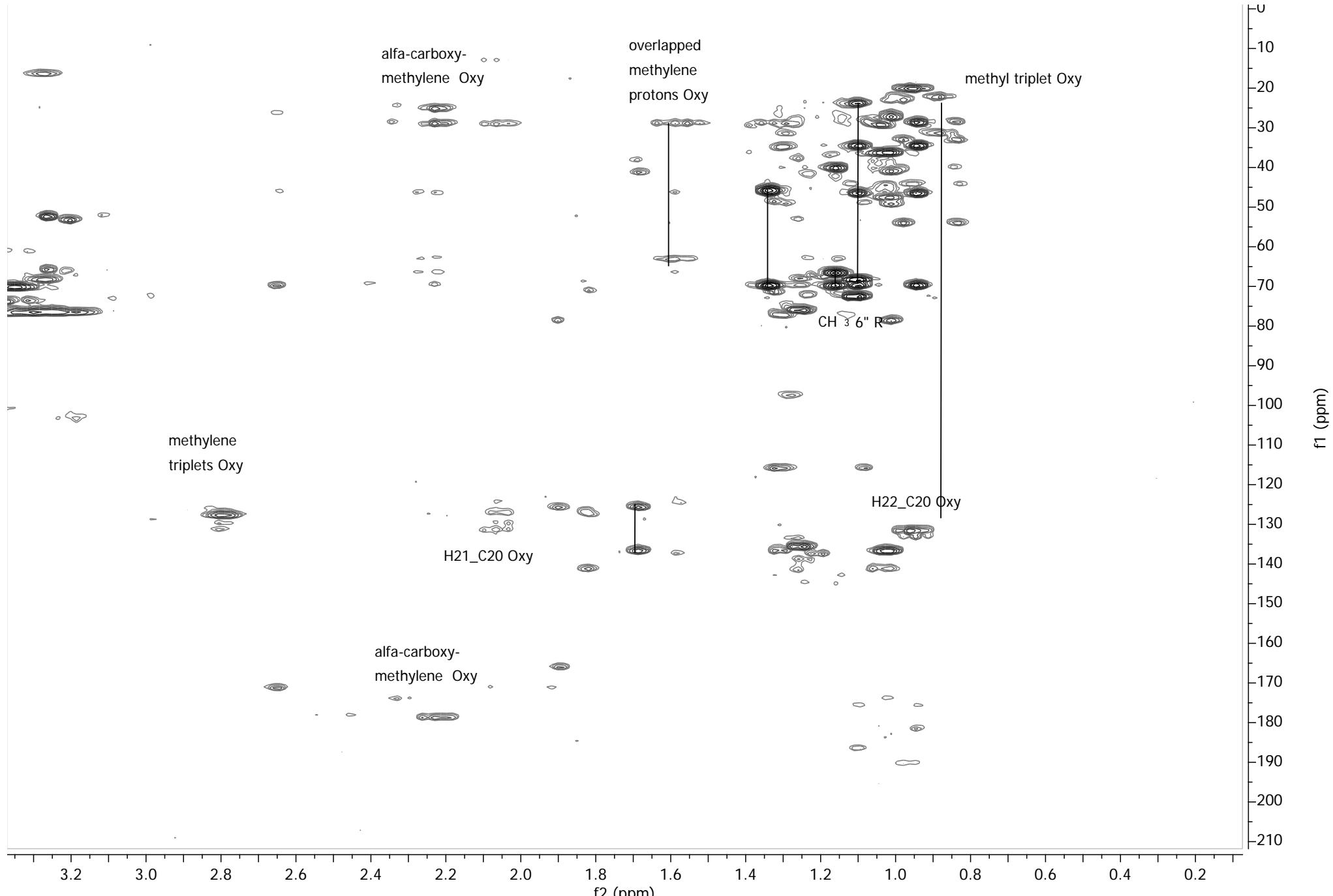


Figure S4. HSQC-TOCSY experiment of *Urtica dioica* crude extract

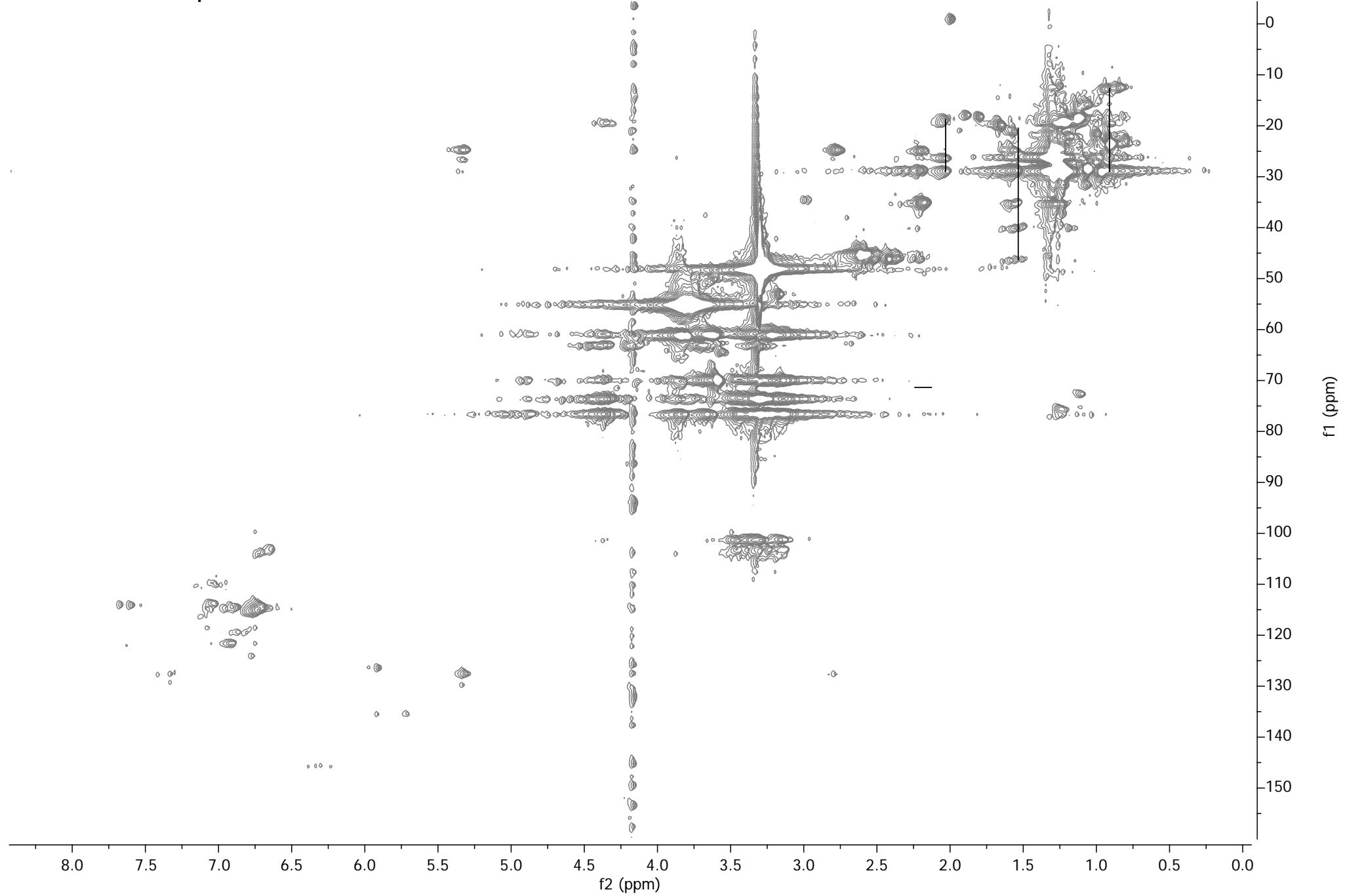


Figure S5. DQCOSY experiment of *Urtica dioica* crude extract. R= rutin. OXY=oxylipins

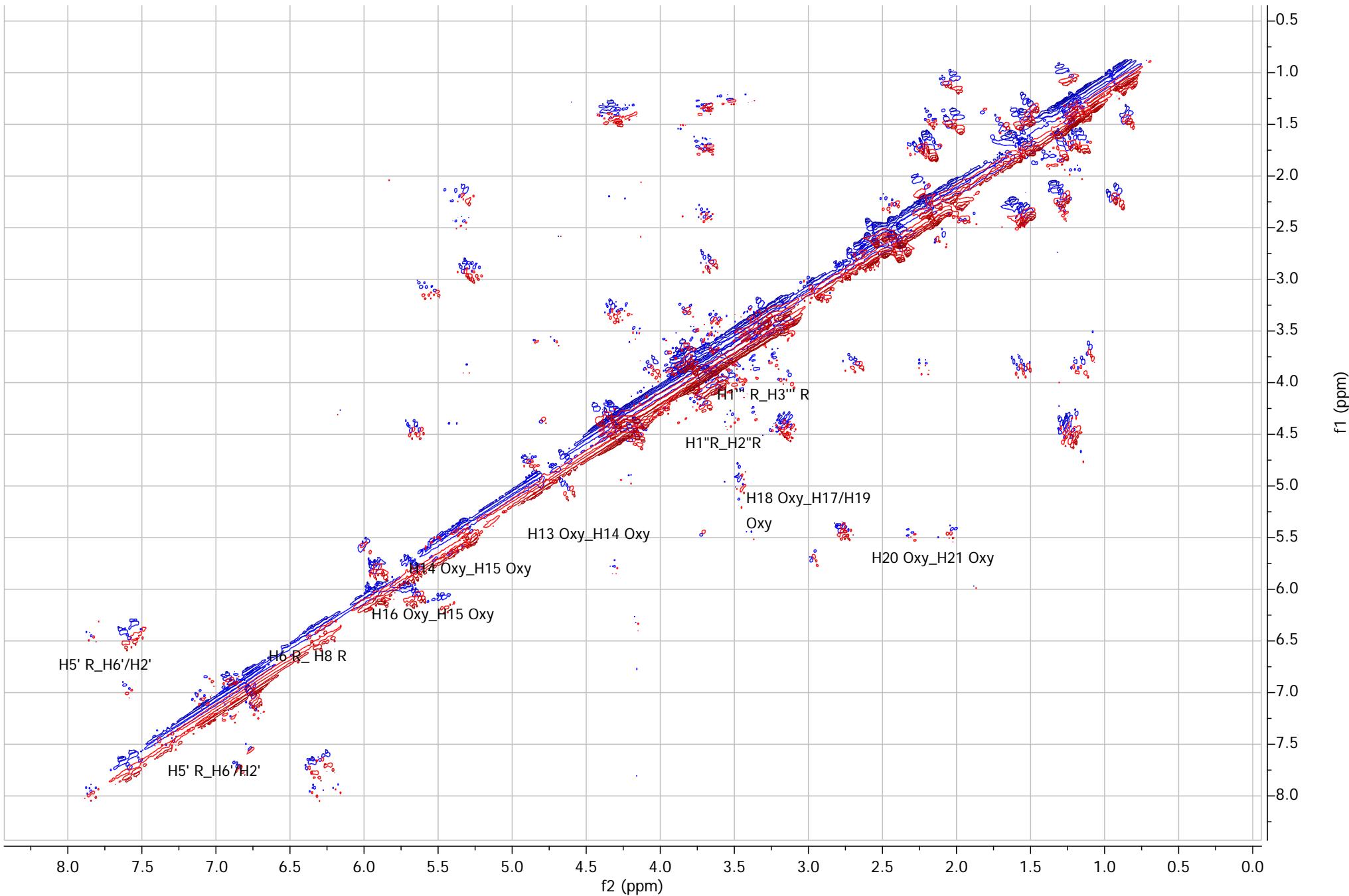


Figure S6. HMBC experiment of rutin

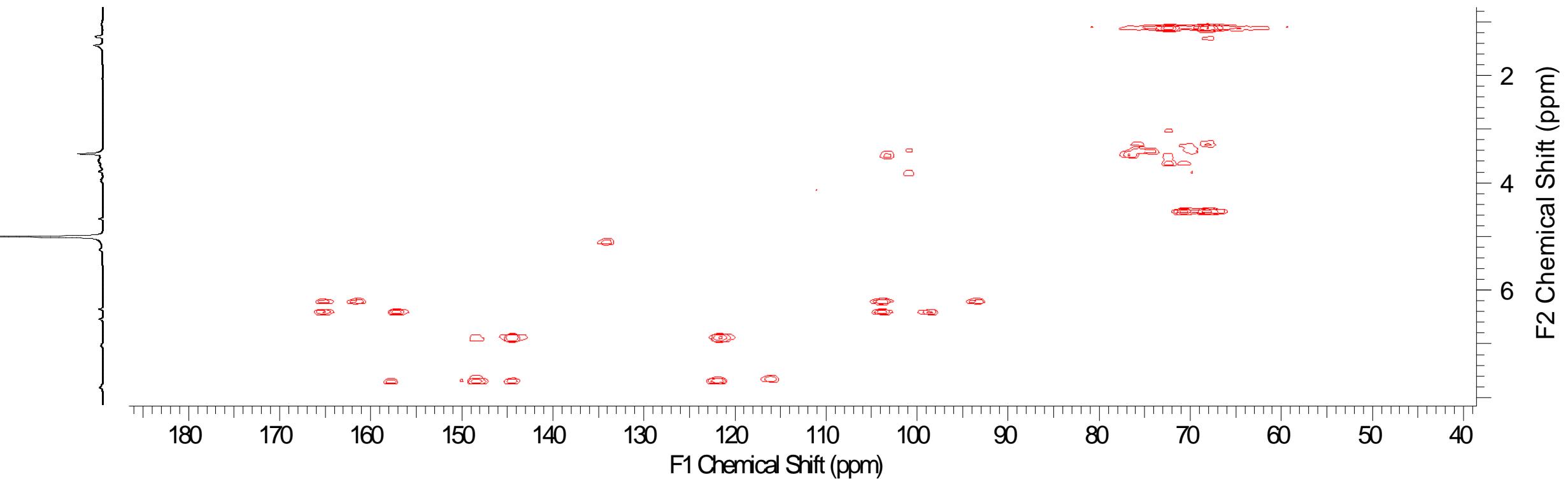


Figure S7. ^1H -NMR spectrum of oxylipins' enriched fraction

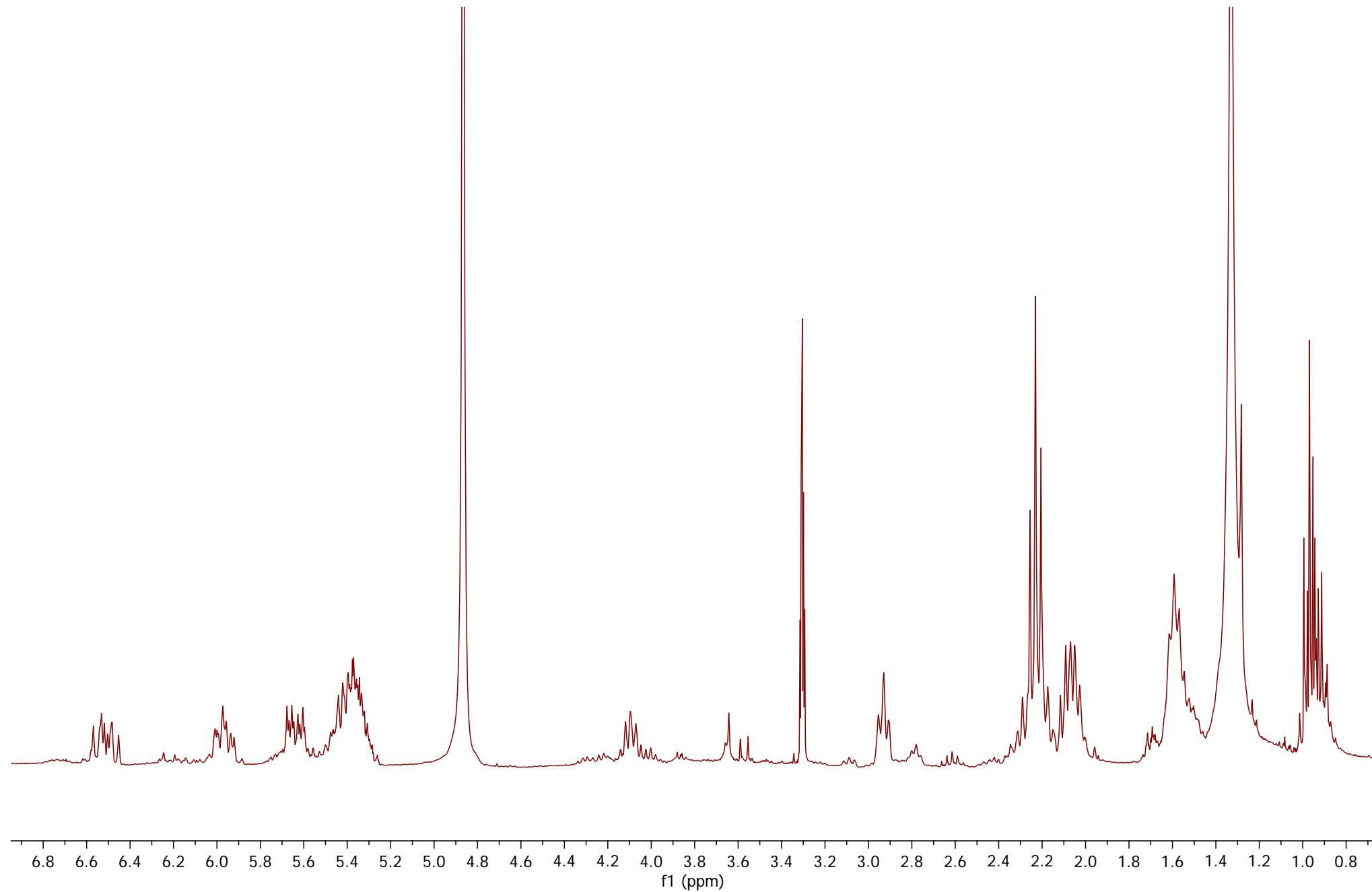


Figure S8. H2BC experiment of oxylipins enriched fraction

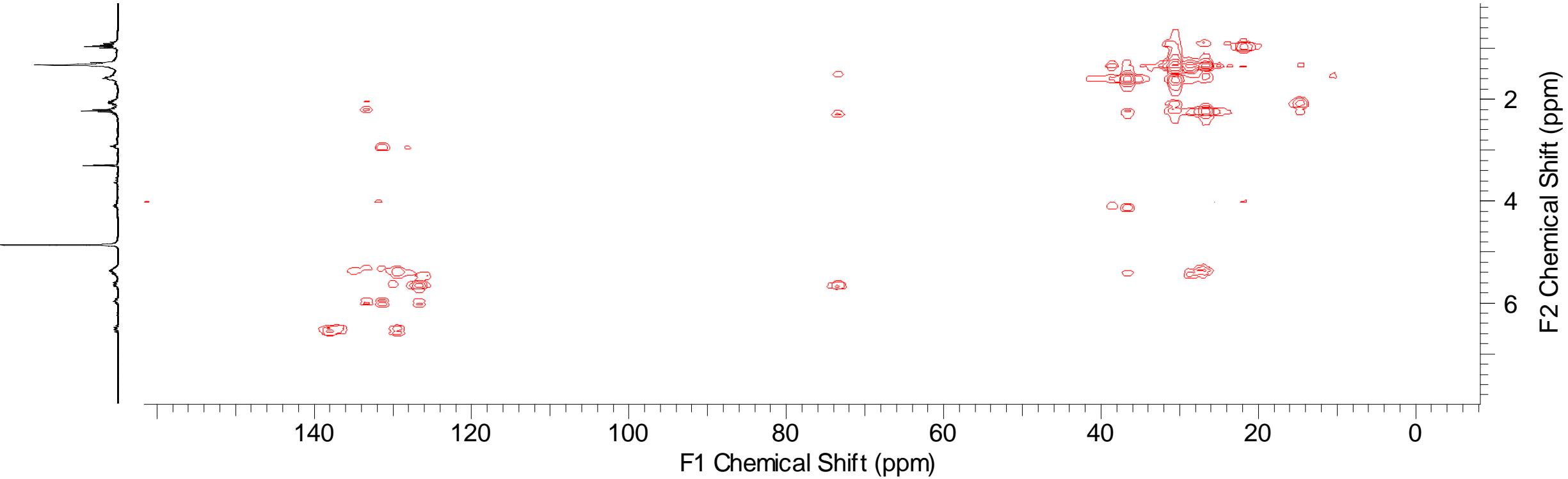
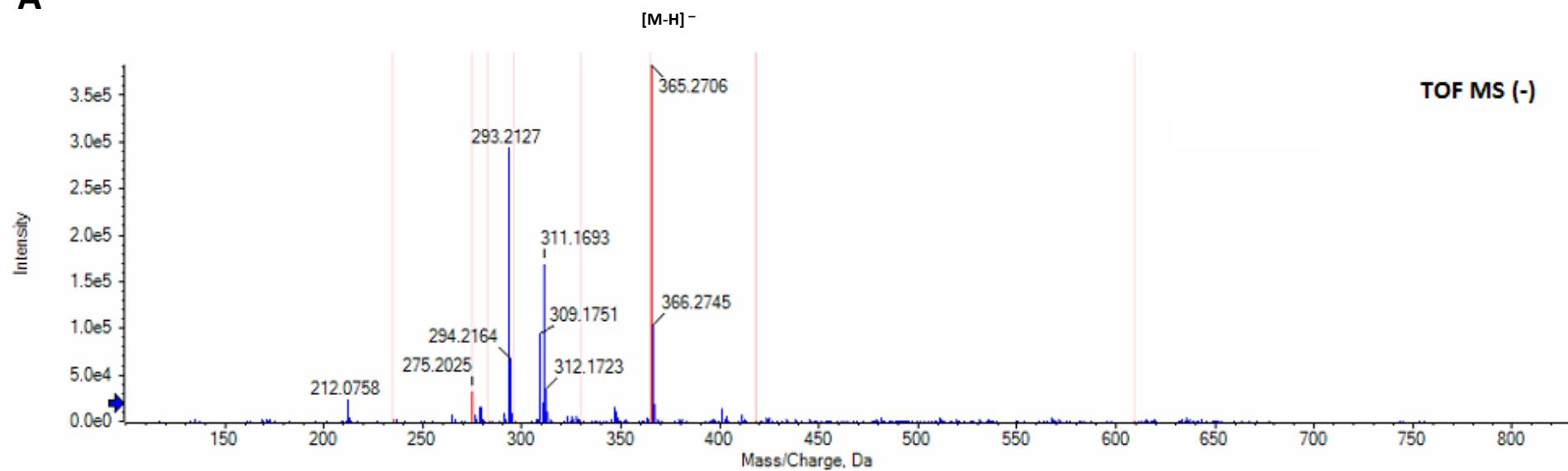


Figure S9. ESI Q-TOF MS SPECTRUM of main oxylipin acquired in negative (A) and positive ion mode (B)

A



B

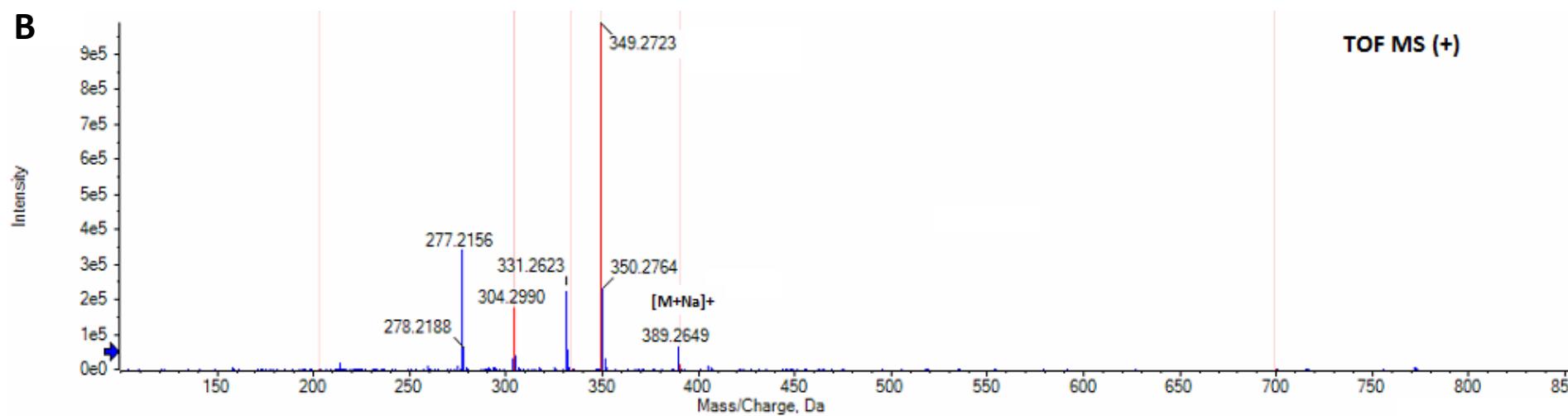


Figure S10 HSQC experiment of the main oxylipin. A aliphatic region. B aromatic region

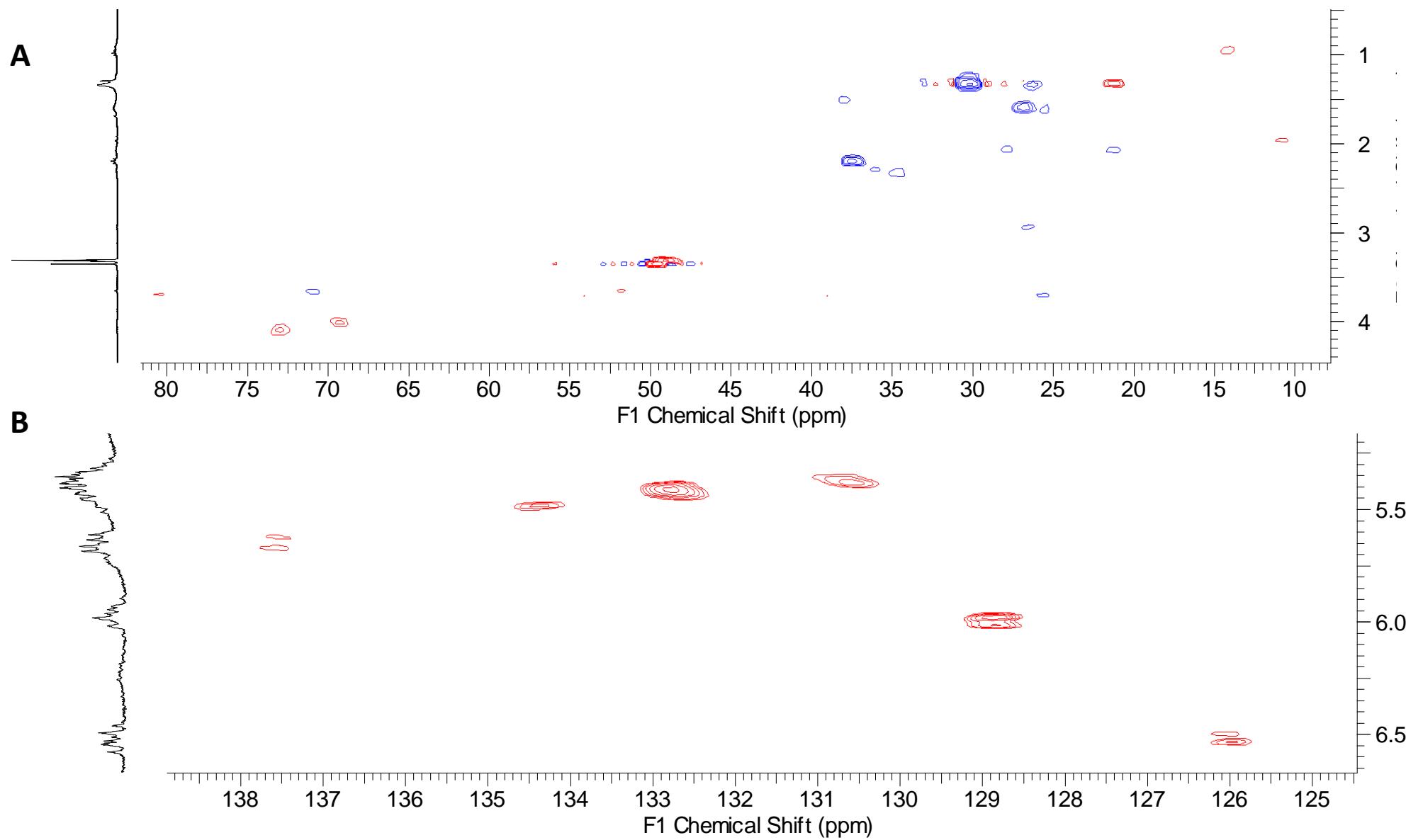


Table S1.NMR data of oxylipin recorded in CD₃OD. δ_H and δ_C are measured in ppm

Position	δ _H ,m (J in Hz)	δ _C , (m)	HMBC (H→C)
1	-	182.4 (C)	
2	2.21 t (7.2)	37.8 (CH ₂)	C1
3	1.61 m	26.8 (CH ₂)	C1
4	1.35 ov	30.6 (CH ₂)	C1
5	1.35 ov	30.5 (CH ₂)	
6	1.35 ov	30.5 (CH ₂)	
7	1.35 ov	30.6 (CH ₂)	
8	1.61 ov	26.1 (CH ₂)	
9	1.51 sv	38.4 (CH ₂)	
10	4.00 ov	69.2(CH)	C9, C8
11	2.22 ov	28.3 (CH ₂)	
12	2.27 ov	36.1 (CH ₂)	
13	4.09 ov	72.9 (CH)	C12, C11, C10
14	5.66 dd (6.6, 15.0 Hz)	137.8 (CH)	C13
15	6.53 dd (15.0 10.8 Hz)	126.1 (CH)	C13, C16, C17
16	5.99 dt (10.8 11.1 Hz)	128.8 (CH)	C14, C15
17	5.35 ov	131.5 (CH)	
18	2.95 t (7.2)	26.6 (CH ₂)	C17, C19, C20
19	5.38 ov	130.8 (CH)	
20	5.43 ov	132.7 (CH)	
21	2.07 m	21.4 (CH ₂)	
22	0.99 t (7.5)	14.2 (CH ₃)	C20, C21

dd=doublet of doublet, m=multiplet, ov=overlapped, t=triplet.