SUPPORTING INFORMATION

Reversed-phase UHPLC/ESI-MS determination of oxylipins in human plasma: case study of female breast cancer

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5 FIGURES, 5 TABLES



Fig. S-1 MS/MS spectra of 63 oxylipins with multiple product ions at optimized collision energy

Fig. S-1 (Continuation)



Fig. S-1 (Continuation)







Fig. S-1 (Continuation)



Fig. S-1 (Continuation)



Fig. S-2 MS/MS spectra of selected standard **a** 12-HETE at various collision energies **b** -15V, **c** -18V, **d** -20V, **e** -23V, **f** -25V, **g** -28V, and **h** -30V





Fig. S-3 Monitoring QC peak areas during the method validation



Fig. S-4 Receiver operating characteristic (ROC) and area under the curve (AUC) values

1- specificity

Fig. S-5 S-plot of 21 quantified oxylipins in plasma samples of breast cancer patients, where the most up-regulated lipid species are in the upper right corner



Table S-1 Oxylipin standards with their systematic names used for UHPLC/MS method development

| Lipid class | [M-H] ⁻ | Oxylipin species | Systematic name | | | | |
|-------------------|--------------------|-----------------------|---|--|--|--|--|
| FA and conjugates | 265 | tetranor-12-HETE | 8R-hydroxy-4Z,6E,10Z-hexadecatrienoic acid | | | | |
| Octadecanoids | 293 | 9-HOTrE | 9S-hydroxy-10E,12Z,15Z-octadecatrienoic acid | | | | |
| | 293 | 13-HOTrE | 13S-hydroxy-9Z,11E,15Z-octadecatrienoic acid | | | | |
| | 293 | 13-OxoODE | 13-keto-9Z,11E-octadecadienoic acid | | | | |
| | 293 | 9-OxoODE | 9-keto-10E,12Z-octadecadienoic acid | | | | |
| | 295 | 13-HODE | 13S-hydroxy-9Z,11E-octadecadienoic acid | | | | |
| | 295 | 9-HODE | 9S-hydroxy-10E,12Z-octadecadienoic acid | | | | |
| | 295 | 12(13)-EpOME | (±)-12(13)-epoxy-9Z-octadecenoic acid | | | | |
| | 295 | 9(10)-EpOME | 9,10-epoxy-12Z-octadecenoic acid | | | | |
| | 313 | 12,13-DiHOME | 12,13-dihydroxy-9Z-octadecenoic acid | | | | |
| | 313 | 9,10-DiHOME | 9,10-dihydroxy-12Z-octadecenoic acid | | | | |
| Eicosanoids | 279 | 12-HHTrE | 12S-hydroxy-5Z,8E,10E-heptadecatrienoic acid | | | | |
| | 317 | 15-HEPE | (±)-15-hydroxy-5Z,8Z,11Z,13E,17Z-eicosapentaenoic acid | | | | |
| | 317 | 11-HEPE | (±)-11-hydroxy-5Z,8Z,12E,14Z,17Z-eicosapentaenoic acid | | | | |
| | 317 | 5-HEPE | (±)-5-hydroxy-6E,8Z,11Z,14Z,17Z-eicosapentaenoic acid | | | | |
| | 317 | 14(15)-EpETE | (±)-14(15)-epoxy-5Z,8Z,11Z,17Z-eicosatetraenoic acid | | | | |
| | 317 | 12-OxoETE | 12-oxo-5Z,8Z,10E,14Z-eicosatetraenoic acid | | | | |
| | 317 | 15-OxoETE | 15-oxo-5Z,8Z,11Z,13E-eicosatetraenoic acid | | | | |
| | 317 | 5-OxoETE | 5-oxo-6E,8Z,11Z,14Z-eicosatetraenoic acid | | | | |
| | 319 | 15-HETE | 15S-hydroxy-5Z,8Z,11Z,13E-eicosatetraenoic acid | | | | |
| | 319 | 11-HETE | 11S-hydroxy-5Z,8Z,12E,14Z-eicosatetraenoic acid | | | | |
| | 319 | 12-HETE | 12S-hydroxy-5Z,8Z,10E,14Z-eicosatetraenoic acid | | | | |
| | 319 | 8-HETE | 8S-hydroxy-5Z,9E,11Z,14Z-eicosatetraenoic acid | | | | |
| | 319 | 9-HETE | 9-hydroxy-5Z,7E,11Z,14Z-eicosatetraenoic acid | | | | |
| | 319 | 5-HETE | 5S-hydroxy-6E,8Z,11Z,14Z-eicosatetraenoic acid | | | | |
| | 319 | 11,12-EET | 11,12-epoxy-5Z,8Z,14Z-eicosatrienoic acid | | | | |
| | 319 | 5,6-EET | 5,6-epoxy-8Z,11Z,14Z-eicosatrienoic acid | | | | |
| | 321 | 15-HETrE | 15S-hydroxy-8Z,11Z,13E-eicosatrienoic acid | | | | |
| | 321 | 5-HETrE | 5S-hydroxy-6E,8Z,11Z-eicosatrienoic acid | | | | |
| | 327 | tetranor-PGDM | 9S-hydroxy-11,15-dioxo-2,3,4,5-tetranor-prostan-1,20-dioic acid | | | | |
| | 333 | PGJ2 | 11-oxo-15S-hydroxy-5Z,9,13E-prostatrienoic acid | | | | |
| | 333 | PGB2 | 15S-hydroxy-9-oxo-5Z,8(12),13E-prostatrienoic acid | | | | |
| | 333 | PGA2 | 9-oxo-15S-hydroxy-5Z,10Z,13E-prostatrienoic acid | | | | |
| | 333 | 15-deoxy-δ-12,14 PGD2 | 9S-hydroxy-11-oxo-5Z,12E,14E-prostatrienoic acid | | | | |
| | 335 | 8,15 DiHETE | 8S,15S-dihydroxy-5Z,9E,11Z,13E-eicosatetraenoic acid | | | | |
| | 335 | 6-trans LTB4 | 5S,12R-dihydroxy-6E,8E,10E,14Z-eicosatetraenoic acid | | | | |
| | 335 | LTB4 | 5S,12R-dihydroxy-6Z,8E,10E,14Z-eicosatetraenoic acid | | | | |
| | 335 | 5,15-DiHETE | 5S,15S-dihydroxy-6E,8Z,10Z,13E-eicosatetraenoic acid | | | | |
| | 335 | 5,6- Dihete | 5S,6S-dihydroxy-8Z,11Z,14Z,17Z-eicosatetraenoic acid | | | | |

| | 337 | 14,15-DiHETrE | 14,15-dihydroxy-5Z,8Z,11Z-eicosatrienoic acid |
|---------------|-----|------------------------------|---|
| | 337 | 5,6 -DiHETrE | 5,6-dihydroxy-8Z,11Z,14Z-eicosatrienoic acid |
| | 351 | PGH2 | 9S,11R-epidioxy-15S-hydroxy-5Z,13E-prostadienoic acid |
| | 351 | PGE2 | 9-oxo-11R,15S-dihydroxy-5Z,13E-prostadienoic acid |
| | 351 | 11β-PGE2 | 9-oxo-11S,15S-dihydroxy-5Z,13E-prostadienoic acid |
| | 351 | 15-keto-PGF2α | 9S,11R-dihydroxy-15-oxo-5Z,13E-prostadienoic acid |
| | 351 | PGD2 | 9S,15S-dihydroxy-11-oxo-5Z,13E-prostadienoic acid |
| | 351 | 13,14-dh-15-k-PGE2 | 9,15-dioxo-11R-hydroxy-5Z-prostenoic acid |
| | 353 | 8-iso-PGF2a | 9S,11R,15S-trihydroxy-5Z,13E-prostadienoic acid-cyclo[8S,12R] |
| | 353 | (+/-) 5-iPF2α-VI | 5,9S,11R-trihydroxy-6E,14Z-prostadienoic acid-cyclo[8S,12R] |
| | 353 | PGF2a | 9S,11R,15S-trihydroxy-5Z,13E-prostadienoic acid |
| | 353 | 13,14-dh-15-k-PGF2α | 9S,11S-dihydroxy-15-oxo-5Z-prostenoic acid |
| | 369 | 6-keto-PGF1α | 6-oxo-9S,11R,15S-trihydroxy-13E-prostenoic acid |
| | 369 | TXB2 | 9S,11,15S-trihydroxy-thromboxa-5Z,13E-dien-1-oic acid |
| Docosanoids | 343 | 20-HDoHE | (±)-20-hydroxy-4Z,7Z,10Z,13Z,16Z,18E-docosahexaenoic acid |
| | 343 | 17-HDoHE | (±)-17-hydroxy-4Z,7Z,10Z,13Z,15E,19Z-docosahexaenoic acid |
| | 343 | 10-HDoHE | (±)-10-hydroxy-4Z,7Z,11E,13Z,16Z,19Z-docosahexaenoic acid |
| | 343 | 14-HDoHE | (±)-14-hydroxy-4Z,7Z,10Z,12E,16Z,19Z-docosahexaenoic acid |
| | 343 | 11-HDoHE | (±)-11-hydroxy-4Z,7Z,9E,13Z,16Z,19Z-docosahexaenoic acid |
| | 343 | 7-HDoHE | (±)-7-hydroxy-4Z,8E,10Z,13Z,16Z,19Z-docosahexaenoic acid |
| | 343 | 8-HDoHE | (±)-8-hydroxy-4Z,6E,10Z,13Z,16Z,19Z-docosahexaenoic acid |
| | 343 | 4-HDoHE | (±)4-hydroxy-5E,7Z,10Z,13Z,16Z,19Z-docosahexaenoic acid |
| | 361 | 19,20-DiHDPE | (±)-19,20-dihydroxy-4Z,7Z,10Z,13Z,16Z-docosapentaenoic acid |
| | 375 | Resolvin D1 | 7S,8R,17S-trihydroxy-4Z,9E,11E,13Z,15E,19Z-docosahexaenoic acid |
| Deuterated IS | 299 | D4-13-HODE | (+)-13-hydroxyl-9Z 11E_octadecadienoic_9 10 12 13-d. acid |
| | 299 | D₄-9-HODE | (\pm) 9-hydroxyl-10E.12Z-octadecadienoic-9.10.12.13-d4 acid |
| | 317 | D4-9 10-DiHOME | 9 10-dihydroxy-127-octadecenoic-9 10 12 13-daacid |
| | 317 | D ₄ -12.13-DiHOME | (\pm) -12.1-dihydroxy-9Z-octadecenoic-9.10.12.13-d4 acid |
| | 323 | D7-5-0X0ETE | 5-oxo-6F.8Z.11Z.14Z-eicosatetraenoic-6.8.9.11.12.14.15-d7 acid |
| | 325 | D ₆ -20-HETE | (±)-20-hvdroxy-5Z.8Z.11Z.14Z-eicosatetraenoic-16.16.17.17.18.18-d ₆ acid |
| | 327 | Ds-15-HETE | (±)-15-hvdroxy-5Z.8Z.11Z.13E-eicosatetraenoic-5.6.8.9.11.12.14.15-ds acid |
| | 327 | D8-12-HETE | (±)-12-hvdroxy-5Z.8Z.10E.14Z-eicosatetraenoic-5.6.8.9.11.12.14.15-d ₈ acid |
| | 327 | D ₈ -5-HETE | (±)-5-hydroxy - 6E.8Z.11Z.14Z-eicosatetraenoic-5.6.8.9.11.12.14.15-d ₈ acid |
| | 330 | D ₁₁ -14,15-EET | (±)-14(15)-epoxy-5Z.8Z.11Z-eicosatrienoic-16,16,17,17,18,18,19,19,20,20,20-d ₁₁ acid |
| | 330 | D11-8,9-EET | (±)-8(9)-epoxy-5Z,8Z,14Z-eicosatrienoic-16.16,17,17,18,18,19,19,20,20,20-du acid |
| | 330 | D ₁₁ -11,12-EET | (±)-11(12)-epoxy-5Z,8Z,14Z-eicosatrienoic-16,16,17,17,18,18,19,19,20,20,20,-du acid |
| | 339 | D4-LTB4 | 5S,12R-dihydroxy-6Z,8E,10E,14Z-eicosatetraenoic-6,7,14,15-d4 acid |
| | 357 | D₄-PGF2α | 9S.11R.15S-trihvdroxy-5Z.13E-prostadienoic-3.3.4.4-deacid |

| Table | S-2 Information table about control and breast cancer patients. $BMI = body mas$ | 3 S |
|--------|---|------------|
| index, | cancer stages: pTis = tumor in situ, pT1 = tumors 1 mm or less, pT2 = tumors from | m |
| 1.01 – | 2 mm, and $pTx = not assessed$ | |

| Healthy volunteers | Age | Breast cancer patients | t cancer Age tients | | Smoker | Stage of cancer | |
|-----------------------|-----------------|---------------------------|------------------------|-------|--------|-----------------|--|
| 1 | 63 | 21 | 60 | 30.12 | no | pTis | |
| 2 | 51 | 22 | 54 | 25.31 | no | pT1b | |
| 3 | 50 | 23 | 56 | 23.05 | yes | pT1c | |
| 4 | 59 | 24 | 59 | 22.06 | yes | pT1b | |
| 5 | 62 | 25 | 65 | 32.05 | yes | pTis | |
| 6 | 55 | 26 | 50 | - | yes | pTis | |
| 7 | 60 | 27 | 65 | 37.34 | yes | pT1c | |
| 8 | 57 | 28 | 59 | 27.45 | no | pT1a | |
| 9 | 49 | 29 | 48 | 20.20 | yes | pT1c | |
| 10 | 55 | 30 | 53 | 33.86 | yes | pT1c | |
| 11 | 59 | 31 | 57 | - | - | pTis | |
| 12 | 57 | 32 | 62 | 26.37 | no | pT1b | |
| 13 | 56 | 33 | 50 | 23.72 | - | pT2 | |
| 14 | 57 | 34 | 52 | 25.39 | no | pT1a | |
| 15 | 61 | 35 | 55 | 25.33 | yes | pT1c | |
| 16 | 58 | 36 | 53 | 24.53 | no | pTx | |
| 17 | 56 | 37 | 54 | 21.56 | yes | pT1c | |
| 18 | 48 | 38 | 61 | 24.68 | no | pT1c | |
| 19 | 53 | 39 | 52 | 22.49 | no | pT1b | |
| 20 | 56 | 40 | 63 | - | - | pT1c | |
| Median ¹ | 56.5 ± 4.10 | Median ¹ | 55.5 ± 4.99 | | | | |

 1 Median \pm standard deviation

Table S-3 Parameters of validation for 14 deuterated oxylipin internal standards for high level

 (HL) and low level (LL) concentrations

| Internal standards | LOD [ng/ml] | LOQ [ng/ml] | Slope | Intercept | Correlation coefficient | Matrix effect ^a [%] | Recovery rate HL [%] | Recovery rate LL [%] |
|----------------------------------|----------------|----------------|-------|-----------|----------------------------|-----------------------------------|-------------------------|-------------------------|
| D ₄ -PGF2a | 1.50 | 5 | 3406 | 5205 | 0.9992 | 67.9 ± 4.8 | 87.5 | 83.3 |
| D ₄ -LTB ₄ | 0.30 | 1 | 3863 | 3133 | 0.9995 | 110 ± 9.3 | 89.5 | 79.8 |
| D ₄ -9-HODE | 0.15 | 0.5 | 4801 | 6209 | 0.9996 | 127.3 ± 9.8 | 94.4 | 77.7 |
| D ₄ -13-HODE | 0.15 | 0.5 | 5529 | 8057 | 0.9995 | 109.3 ± 13.9 | 98.3 | 79.2 |
| D ₄ -9,10-DiHOME | 0.15 | 0.5 | 3071 | 4324 | 0.9994 | 130.5 ± 16.4 | 97.8 | 91.1 |
| D ₄ -12,13-DiHOME | 0.60 | 2 | 3406 | 5205 | 0.9994 | 120.6 ± 19.5 | 93.7 | 102.8 |
| D7-5-0x0ETE | 0.60 | 2 | 225.3 | 162.2 | 0.9996 | 110.5 ± 18.3 | 89.0 | 78.6 |
| D ₆ -20-HETE | 0.60 | 2 | 1515 | 1253 | 0.9994 | 145.5 ± 10.8 | 95.4 | 71.2 |
| D ₈ -15-HETE | 0.15 | 0.5 | 4697 | 3212 | 0.9991 | 152.8 ± 15.8 | 96.0 | 92.7 |
| D ₈ -12-HETE | 0.15 | 0.5 | 1171 | 1661 | 0.9995 | 159 ± 15.8 | 88.1 | 83.9 |
| D ₈ -5-HETE | 0.60 | 2 | 2297 | 2307 | 0.9993 | 162.1 ± 11 | 94.6 | 81.1 |
| D ₁₁ -8,9-EET | 0.15 | 0.5 | 290.6 | 280.2 | 0.9995 | 159.1 ± 12.6 | 84.3 | 95.9 |
| D11-11,12-EET | 0.15 | 0.5 | 1117 | 2608 | 0.9995 | 107 ± 18.7 | 89.5 | 94.0 |
| D ₁₁ -14,15-EET | 0.15 | 0.5 | 473.6 | 558.4 | 0.9994 | 120 ± 19.1 | 89.4 | 86.2 |

^aMean \pm standard deviation

Table S-4 Precision and accuracy for 14 deuterated IS for high level (HL) and low level (LL)

 concentrations

| Internal standards | Within-run precision HL [%] | Between-run precision HL [%] | Within-run precision LL [%] | Between-run precision LL [%] | Within-run accuracy HL [%] | Between-run accuracy HL [%] | Within-run accuracy LL [%] | Between-run accuracy LL [%] |
|----------------------------------|-----------------------------------|------------------------------------|-----------------------------------|------------------------------------|----------------------------------|-----------------------------------|----------------------------------|-----------------------------------|
| D₄-PGF2α | 9.5 | 12.8 | 5.3 | 8.3 | 91.0 | 93.6 | 80.1 | 79.5 |
| D ₄ -LTB ₄ | 7.7 | 8.7 | 9.2 | 10.9 | 101.6 | 100.2 | 103.7 | 108.6 |
| D ₄ -9-HODE | 7.7 | 11.6 | 11.2 | 11.7 | 103.1 | 111.8 | 111.4 | 113.0 |
| D ₄ -13-HODE | 6.0 | 9.4 | 12.7 | 8.3 | 101.1 | 110.7 | 92.6 | 103.7 |
| D ₄ -9,10-DiHOME | 6.2 | 13.5 | 6.6 | 8.3 | 94.6 | 88.1 | 95.2 | 106.6 |
| D ₄ -12,13-DiHOME | 5.4 | 9.2 | 5.6 | 6.7 | 99.4 | 98.4 | 107.3 | 111.7 |
| D7-5-0x0ETE | 5.2 | 6.8 | 15.0 | 10.1 | 96.5 | 104.2 | 100.8 | 104.7 |
| D ₆ -20-HETE | 8.2 | 12.6 | 24.8 | 25.3 | 112.5 | 117.0 | 115.6 | 123.0 |
| D ₈ -15-HETE | 9.2 | 10.4 | 10.5 | 11.5 | 99.3 | 101.9 | 104.6 | 92.8 |
| D ₈ -12-HETE | 6.1 | 7.6 | 8.9 | 5.72 | 101.7 | 98.6 | 94.7 | 104.1 |
| D ₈ -5-HETE | 5.9 | 8.7 | 11.5 | 9.3 | 89.3 | 88.9 | 98.2 | 86.0 |
| D ₁₁ -8,9-EET | 6.8 | 7.5 | 8.9 | 9.7 | 109.9 | 110.2 | 104.9 | 110.5 |
| D ₁₁ -11,12-EET | 6.1 | 7.3 | 5.5 | 10.5 | 102.2 | 105.2 | 97.8 | 114.9 |
| D ₁₁ -14,15-EET | 6.3 | 7.1 | 5.8 | 5.0 | 112.6 | 86.1 | 112.4 | 113.1 |

Table S-6 Average concentrations (pmol/mL) of oxylipins in human plasma of healthy volunteers (normal) and breast cancer patients (tumor) with statistical parameters; VIP – variable importance in projection

| Oxylipin species | Normal ^a | Tumor ^a | Fold change | p-value ^b | T-value | VIP- value ^c |
|--------------------|---------------------|--------------------|----------------|----------------------|---------|----------------------------|
| 9-HODE | 18.13 ± 7.16 | 30.85 ± 15.89 | 1.70 | 3.75E-03 | -3.26 | 1.77 |
| 13-HOTrE | 2.04 ± 1.11 | 3.10 ± 1.14 | 1.52 | 6.16E-03 | -2.98 | 1.76 |
| 13-HODE | 16.50 ± 5.97 | 22.40 ± 8.57 | 1.36 | 1.91E-02 | -2.53 | 1.40 |
| 19,20-DiHDPE | 1.79 ± 1.03 | 2.58 ± 1.02 | 1.44 | 2.28E-02 | -2.44 | 1.26 |
| 9-HOTrE | 4.00 ± 2.12 | 5.96 ± 3.48 | 1.53 | 2.61E-02 | -2.39 | 1.18 |
| 12-HHTrE | 0.51 ± 0.60 | 0.96 ± 0.66 | 1.90 | 3.36E-02 | -2.26 | 1.59 |
| 14,15-DiHETrE | 3.17 ± 1.35 | 4.04 ± 1.58 | 1.28 | 7.62E-02 | -1.87 | 1.18 |
| 15-HETE | 0.93 ± 0.50 | 1.36 ± 0.93 | 1.46 | 8.73E-02 | -1.82 | 1.11 |
| 12-HETE | 13.96 ± 13.47 | 27.32 ± 31.88 | 1.73 | 1.37E-01 | -1.57 | 0.67 |
| LTB4 | 0.24 ± 0.20 | 0.33 ± 0.22 | 1.38 | 1.89E-01 | -1.37 | 1.14 |
| 9-HETE | 1.18 ± 0.36 | 1.91 ± 2.67 | 1.39 | 2.01E-01 | -1.34 | 0.80 |
| 11-HETE | 0.88 ± 0.45 | 1.54 ± 2.32 | 1.74 | 2.42E-01 | -1.24 | 0.84 |
| 14-HDoHE | 2.81 ± 3.53 | 4.38 ± 4.54 | 1.56 | 2.42E-01 | -1.22 | 0.65 |
| 7-HDoHE | 9.03 ± 10.82 | 13.21 ± 12.35 | 1.46 | 2.75E-01 | -1.14 | 0.78 |
| 20-HDoHE | 1.28 ± 0.65 | 1.53 ± 0.77 | 1.20 | 2.80E-01 | -1.12 | 0.52 |
| 4-HDoHE | 3.34 ± 2.03 | 3.12 ± 1.65 | 0.91 | 6.32E-01 | 0.50 | 0.16 |
| 5,6-EET | 1.35 ± 0.82 | 1.25 ± 0.74 | 0.93 | 6.92E-01 | 0.41 | 0.27 |
| 6-trans LTB4 | 0.26 ± 0.08 | 0.25 ± 0.07 | 0.96 | 6.94E-01 | 0.41 | 0.23 |
| 11,1 2-EE T | 3.47 ± 2.84 | 3.31 ± 2.45 | 0.92 | 7.35E-01 | 0.35 | 0.07 |
| 11-HDoHE | 0.48 ± 0.47 | 0.50 ± 0.36 | 1.04 | 9.01E-01 | -0.13 | 0.32 |
| 5,6-DiHETrE | 0.52 ± 0.28 | 0.51 ± 0.25 | 0.99 | 9.35E-01 | 0.08 | 0.09 |

^a Mean \pm standard deviation

^bCalculated using T-test

^c Generated from OPLS-DA model.