**Analytical and Bioanalytical Chemistry**
**Electronic Supplementary Material**

# Quantitative profiling of oxylipins through comprehensive LC-MS/MS analysis: application in cardiac surgery

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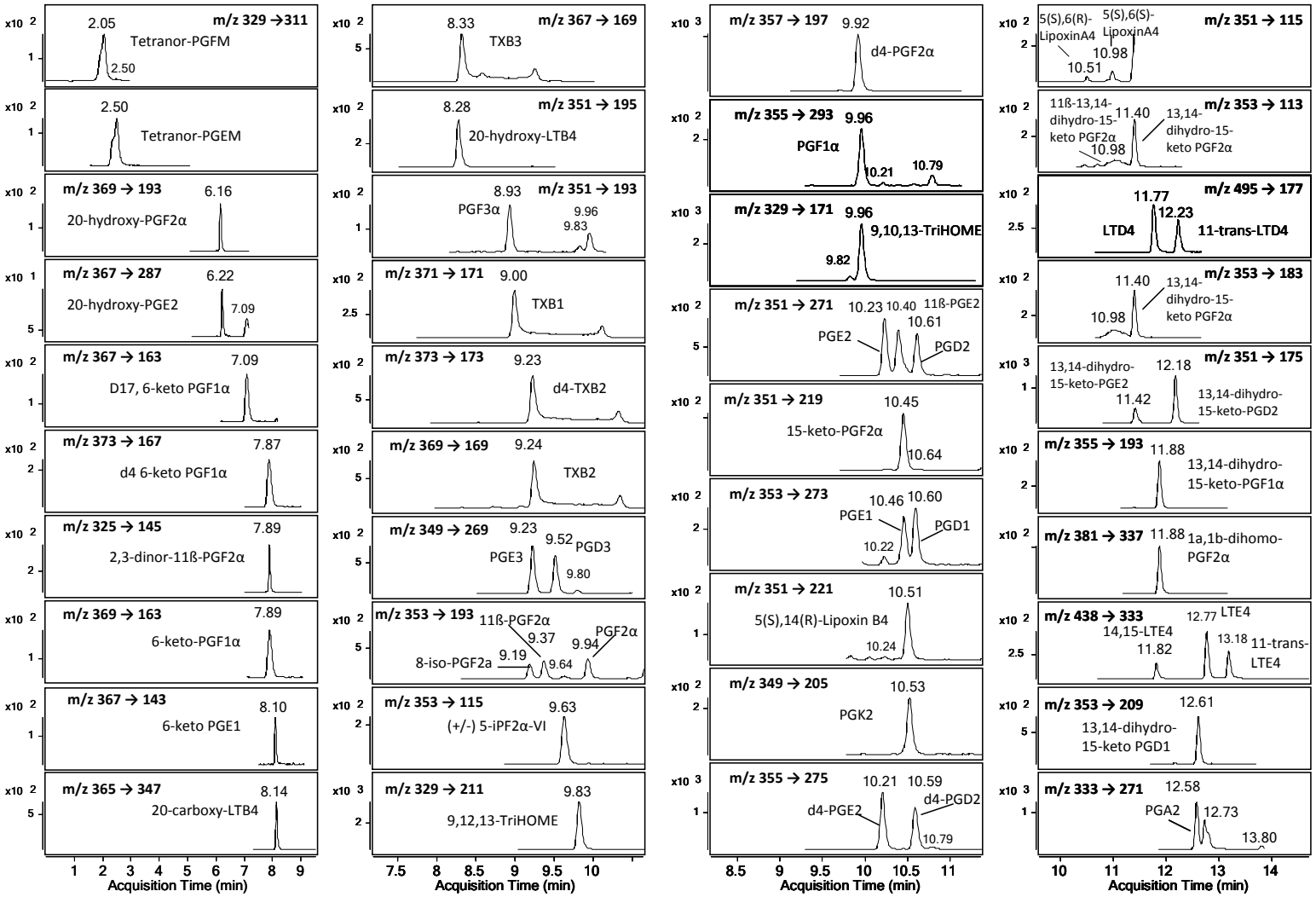


**Fig. S-1a** LC-MS/MS extracted chromatograms of oxylipin library in the range from 2 to 13. 8 min

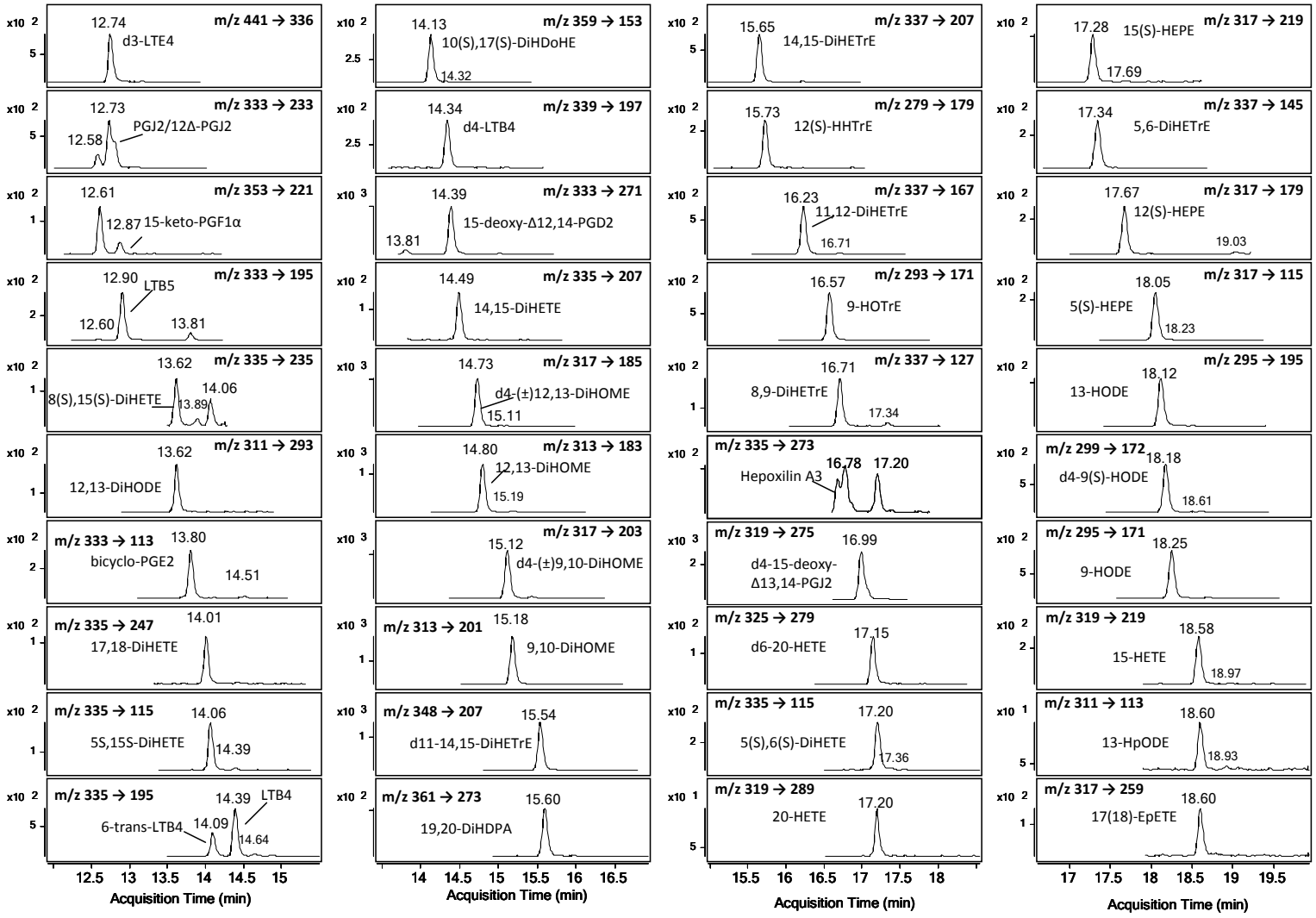


**Fig. S-1b** LC-MS/MS extracted chromatograms of oxylipin library in the range from 10.51 to 18.6 min

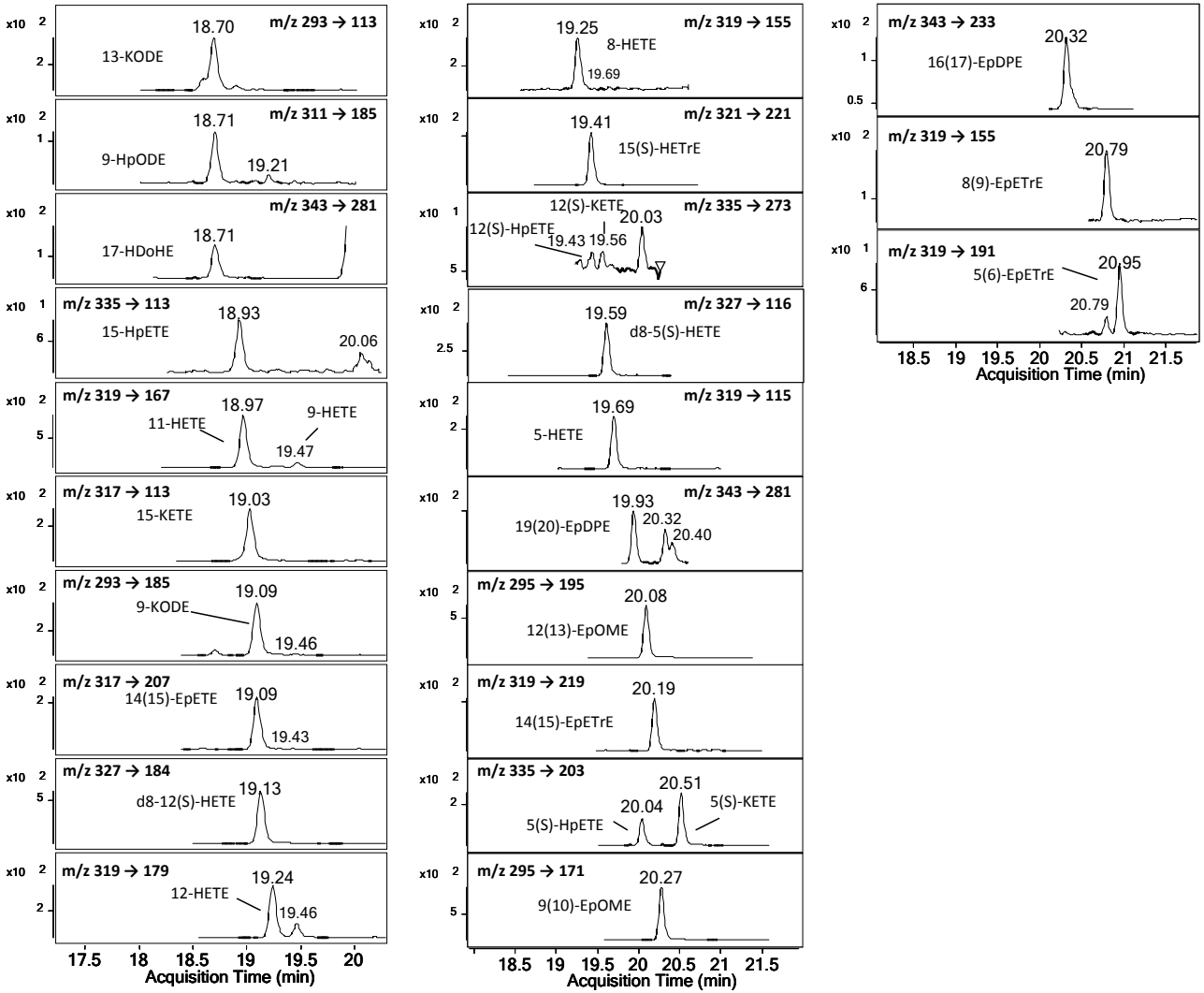


**Fig. S-1c** LC-MS/MS extracted chromatograms of oxylipin library in the range from 15.65 to 20.95 min

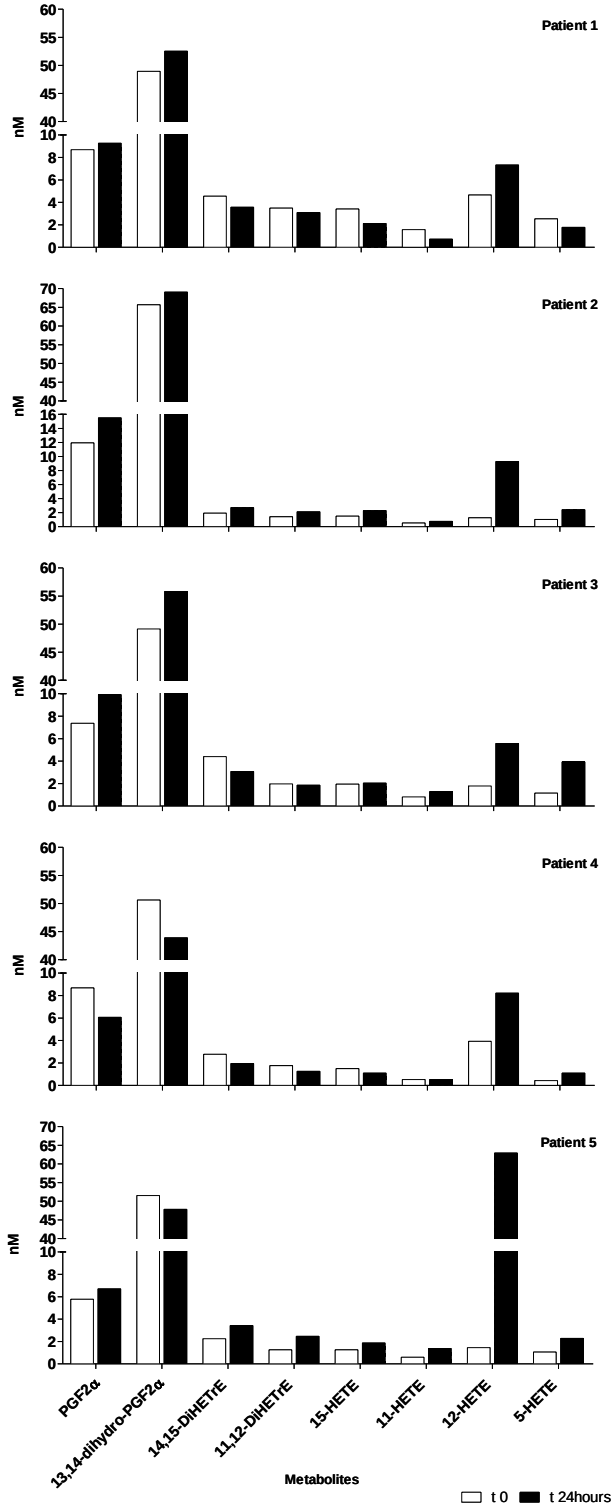


**Fig. S-2** Oxylipins generated in the arachidonic acid pathway with their abundance in the patients (n=5, presented as Patient 1 to Patient 5); the baseline level before surgery (white bars) and levels 24h after the surgery (black bars) are shown

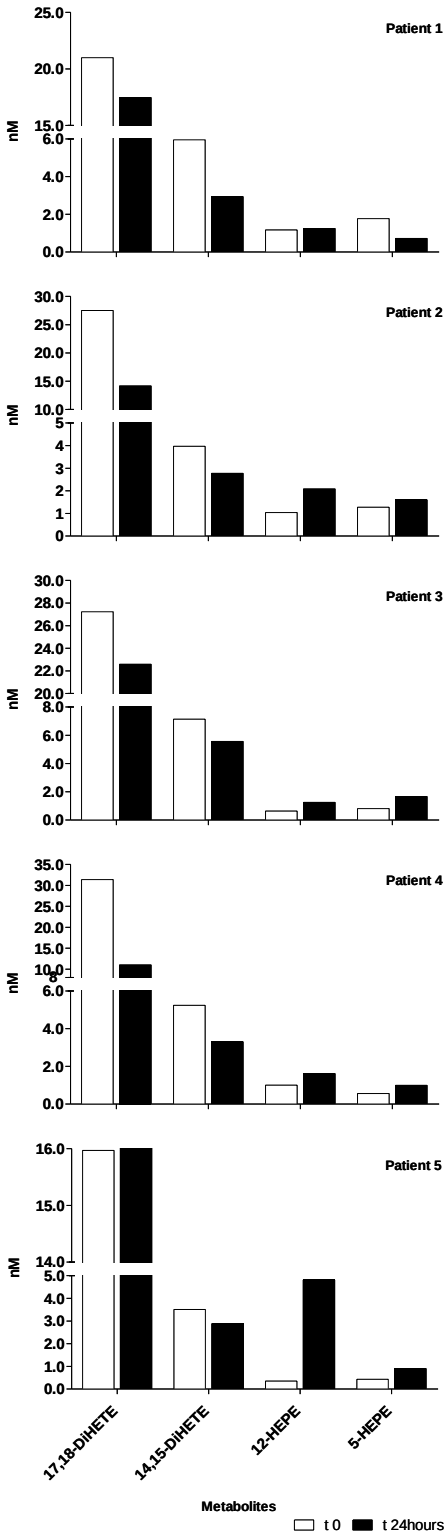


**Fig. S-3** Oxylipins generated in the eicosapentaenoic acid pathway with their abundance in the patients (n=5, presented as Patient 1 to Patient 5); the baseline level before surgery (white bars) and levels 24h after the surgery (black bars) are shown

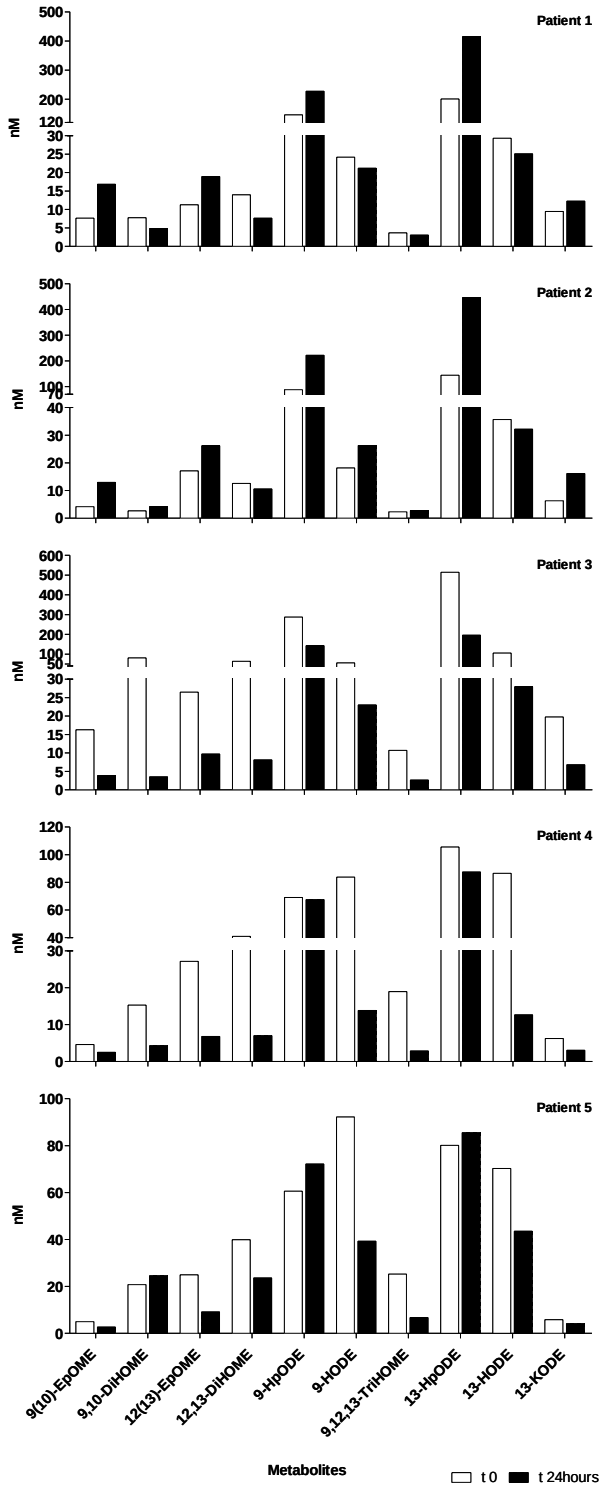


**Fig. S-4** Oxylipins generated in the linoleic acid pathway with their abundance in the patients (n=5, presented as Patient 1 to Patient 5); the baseline level before surgery (white bars) and levels 24h after the surgery (black bars) are shown

**Table S-1** Oxylipin library. The table contains 104 compounds considered in the validation with information of precursor and product ions, as well as retention time and the ISTD for correction.

| Compound Name | Lipid Maps ID | Precursor Ion | Product Ion | Retention Time | Internal Standard |
|---|---|---|---|---|---|
| Tetranor-PGFM | LMFA03010139 | 329.2 | 311.2 | 2.05 | (d4) PGF2α |
| Tetranor-PGEM | LMFA03010032 | 327.1 | 309.2 | 2.50 | (d4) PGE2 |
| 20-hydroxy PGF2α | LMFA03010029 | 369.2 | 193.1 | 6.16 | (d4) PGF2α |
| 20-hydroxy PGE2 | LMFA03010014 | 367.2 | 287.2 | 6.22 | (d4) PGE2 |
| Δ17-6-keto PGF1α | LMFA03010149 | 367.2 | 163.1 | 7.09 | (d4) 6-keto PGF1α |
| **(d4) 6-keto PGF1α** | LMFA03010037 | 373.2 | 167.2 | 7.87 | - |
| 2,3-dinor-11b PGF2α | LMFA03010011 | 325.2 | 145.1 | 7.89 | (d4) PGF2α |
| 6-keto PGF1α | LMFA03010001 | 369.2 | 163.1 | 7.89 | (d4) 6-keto PGF1α |
| 6-keto PGE1 | LMFA03010012 | 367.2 | 143.1 | 8.10 | (d4) PGE2 |
| 20-carboxy LTB4 | LMFA03020016 | 365.2 | 347.2 | 8.14 | (d4) LTB4 |
| 20-hydroxy LTB4 | LMFA03020018 | 351.2 | 195.1 | 8.28 | (d4) LTB4 |
| TXB3 | LMFA03030006 | 367.2 | 169.1 | 8.33 | (d4) TXB2 |
| PGF3α | LMFA03010138 | 351.2 | 193.2 | 8.93 | (d4) PGF2α |
| TXB1 | LMFA03030008 | 371.2 | 171.1 | 9.04 | (d4) TXB2 |
| **(d4) TXB2** | LMFA03030010 | 373.2 | 173.1 | 9.23 | - |
| 8-iso PGF2α | LMFA03110001 | 353.2 | 193.2 | 9.23 | (d4) PGF2α |
| PGE3 | LMFA03010135 | 349.2 | 269.2 | 9.24 | (d4) PGE2 |
| TXB2 | LMFA03030002 | 369.2 | 169.1 | 9.24 | (d4) TXB2 |
| 11β-PGF2α | LMFA03010036 | 353.2 | 193.2 | 9.37 | (d4) PGF2α |
| PGD3 | LMFA03010142 | 349.2 | 269.2 | 9.52 | (d4) PGD2 |
| (+/-) 5-iPF2α-VI | LMFA03110011 | 353.2 | 115.1 | 9.63 | (d4) PGF2α |
| 9,12,13-TriHOME | LMFA02000014 | 329.2 | 211.2 | 9.83 | (d4) 9(S)-HODE |
| **(d4) PGF2α** | LMFA03010006 | 357.3 | 197.2 | 9.92 | - |
| PGF2α | LMFA03010002 | 353.2 | 193.2 | 9.94 | (d4) PGF2α |
| 9,10,13-TriHOME | - | 329.2 | 171.1 | 9.96 | (d4) 9(S)-HODE |
| PGF1α | LMFA03010137 | 355.2 | 293.2 | 9.96 | (d4) PGF2α |
| **(d4) PGE2** | LMFA03010007 | 355.2 | 275.2 | 10.21 | - |
| PGE2 | LMFA03010003 | 351.2 | 271.2 | 10.23 | (d4) PGE2 |
| 11β-PGE2 | LMFA03010060 | 351.2 | 271.2 | 10.40 | (d4) PGE2 |
| 15-keto PGF2α | LMFA03010026 | 351.2 | 219.1 | 10.45 | (d4) PGF2α |
| PGE1 | LMFA03010134 | 353.2 | 273.2 | 10.46 | (d4) PGE2 |
| 5(S),14(R)-Lipoxin B4 | LMFA03040002 | 351.2 | 221.2 | 10.51 | (d4) LTB4 |
| 5(S),6(R)-Lipoxin A4 | LMFA03040001 | 351.2 | 115.1 | 10.51 | (d4) LTB4 |
| PGK2 | LMFA03010023 | 349.2 | 205.1 | 10.53 | (d4) PGE2 |
| **(d4) PGD2** | LMFA03010008 | 355.2 | 275.2 | 10.59 | - |
| PGD1 | LMFA03010049 | 353.2 | 273.2 | 10.60 | (d4) PGD2 |
| PGD2 | LMFA03010004 | 351.2 | 271.2 | 10.61 | (d4) PGD2 |
| 15-keto PGF1α | LMFA03010150 | 353.2 | 221.1 | 10.70 | (d4) 6-keto PGF1α |
| 13,14-dihydro PGF2α | LMFA03010079 | 355.2 | 275.2 | 10.79 | (d4) PGF2α |
| 11β-13,14-dihydro-15-keto PGF2α | LMFA03010203 | 353.2 | 113.2 | 10.98 | (d4) PGF2α |
| 5(S),6(S)-Lipoxin A4 | LMFA03040003 | 351.2 | 115.1 | 10.98 | (d4) LTB4 |
| 13,14-dihydro-15-keto PGF2α | LMFA03010027 | 353.2 | 183.1 | 11.40 | (d4) PGF2α |
| 13,14-dihydro-15-keto PGE2 | LMFA03010031 | 351.2 | 175.2 | 11.42 | (d4) PGE2 |
| 14,15-LTC4 | LMFA03020031 | 624.3 | 272.1 | 11.48 | (d3) LTE4 |

## Continue Table S-1

| Compound Name | Lipid Maps ID | Precursor Ion | Product Ion | Retention Time | Internal Standard |
|---|---|---|---|---|---|
| LTD4 | LMFA03020006 | 495.2 | 177.1 | 11.77 | (d3) LTE4 |
| 14,15-LTE4 | LMFA03020033 | 438.2 | 333.2 | 11.82 | (d3) LTE4 |
| 13,14-dihydro-15-keto PGF1α | - | 355.2 | 193.2 | 11.88 | (d4) PGF2α |
| 1α,1b-dihomo PGF2α | LMFA03010157 | 381.3 | 337.2 | 11.88 | (d4) PGF2α |
| 13,14-dihydro-15-keto PGD2 | LMFA03010022 | 351.2 | 175.2 | 12.18 | (d4) PGD2 |
| 11-trans LTD4 | LMFA03020021 | 495.2 | 177.1 | 12.23 | (d3) LTE4 |
| PGA2 | LMFA03010035 | 333.2 | 271.2 | 12.58 | (d4) PGE2 |
| 13,14-dihydro-15-keto PGD1 | - | 353.2 | 209.1 | 12.61 | (d4) PGD2 |
| PGJ2 | LMFA03010019 | 333.2 | 233.1 | 12.73 | (d4) PGD2 |
| Δ12-PGJ2 | LMFA03010020 | 333.2 | 233.1 | 12.73 | (d4) 15-deoxy-Δ12,14-PGJ2 |
| **(d3) LTE4** | - | 441.2 | 336.2 | 12.74 | - |
| LTE4 | LMFA03020002 | 438.2 | 333.2 | 12.77 | (d3) LTE4 |
| LTB5 | LMFA03020010 | 333.2 | 195.1 | 12.90 | (d4) LTB4 |
| LTC4 | LMFA03020003 | 624.3 | 272.1 | 13.04 | (d3) LTE4 |
| 11-trans LTE4 | LMFA03020022 | 438.2 | 333.2 | 13.18 | (d3) LTE4 |
| 11-trans LTC4 | LMFA03020020 | 624.3 | 272.1 | 13.57 | (d3) LTE4 |
| 12,13-DiHODE | LMFA02000046 | 311.2 | 293 | 13.62 | (d4) 9(S)-HODE |
| 8(S),15(S)-DiHETE | LMFA03060050 | 335.2 | 235.2 | 13.62 | (d4) LTB4 |
| bicyclo-PGE2 | LMFA03010034 | 333.2 | 113.2 | 13.80 | (d4) PGE2 |
| 17,18-DiHETE | LMFA03060078 | 335.2 | 247.2 | 14.01 | (d11) 14,15-DiHETrE |
| 5(S),15(S)-DiHETE | LMFA03060049 | 335.2 | 115.2 | 14.06 | (d4) LTB4 |
| 6-trans-LTB4 | LMFA03020013 | 335.2 | 195.1 | 14.09 | (d4) LTB4 |
| 10(S),17(S)-DiHDoHE | LMFA04000047 | 359.2 | 153.2 | 14.13 | (d8) 12(S)-HETE |
| **(d4) LTB4** | LMFA03020030 | 339.2 | 197.1 | 14.34 | - |
| 15-deoxy-Δ12,14-PGD2 | LMFA03010051 | 333.2 | 271.2 | 14.39 | (d4) PGD2 |
| LTB4 | LMFA03020001 | 335.2 | 195.1 | 14.39 | (d4) LTB4 |
| 14,15-DiHETE | LMFA03060077 | 335.2 | 207.1 | 14.49 | (d11) 14,15-DiHETrE |
| **(d4)(±)12,13-DiHOME** | LMFA01050357 | 317.3 | 185.2 | 14.73 | - |
| 12,13-DiHOME | LMFA01050351 | 313.2 | 183.2 | 14.80 | (d4) 12,13-DiHOME |
| **(d4)-(±)9,10-DiHOME** | LMFA01050358 | 317.3 | 203.2 | 15.12 | - |
| 9,10-DiHOME | LMFA01050350 | 313.2 | 201.1 | 15.18 | (d4) 9,10-DiHOME |
| **(d11) 14,15-DiHETrE** | - | 348.3 | 207.1 | 15.54 | - |
| 19,20-DiHDPA | LMFA04000043 | 361.2 | 273.3 | 15.60 | (d8) 12(S)-HETE |
| 14,15-DiHETrE | LMFA03050010 | 337.2 | 207.2 | 15.65 | (d11) 14,15-DiHETrE |
| 12S-HHTrE | LMFA03050002 | 279.2 | 179.2 | 15.73 | (d8) 12(S)-HETE |
| 11,12-DiHETrE | LMFA03050008 | 337.2 | 167.2 | 16.23 | (d11) 14,15-DiHETrE |
| 9-HOTrE | LMFA02000024 | 293.2 | 171.1 | 16.57 | (d4) 9(S)-HODE |
| 8,9-DiHETrE | LMFA03050006 | 337.2 | 127 | 16.71 | (d11) 14,15-DiHETrE |
| Hepoxilin A3 | LMFA03090005 | 335.2 | 273.2 | 16.78 | (d8) 12(S)-HETE |
| **(d4) 15-deoxy-Δ12,14-PGJ2** | LMFA03010177 | 319.2 | 275.3 | 16.99 | - |
| **(d6) 20-HETE** | LMFA03060082 | 325.3 | 279.2 | 17.15 | - |
| 20-HETE | LMFA03060009 | 319.2 | 289.2 | 17.20 | d6-20-HETE |
| 5(S),6(S)-DiHETE | LMFA03060018 | 335.2 | 115.1 | 17.20 | (d4) LTB4 |

## Continue Table S-1

| Compound Name | Lipid Maps ID | Precursor Ion | Product Ion | Retention Time | Internal Standard |
|---|---|---|---|---|---|
| 15(S)-HEPE | LMFA03070009 | 317.2 | 219.2 | 17.28 | (d8) 5(S)-HETE |
| 5,6-DiHETrE | LMFA03050004 | 337.2 | 145.1 | 17.34 | (d11) 14,15-DiHETrE |
| 12(S)-HEPE | LMFA03070008 | 317.2 | 179.1 | 17.67 | (d8) 12(S)-HETE |
| 5(S)-HEPE | LMFA03070010 | 317.2 | 115.1 | 18.05 | (d8) 5(S)-HETE |
| 13-HODE | LMFA01050349 | 295.2 | 195.2 | 18.12 | (d4) 9(S)-HODE |
| **(d4) 9(S)-HODE** | LMFA01050353 | 299.2 | 172.1 | 18.18 | - |
| 9-HODE | LMFA01050278 | 295.2 | 171.1 | 18.25 | (d4) 9(S)-HODE |
| 15-HETE | LMFA03060001 | 319.2 | 219.2 | 18.58 | (d8) 5(S)-HETE |
| 13-HpODE | LMFA02000034 | 311.2 | 113.2 | 18.60 | (d4) 9(S)-HODE |
| 17(18)-EpETE | LMFA03000004 | 317.2 | 259.2 | 18.60 | (d11) 14,15-DiHETrE |
| 13-KODE | LMFA02000016 | 293.2 | 113.1 | 18.70 | (d4) 9(S)-HODE |
| 9-HpODE | LMFA02000012 | 311.2 | 185.2 | 18.71 | (d4) 9(S)-HODE |
| 17-HDoHE | LMFA04000032 | 343.2 | 281.3 | 18.71 | (d8) 12(S)-HETE |
| 15-HpETE | LMFA03060014 | 335.2 | 113.1 | 18.93 | (d8) 5(S)-HETE |
| 11-HETE | LMFA03060028 | 319.2 | 167.1 | 18.97 | (d8) 12(S)-HETE |
| 15-KETE | LMFA03060051 | 317.2 | 113.2 | 19.03 | (d8) 5(S)-HETE |
| 9-KODE | LMFA01060177 | 293.2 | 185.2 | 19.09 | (d4) 9(S)-HODE |
| 14(15)-EpETE | LMFA03000003 | 317.2 | 207.1 | 19.09 | (d11) 14,15-DiHETrE |
| **(d8) 12(S)-HETE** | LMFA03060081 | 327.3 | 184.2 | 19.13 | - |
| 12-HETE | LMFA03060088 | 319.2 | 179.2 | 19.24 | (d8) 12(S)-HETE |
| 8-HETE | LMFA03060006 | 319.2 | 155.1 | 19.25 | (d8) 5(S)-HETE |
| 15(S)-HETrE | LMFA03050007 | 321.2 | 221.2 | 19.41 | (d11) 14,15-DiHETrE |
| 12(S)-HpETE | LMFA03060013 | 335.2 | 273.3 | 19.43 | (d8) 12(S)-HETE |
| 9-HETE | LMFA03060089 | 319.2 | 167.1 | 19.47 | (d8) 12(S)-HETE |
| 12-KETE | LMFA03060019 | 317.2 | 273.3 | 19.56 | (d8) 12(S)-HETE |
| **(d8) 5(S)-HETE** | LMFA03060005 | 327.3 | 116.1 | 19.59 | - |
| 5-HETE | LMFA03060002 | 319.2 | 115.1 | 19.69 | (d8) 5(S)-HETE |
| 19(20)-EpDPE | LMFA04000038 | 343.2 | 281.3 | 19.93 | (d8) 12(S)-HETE |
| 5(S)-HpETE | LMFA03060012 | 335.2 | 203.2 | 20.04 | (d8) 5(S)-HETE |
| 12(13)-EpOME | LMFA02000038 | 295.2 | 195.2 | 20.08 | (d4) 12,13-DiHOME |
| 14(15)-EpETrE | LMFA03080005 | 319.2 | 219.2 | 20.19 | (d11) 14,15-DiHETrE |
| 9(10)-EpOME | LMFA02000037 | 295.2 | 171.2 | 20.27 | (d4) 9,10-DiHOME |
| 16(17)-EpDPE | LMFA04000037 | 343.2 | 233.2 | 20.32 | (d8) 12(S)-HETE |
| 5-KETE | LMFA03060011 | 317.2 | 203.2 | 20.51 | (d8) 5(S)-HETE |
| 11(12)-EpETrE | LMFA03080004 | 319.2 | 167.1 | 20.71 | (d11) 14,15-DiHETrE |
| 8(9)-EpETrE | LMFA03080003 | 319.2 | 155.1 | 20.79 | (d11) 14,15-DiHETrE |
| 5(6)-EpETrE | LMFA03080002 | 319.2 | 191.2 | 20.95 | (d11) 14,15-DiHETrE |

**Table S-2** Concentration levels of 104 oxylipins for calibration lines. The values display the amount on column.

| | conc [nM] | | | | | | | | |
|---|---|---|---|---|---|---|---|---|---|
| **compound name** | **C0** | **C1** | **C2** | **C3** | **C4** | **C5** | **C6** | **C7** | **C8** |
| Tetranor-PGFM | 0 | 3 | 6 | 13 | 25 | 50 | 101 | 201 | 403 |
| Tetranor-PGEM | 0 | 3 | 6 | 13 | 25 | 50 | 100 | 201 | 401 |
| 20-hydroxy PGF2α | 0 | 2 | 4 | 9 | 18 | 35 | 71 | 142 | 284 |
| 20-hydroxy PGE2 | 0 | 3 | 6 | 12 | 25 | 50 | 100 | 200 | 399 |
| Δ17-6-keto PGF1α | 0 | 2 | 4 | 8 | 17 | 33 | 67 | 133 | 266 |
| 2,3-dinor-11b PGF2α | 0 | 3 | 6 | 13 | 25 | 51 | 102 | 204 | 408 |
| 6-keto PGF1α | 0 | 2 | 4 | 9 | 18 | 35 | 71 | 142 | 284 |
| 20-carboxy LTB4 | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 138 | 275 |
| 6-keto PGE1 | 0 | 3 | 6 | 12 | 24 | 48 | 95 | 190 | 380 |
| 20-hydroxy LTB4 | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 278 |
| TXB3 | 0 | 3 | 6 | 12 | 25 | 50 | 100 | 200 | 399 |
| PGF3α | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 278 |
| TXB1 | 0 | 3 | 6 | 12 | 24 | 47 | 94 | 188 | 376 |
| 8-iso PGF2α | 0 | 2 | 4 | 9 | 17 | 35 | 69 | 138 | 277 |
| PGE3 | 0 | 3 | 6 | 12 | 25 | 50 | 100 | 200 | 400 |
| TXB2 | 0 | 3 | 6 | 12 | 24 | 47 | 95 | 189 | 378 |
| 11β-PGF2α | 0 | 3 | 6 | 12 | 25 | 49 | 99 | 198 | 395 |
| PGD3 | 0 | 3 | 6 | 12 | 25 | 50 | 100 | 200 | 400 |
| 5-iPF2α-VI | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 140 | 281 |
| 9,12,13-TriHOME | 0 | 3 | 6 | 13 | 25 | 50 | 100 | 201 | 401 |
| PGF1α | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 138 | 275 |
| 9,10,13-TriHOME | 0 | 3 | 6 | 13 | 25 | 50 | 101 | 201 | 402 |
| PGF2α | 0 | 2 | 4 | 9 | 17 | 35 | 69 | 138 | 277 |
| PGE2 | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 278 |
| 11β-PGE2 | 0 | 3 | 6 | 12 | 25 | 50 | 99 | 199 | 397 |
| 15-keto PGF2α | 0 | 3 | 6 | 12 | 25 | 50 | 99 | 199 | 397 |
| PGE1 | 0 | 2 | 4 | 9 | 17 | 35 | 69 | 138 | 277 |
| PGK2 | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 5(S),14(R)-Lipoxin B4 | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 278 |
| PGD1 | 0 | 3 | 6 | 12 | 25 | 49 | 99 | 198 | 395 |
| PGD2 | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 278 |
| 15-keto PGF1α | 0 | 2 | 4 | 9 | 17 | 35 | 69 | 138 | 277 |
| 13,14-dihydro PGF2α | 0 | 3 | 6 | 12 | 25 | 49 | 98 | 196 | 393 |
| 11β-13,14-dihydro-15-keto PGF2α | 0 | 2 | 4 | 9 | 17 | 35 | 69 | 138 | 277 |
| LTD4 | 0 | 2 | 4 | 8 | 17 | 34 | 68 | 135 | 271 |
| 13,14-dihydro-15-keto PGF2α | 0 | 3 | 6 | 12 | 25 | 49 | 99 | 198 | 395 |
| 14,15-LTC4 | 0 | 3 | 6 | 13 | 25 | 50 | 101 | 202 | 403 |
| 5(S),6(R)-Lipoxin A4 | 0 | 5 | 9 | 18 | 36 | 73 | 146 | 291 | 583 |
| 13,14-dihydro-15-keto PGE2 | 0 | 3 | 6 | 12 | 25 | 50 | 99 | 199 | 397 |
| 5(S),6(S)-Lipoxin A4 | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 278 |
| 14,15-LTE4 | 0 | 3 | 6 | 13 | 25 | 50 | 101 | 201 | 403 |
| 11-trans LTD4 | 0 | 2 | 4 | 8 | 17 | 34 | 68 | 135 | 271 |
| 13,14-dihydro-15-keto PGF1α | 0 | 3 | 6 | 12 | 24 | 48 | 95 | 190 | 380 |
| 1a,1b-dihomo PGF2α | 0 | 3 | 6 | 11 | 23 | 46 | 92 | 183 | 366 |
| 13,14-dihydro-15-keto PGD2 | 0 | 3 | 6 | 12 | 25 | 50 | 99 | 199 | 397 |
| PGA2 | 0 | 3 | 7 | 13 | 26 | 52 | 105 | 209 | 419 |
| 13,14-dihydro-15-keto PGD1 | 0 | 2 | 4 | 9 | 17 | 35 | 69 | 138 | 276 |
| PGJ2 | 0 | 3 | 7 | 13 | 26 | 52 | 105 | 209 | 419 |
| LTE4 | 0 | 3 | 6 | 13 | 25 | 50 | 100 | 201 | 402 |
| Δ12-PGJ2 | 0 | 2 | 3 | 7 | 13 | 26 | 52 | 105 | 209 |
| LTB5 | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 140 | 281 |
| LTC4 | 0 | 3 | 6 | 13 | 25 | 50 | 101 | 202 | 403 |
| 11-trans LTE4 | 0 | 3 | 6 | 13 | 25 | 50 | 100 | 201 | 402 |
| 11-trans LTC4 | 0 | 3 | 6 | 13 | 25 | 50 | 101 | 202 | 403 |

## Continue Table S-2

| | conc [nM] | | | | | | | | |
|---|---|---|---|---|---|---|---|---|---|
| **compound name** | **C0** | **C1** | **C2** | **C3** | **C4** | **C5** | **C6** | **C7** | **C8** |
| 12,13-DiHODE | 0 | 2 | 4 | 8 | 17 | 34 | 67 | 135 | 269 |
| 8(S),15(S)-DiHETE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 279 |
| bicyclo-PGE2 | 0 | 3 | 7 | 13 | 26 | 52 | 105 | 209 | 419 |
| 17,18-DiHETE | 0 | 3 | 6 | 12 | 25 | 50 | 100 | 200 | 400 |
| 5(S),15(S)-DiHETE | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 137 | 275 |
| 6-trans-LTB4 | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 137 | 275 |
| 10(S),17(S)-DiHDoHE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 15-deoxy-Δ12,14-PGD2 | 0 | 3 | 5 | 10 | 20 | 41 | 81 | 163 | 326 |
| LTB4 | 0 | 4 | 9 | 17 | 35 | 70 | 139 | 279 | 558 |
| 14,15-DiHETE | 0 | 3 | 6 | 12 | 25 | 50 | 100 | 200 | 400 |
| 12,13-DiHOME | 0 | 2 | 4 | 9 | 17 | 35 | 69 | 138 | 276 |
| 9,10-DiHOME | 0 | 2 | 4 | 9 | 17 | 35 | 69 | 138 | 276 |
| 19,20-DiHDPA | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 278 |
| 14,15-DiHETrE | 0 | 2 | 5 | 9 | 18 | 37 | 74 | 148 | 295 |
| 12S-HHTrE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 11,12-DiHETrE | 0 | 2 | 5 | 9 | 19 | 37 | 74 | 149 | 297 |
| 9-HOTrE | 0 | 2 | 4 | 9 | 18 | 36 | 71 | 143 | 285 |
| 8,9-DiHETrE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| Hepoxilin A3 | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 139 | 279 |
| 20-HETE | 0 | 2 | 4 | 8 | 17 | 33 | 66 | 132 | 265 |
| 5(S),6(S)-DiHETE | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 137 | 275 |
| 15(S)-HEPE | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 141 | 282 |
| 5,6-DiHETrE | 0 | 2 | 5 | 9 | 19 | 37 | 74 | 149 | 297 |
| 12(S)-HEPE | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 141 | 282 |
| 5(S)-HEPE | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 141 | 282 |
| 13-HODE | 0 | 2 | 4 | 9 | 18 | 35 | 71 | 142 | 284 |
| 9-HODE | 0 | 2 | 4 | 9 | 18 | 35 | 71 | 142 | 284 |
| 15-HETE | 0 | 2 | 4 | 8 | 17 | 33 | 67 | 133 | 266 |
| 13-HpODE | 0 | 4 | 7 | 14 | 28 | 56 | 112 | 224 | 448 |
| 17(18)-EpETE | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 141 | 282 |
| 13-KODE | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 138 | 276 |
| 9-HpODE | 0 | 2 | 4 | 8 | 17 | 34 | 67 | 135 | 269 |
| 17-HDoHE | 0 | 2 | 4 | 9 | 18 | 36 | 71 | 142 | 285 |
| 15-HpETE | 0 | 2 | 5 | 9 | 18 | 36 | 73 | 145 | 290 |
| 11-HETE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 15-KETE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 9-KODE | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 138 | 276 |
| 14(15)-EpETE | 0 | 2 | 4 | 8 | 17 | 33 | 66 | 132 | 265 |
| 12-HETE | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 141 | 282 |
| 8-HETE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 15(S)-HETrE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 12S-HpETE | 0 | 2 | 5 | 9 | 18 | 36 | 73 | 145 | 290 |
| 9-HETE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 12-KETE | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 141 | 282 |
| 5-HETE | 0 | 2 | 4 | 9 | 18 | 35 | 70 | 141 | 282 |
| 19(20)-EpDPE | 0 | 2 | 4 | 9 | 18 | 36 | 71 | 142 | 285 |
| 5(S)-HpETE | 0 | 2 | 5 | 9 | 18 | 36 | 72 | 144 | 289 |
| 12(13)-EpOME | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 137 | 274 |
| 14(15)-EpETrE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 9(10)-EpOME | 0 | 2 | 4 | 9 | 17 | 34 | 69 | 137 | 274 |
| 16(17)-EpDPE | 0 | 3 | 6 | 13 | 25 | 51 | 102 | 203 | 407 |
| 5-KETE | 0 | 1 | 2 | 4 | 8 | 16 | 32 | 64 | 127 |
| 11(12)-EpETrE | 0 | 2 | 4 | 9 | 17 | 35 | 70 | 140 | 280 |
| 8(9)-EpETrE | 0 | 3 | 6 | 13 | 25 | 50 | 101 | 201 | 402 |
| 5(6)-EpETrE | 0 | 2 | 5 | 10 | 19 | 38 | 77 | 153 | 306 |

**Table S-3** Statistics of the validation, including linearity ($R^2$), sensitivity (LOD/LOQ), reproducibility (precision and batch-to-batch effect) for the oxylipin library

| Compound Name | Retention Time | $R^2$ | LOD [nM] | LOQ [nM] | Precision RSD [%] | Batch-to-Batch Effect RSD [%] |
|---|---|---|---|---|---|---|
| Tetranor-PGFM | 2.05 | 0.985 | 0.6 | 2.1 | < 16 | 11-28 |
| Tetranor-PGEM | 2.50 | 0.995 | 1.2 | 4.0 | < 10 | 5-17 |
| 20-hydroxy PGF2α | 6.16 | 0.992 | 0.6 | 1.9 | < 15 | 6-15 |
| 20-hydroxy PGE2 | 6.22 | 0.994 | 1.4 | 4.5 | < 17 | 6-29 |
| Δ17-6-keto PGF1α | 7.09 | 0.993 | 0.5 | 1.5 | < 7 | 35-38 |
| 2,3-dinor-11b PGF2α | 7.89 | 0.998 | 0.2 | 0.7 | < 11 | 4-12 |
| 6-keto PGF1α | 7.89 | 0.994 | 0.8 | 2.6 | < 10 | 3-16 |
| 20-carboxy LTB4 | 8.10 | 0.991 | 3.6 | 12.1 | < 17 | 7-23 |
| 6-keto PGE1 | 8.14 | 0.993 | 0.5 | 1.6 | < 6 | 3-22 |
| 20-hydroxy LTB4 | 8.28 | 0.997 | 0.7 | 2.4 | < 9 | 6-26 |
| TXB3 | 8.33 | 0.989 | 0.2 | 0.6 | < 12 | 8-30 |
| PGF3α | 8.93 | 0.992 | 2.7 | 9.0 | < 25 | 5-67 |
| TXB1 | 9.04 | 0.989 | 0.4 | 1.3 | < 13 | 6-33 |
| 8-iso PGF2α | 9.23 | 0.995 | 1.0 | 3.2 | < 17 | 4-18 |
| PGE3 | 9.24 | 0.992 | 0.5 | 1.7 | < 14 | 5-26 |
| TXB2 | 9.24 | 0.995 | 0.3 | 1.0 | < 7 | 13-22 |
| 11β-PGF2α | 9.37 | - | - | - | - | - |
| PGD3 | 9.52 | 0.993 | 0.5 | 1.6 | < 16 | 6-16 |
| 5-iPF2a-VI | 9.63 | 0.994 | 0.4 | 1.5 | < 14 | 8-16 |
| 9,12,13-TriHOME | 9.83 | 0.946 | 0.2 | 0.8 | < 32 | 9-16 |
| PGF1α | 9.94 | 0.996 | 0.8 | 2.5 | < 14 | 5-12 |
| 9,10,13-TriHOME | 9.96 | 0.953 | 0.5 | 1.8 | < 10 | 10-17 |
| PGF2α | 9.96 | 0.998 | 0.9 | 2.9 | < 10 | 4-15 |
| PGE2 | 10.23 | 0.999 | 0.2 | 0.6 | < 13 | 4-20 |
| 11β-PGE2 | 10.40 | 0.998 | 0.9 | 3.1 | < 8 | 3-19 |
| 15-keto PGF2α | 10.45 | 0.998 | 0.5 | 1.7 | < 12 | 4-12 |
| PGE1 | 10.46 | 0.995 | 1.7 | 5.6 | < 26 | 4-28 |
| 5(S),14(R)-Lipoxin B4 | 10.51 | 0.990 | 2.6 | 8.5 | < 18 | 7-21 |
| 5(S),6(R)-Lipoxin A4 | 10.51 | 0.983 | 26.8 | 89.4 | < 43 | 12-53 |
| PGK2 | 10.53 | 0.373 | 21.6 | 72.1 | < 87 | 45-70 |
| PGD1 | 10.60 | 0.960 | 2.1 | 7.0 | < 24 | 5-25 |
| PGD2 | 10.61 | 0.994 | 0.3 | 0.9 | < 11 | 6-18 |
| 15-keto PGF1α | 10.70 | 0.995 | 1.0 | 3.3 | < 15 | 3-11 |
| 13,14-dihydro PGF2α | 10.79 | 0.990 | 11.4 | 38.2 | < 8 | 12-26 |
| 11β-13,14-dihydro-15-keto PGF2α | 10.98 | 0.997 | 5.7 | 18.9 | < 44 | 9-28 |
| 5(S),6(S)-Lipoxin A4 | 10.98 | 0.987 | 30.6 | 102.0 | < 23 | 8-21 |
| 13,14-dihydro-15-keto PGF2α | 11.40 | 0.996 | 1.0 | 3.5 | < 19 | 8-14 |
| 13,14-dihydro-15-keto PGE2 | 11.42 | 0.985 | 0.9 | 3.1 | < 8 | 4-28 |
| 14,15-LTC4 | 11.48 | - | - | - | - | - |
| LTD4 | 11.77 | 0.995 | 0.9 | 2.8 | < 17 | 4-22 |
| 14,15-LTE4 | 11.82 | 0.993 | 3.6 | 12.1 | < 22 | 11-30 |
| 13,14-dihydro-15-keto PGF1α | 11.88 | 0.995 | 0.7 | 2.2 | < 10 | 4-13 |
| 1α,1b-dihomo PGF2α | 11.88 | 0.997 | 0.6 | 2.1 | < 11 | 4-10 |
| 13,14-dihydro-15-keto PGD2 | 12.18 | 0.995 | 0.6 | 2.0 | < 11 | 5-16 |
| 11-trans LTD4 | 12.23 | 0.981 | 0.4 | 1.4 | < 34 | 7-32 |
| PGA2 | 12.58 | 0.998 | 0.3 | 1.1 | < 9 | 5-19 |
| 13,14-dihydro-15-keto PGD1 | 12.61 | 0.995 | 0.4 | 1.4 | < 12 | 4-16 |
| PGJ2 | 12.73 | 0.994 | 14.6 | 48.7 | < 17 | 4-15 |
| Δ12-PGJ2 | 12.73 | 0.996 | 0.5 | 1.5 | < 8 | 7-10 |
| LTE4 | 12.77 | 0.997 | 0.7 | 2.2 | < 8 | 5-51 |
| LTB5 | 12.90 | 0.983 | 1.5 | 5.0 | < 29 | 23-31 |
| LTC4 | 13.04 | - | - | - | - | - |
| 11-trans LTE4 | 13.18 | 0.987 | 3.2 | 10.6 | < 43 | 8-24 |
| 11-trans LTC4 | 13.57 | - | - | - | - | - |

## Continue Table S-3

| Compound Name | Retention Time | $R^2$ | LOD [nM] | LOQ [nM] | Precision RSD [%] | Batch-to-Batch Effect RSD [%] |
|---|---|---|---|---|---|---|
| 12,13-DiHODE | 13.62 | 0.953 | 2.7 | 9.0 | < 12 | 10-19 |
| 8(S),15(S)-DiHETE | 13.62 | 0.983 | 4.6 | 15.4 | < 9 | 10-17 |
| bicyclo-PGE2 | 13.80 | 0.995 | 1.7 | 5.6 | < 14 | 4-14 |
| 17,18-DiHETE | 14.01 | 0.990 | 4.7 | 15.5 | < 11 | 8-26 |
| 5(S),15(S)-DiHETE | 14.06 | 0.992 | 0.8 | 2.5 | < 8 | 7-16 |
| 6-trans-LTB4 | 14.09 | 0.995 | 1.2 | 4.1 | < 20 | 7-18 |
| 10(S),17(S)-DiHDoHE | 14.13 | 0.993 | 0.6 | 1.9 | < 11 | 5-15 |
| 15-deoxy-Δ12,14-PGD2 | 14.39 | 0.996 | 0.5 | 1.6 | < 6 | 6-14 |
| LTB4 | 14.39 | 0.997 | 1.1 | 3.6 | < 6 | 4-23 |
| 14,15-DiHETE | 14.49 | 0.993 | 0.8 | 2.7 | < 13 | 11-22 |
| 12,13-DiHOME | 14.80 | 0.975 | 0.3 | 1.0 | < 4 | 3-6 |
| 9,10-DiHOME | 15.18 | 0.981 | 0.5 | 1.8 | < 8 | 5-9 |
| 19,20-DiHDPA | 15.60 | 0.991 | 1.7 | 5.7 | < 11 | 6-30 |
| 14,15-DiHETrE | 15.65 | 0.997 | 0.3 | 1.0 | < 6 | 2-21 |
| 12S-HHTrE | 15.73 | 0.990 | 1.5 | 5.1 | < 12 | 8-36 |
| 11,12-DiHETrE | 16.23 | 0.993 | 0.5 | 1.7 | < 12 | 6-21 |
| 9-HOTrE | 16.57 | 0.967 | 0.3 | 1.1 | < 7 | 5-23 |
| 8,9-DiHETrE | 16.71 | 0.993 | 0.3 | 1.0 | < 15 | 10-17 |
| Hepoxilin A3 | 16.78 | 0.934 | 4.2 | 14.0 | < 76 | 22-50 |
| 20-HETE | 17.20 | 0.983 | 0.9 | 3.0 | < 41 | 8-33 |
| 5(S),6(S)-DiHETE | 17.20 | 0.992 | 0.4 | 1.4 | < 20 | 5-22 |
| 15(S)-HEPE | 17.28 | 0.983 | 0.9 | 3.1 | < 13 | 11-19 |
| 5,6-DiHETrE | 17.34 | 0.994 | 0.4 | 1.2 | < 9 | 7-21 |
| 12(S)-HEPE | 17.67 | 0.985 | 0.3 | 0.9 | < 12 | 6-23 |
| 5(S)-HEPE | 18.05 | 0.983 | 0.1 | 0.3 | < 12 | 8-24 |
| 13-HODE | 18.12 | 0.940 | 4.2 | 13.9 | < 9 | 7-16 |
| 9-HODE | 18.25 | 0.942 | 0.2 | 0.5 | < 7 | 7-22 |
| 15-HETE | 18.58 | 0.982 | 0.2 | 0.8 | < 11 | 6-18 |
| 13-HpODE | 18.60 | 0.939 | 3.5 | 11.8 | < 6 | 23-72 |
| 17(18)-EpETE | 18.60 | 0.994 | 0.5 | 1.7 | < 17 | 8-22 |
| 13-KODE | 18.70 | 0.978 | 0.7 | 2.4 | < 14 | 8-48 |
| 9-HpODE | 18.71 | 0.953 | 8.7 | 29.1 | < 7 | 29-72 |
| 17-HDoHE | 18.71 | 0.985 | 11.3 | 37.8 | < 17 | 7-29 |
| 15-HpETE | 18.93 | 0.985 | 2.3 | 7.7 | < 24 | 30-75 |
| 11-HETE | 18.97 | 0.987 | 0.3 | 0.9 | < 9 | 4-21 |
| 15-KETE | 19.03 | 0.988 | 0.8 | 2.8 | < 13 | 8-34 |
| 9-KODE | 19.09 | 0.951 | 3.2 | 10.8 | < 11 | 10-30 |
| 14(15)-EpETE | 19.09 | 0.988 | 0.8 | 2.6 | < 19 | 9-23 |
| 12-HETE | 19.24 | 0.986 | 0.2 | 0.6 | < 9 | 4-17 |
| 8-HETE | 19.25 | 0.990 | 1.4 | 4.7 | < 8 | 8-21 |
| 15(S)-HETrE | 19.41 | 0.988 | 0.5 | 1.7 | < 11 | 7-21 |
| 12S-HpETE | 19.43 | 0.539 | - | - | < 20 | 51-95 |
| 9-HETE | 19.47 | 0.963 | 6.5 | 21.7 | < 12 | 9-25 |
| 12-KETE | 19.56 | - | - | - | - | - |
| 5-HETE | 19.69 | 0.988 | 0.2 | 0.6 | < 11 | 14-20 |
| 19(20)-EpDPE | 19.93 | 0.988 | 1.2 | 4.0 | < 12 | 6-18 |
| 5(S)-HpETE | 20.04 | 0.968 | 8.1 | 27.0 | < 19 | 34-71 |
| 12(13)-EpOME | 20.08 | 0.964 | 2.5 | 8.4 | < 7 | 14-39 |
| 14(15)-EpETrE | 20.19 | 0.986 | 0.9 | 3.1 | < 27 | 7-21 |
| 9(10)-EpOME | 20.27 | 0.932 | 0.6 | 1.9 | < 11 | 10-38 |
| 16(17)-EpDPE | 20.32 | 0.994 | 1.0 | 3.5 | < 21 | 8-15 |
| 5-KETE | 20.51 | 0.990 | 0.4 | 1.2 | < 15 | 9-17 |
| 11(12)-EpETrE | 20.71 | 0.988 | 0.4 | 1.3 | < 23 | 16-22 |
| 8(9)-EpETrE | 20.79 | 0.985 | 1.4 | 4.5 | < 17 | 10-19 |
| 5(6)-EpETrE | 20.95 | - | - | - | - | - |

**Table S-4** Detected amounts [nM] of oxylipins before and 24 hours after surgery.

| | Patient 1 | | Patient 2 | | Patient 3 | | Patient 4 | | Patient 5 | |
|---|---|---|---|---|---|---|---|---|---|---|
| **Compound name** | **T0** | **T 24h** | **T0** | **T 24h** | **T0** | **T 24h** | **T0** | **T 24h** | **T0** | **T 24h** |
| **Arachidonic Acid** | | | | | | | | | | |
| TXB2 | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | 1.5 (±0.3) |
| PGF2α | 8.7 (±1.5) | 9.3 (±2.0) | 11.9 (±3.1) | 15.5 (±10.1) | 7.4 (±2.0) | 9.9 (±2.0) | 8.7 (±3.3) | 6.1 (±0.4) | 5.8 (±1.1) | 6.7 (±0.7) |
| PGE2 | <LOQ | 0.7 (±0.2) | 0.7 (±0.3) | 0.8 (±0.1) | 1.2 (±1.0) | <LOQ | <LOQ | 1.5 (±1.0) | <LOQ | 0.9 (±0.2) |
| 11β-PGE2 | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ |
| 13,14-dihydro PGF2α | 48.9 (±3.0) | 52.5 (±6.2) | 65.7 (±18.8) | 69.1 (±18.5) | 49.2 (±5.8) | 55.8 (±6.2) | 50.6 (±14.1) | 43.9 (±4.8) | 51.5 (±4.9) | 47.8 (±13.0) |
| 14,15-DiHETrE | 4.6 (±0.7) | 3.6 (±0.2) | 1.9 (±0.2) | 2.7 (±0.3) | 4.4 (±0.2) | 3.1 (±0.4) | 2.8 (±0.5) | 1.9 (±0.2) | 2.3 (±0.2) | 3.4 (±0.4) |
| 11,12-DiHETrE | 3.5 (±0.4) | 3.1 (±0.8) | <LOQ | 2.1 (±0.9) | 2.0 (±0.2) | 1.9 (±0.3) | 1.8 (±0.3) | <LOQ | <LOQ | 2.5 (±0.9) |
| 8,9-DiHETrE | 1.6 (±0.2) | 1.2 (±0.5) | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ |
| 5,6-DiHETrE | 1.7 (±0.8) | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ |
| 15-HETE | 3.4 (±1.1) | 2.1 (±0.2) | 1.5 (±0.5) | 2.3 (±0.5) | 1.9 (±0.9) | 2.0 (±0.4) | 1.5 (±0.3) | 1.1 (±0.5) | 1.3 (±0.3) | 1.9 (±1.3) |
| 11-HETE | 1.6 (±0.7) | <LOQ | <LOQ | <LOQ | <LOQ | 1.3 (±0.4) | <LOQ | <LOQ | <LOQ | 1.4 (±0.2) |
| 12-HETE | 4.6 (±0.8) | 7.3 (±2.4) | 1.3 (±0.4) | 9.3 (±2.4) | 1.8 (±0.4) | 5.6 (±0.6) | 3.9 (±1.6) | 8.2 (±3.2) | 1.5 (±0.9) | 62.9 (±8.4) |
| 8-HETE | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ |
| 5-HETE | 2.5 (±0.6) | 1.8 (±0.2) | 1.0 (±0.6) | 2.4 (±0.4) | 1.2 (±0.4) | 3.9 (±0.6) | <LOQ | 1.1 (±0.4) | 1.1 (±0.3) | 2.3 (±0.4) |
| **Linoleic Acid** | | | | | | | | | | |
| 9,12,13-TriHOME | 3.6 (±0.8) | 3.1 (±0.6) | 2.3 (±0.3) | 2.7 (±0.4) | 10.7 (±1.0) | 2.6 (±0.5) | 18.9 (±2.1) | 2.8 (±0.6) | 25.2 (±2.7) | 6.7 (±5.2) |
| 9,10,13-TriHOME | 2.1 (±0.6) | 1.8 (±0.2) | <LOQ | <LOQ | 6.5 (±0.7) | <LOQ | 5.4 (±1.2) | <LOQ | 4.9 (±0.2) | 5.1 (±3.9) |
| 12,13-DiHOME | 14.0 (±2.4) | 7.6 (±0.7) | 12.6 (±0.6) | 10.6 (±1.1) | 63.5 (±4.0) | 8.1 (±0.6) | 40.9 (±2.5) | 7.0 (±1.8) | 39.9 (±2.9) | 23.6 (±16.0) |
| 9,10-DiHOME | 7.7 (±2.5) | 4.9 (±1.4) | 2.6 (±0.5) | 4.2 (±1.2) | 80.7 (±8.4) | 3.5 (±0.7) | 15.3 (±2.6) | 4.3 (±0.5) | 20.8 (±0.9) | 24.5 (±23.5) |
| 13-HODE | 29.3 (±12.6) | 25.1 (±3.7) | 35.7 (±3.2) | 32.2 (±6.3) | 104.8 (±4.9) | 28.0 (±3.1) | 86.5 (±3.8) | <LOQ | 70.3 (±6.3) | 43.5 (±18.0) |
| 9-HODE | 24.2 (±1.1) | 21.2 (±2.6) | 18.1 (±1.3) | 26.3 (±3.3) | 55.0 (±4.3) | 23.0 (±2.5) | 83.7 (±7.5) | 13.9 (±0.8) | 92.2 (±3.4) | 39.3 (±13.6) |
| 13-HpODE | 200.3 (±86.1) | 415.0 (±220.5) | 143.7 (±96.0) | 445.8 (±234.2) | 513.8 (±82.8) | 195.9 (±97.4) | 105.5 (±38.4) | 87.4 (±27.1) | 80.1 (±19.5) | 85.5 (±45.3) |
| 13-KODE | 9.5 (±3.1) | 12.3 (±6.1) | 6.3 (±1.6) | 16.1 (±8.7) | 19.7 (±1.5) | 6.8 (±2.5) | 6.2 (±0.4) | 3.0 (±1.2) | 5.8 (±0.3) | 4.1 (±0.9) |
| 9-HpODE | 146.2 (±26.4) | 227.2 (±97.0) | 88.2 (±37.3) | 221.7 (±141.4) | 287.0 (±32.9) | 142.5 (±54.4) | 69.1 (±36.5) | 67.5 (±44.3) | 60.5 (±21.0) | 72.1 (±15.5) |
| 9-KODE | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ |
| 12(±13)-EpOME | 11.3 (±8.2) | 18.9 (±6.4) | 17.1 (±4.5) | 26.1 (±16.2) | 26.5 (±5.5) | 9.7 (±1.3) | 27.2 (±4.6) | <LOQ | 24.9 (±4.1) | 9.1 (±2.9) |
| 9(±10)-EpOME | 7.6 (±4.2) | 16.8 (±4.1) | 4.2 (±1.1) | 13.0 (±5.9) | 16.3 (±2.9) | 3.8 (±2.0) | 4.6 (±1.6) | 2.5 (±0.9) | 5.0 (±1.1) | 2.7 (±2.1) |
| **Dihomo-gamma-linolenic acid** | | | | | | | | | | |
| PGF1α | <LOQ | <LOQ | 5.9 (±5.5) | 4.7 (±3.5) | <LOQ | <LOQ | 2.7 (±1.6) | <LOQ | <LOQ | <LOQ |
| 15(±S)-HETrE | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ |
| **Alpha-linolenic acid** | | | | | | | | | | |
| 9-HOTrE | 1.2 (±0.5) | <LOQ | 1.5 (±0.4) | 1.1 (±0.2) | 3.2 (±0.7) | <LOQ | 7.7 (±1.1) | <LOQ | 8.4 (±0.7) | <LOQ |
| **Eicosapentaenoic acid** | | | | | | | | | | |
| 17,18-DiHETE | 21.0 (±6.3) | 17.5 (±2.8) | 27.5 (±1.0) | <LOQ | 27.2 (±3.3) | 22.6 (±2.2) | 31.4 (±9.3) | <LOQ | 16.0 (±3.0) | 16.5 (±0.6) |
| 14,15-DiHETE | 6.0 (±2.1) | 2.9 (±1.0) | 4.0 (±1.2) | 2.8 (±1.2) | 7.1 (±0.7) | 5.6 (±2.7) | 5.2 (±1.5) | 3.3 (±1.1) | 3.5 (±0.9) | 2.9 (±1.5) |
| 15(±S)-HEPE | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ | <LOQ |
| 12(±S)-HEPE | 1.2 (±0.1) | 1.3 (±0.5) | 1.0 (±0.8) | 2.1 (±0.8) | <LOQ | 1.2 (±0.3) | 1.0 (±0.4) | 1.6 (±0.8) | <LOQ | 4.8 (±0.8) |
| 5(±S)-HEPE | 1.8 (±0.9) | 0.7 (±0.1) | 1.3 (±0.1) | 1.6 (±0.5) | 0.8 (±0.3) | 1.7 (±0.3) | 0.6 (±0.2) | 1.0 (±0.2) | 0.4 (±0.2) | 0.9 (±0.2) |
| **Docosahexaenoic acid** | | | | | | | | | | |
| 19,20-DiHDPA | 10.3 (±1.5) | 10.9 (±1.0) | 9.9 (±1.4) | 8.8 (±3.1) | 13.3 (±0.6) | 12.3 (±2.2) | 10.8 (±1.0) | <LOQ | <LOQ | 6.7 (±0.7) |
| 19(±20)-EpDPE | 7.2 (±2.2) | 4.6 (±1.1) | <LOQ | 8.6 (±7.7) | <LOQ | 4.0 (±0.3) | <LOQ | <LOQ | <LOQ | 4.7 (±1.4) |