

2011/10/01

第2回 機能油脂懇話会

脂肪酸由来の生理活性物質と 炎症の制御

東京大学大学院薬学系研究科
JSTさがけ「炎症の慢性化機構の解明と制御」

有田 誠



Lipid Mediators from Polyunsaturated Fatty Acids

Arachidonic acid (AA) → Prostaglandins
C20:4 ω 6
Leukotrienes
Lipoxins



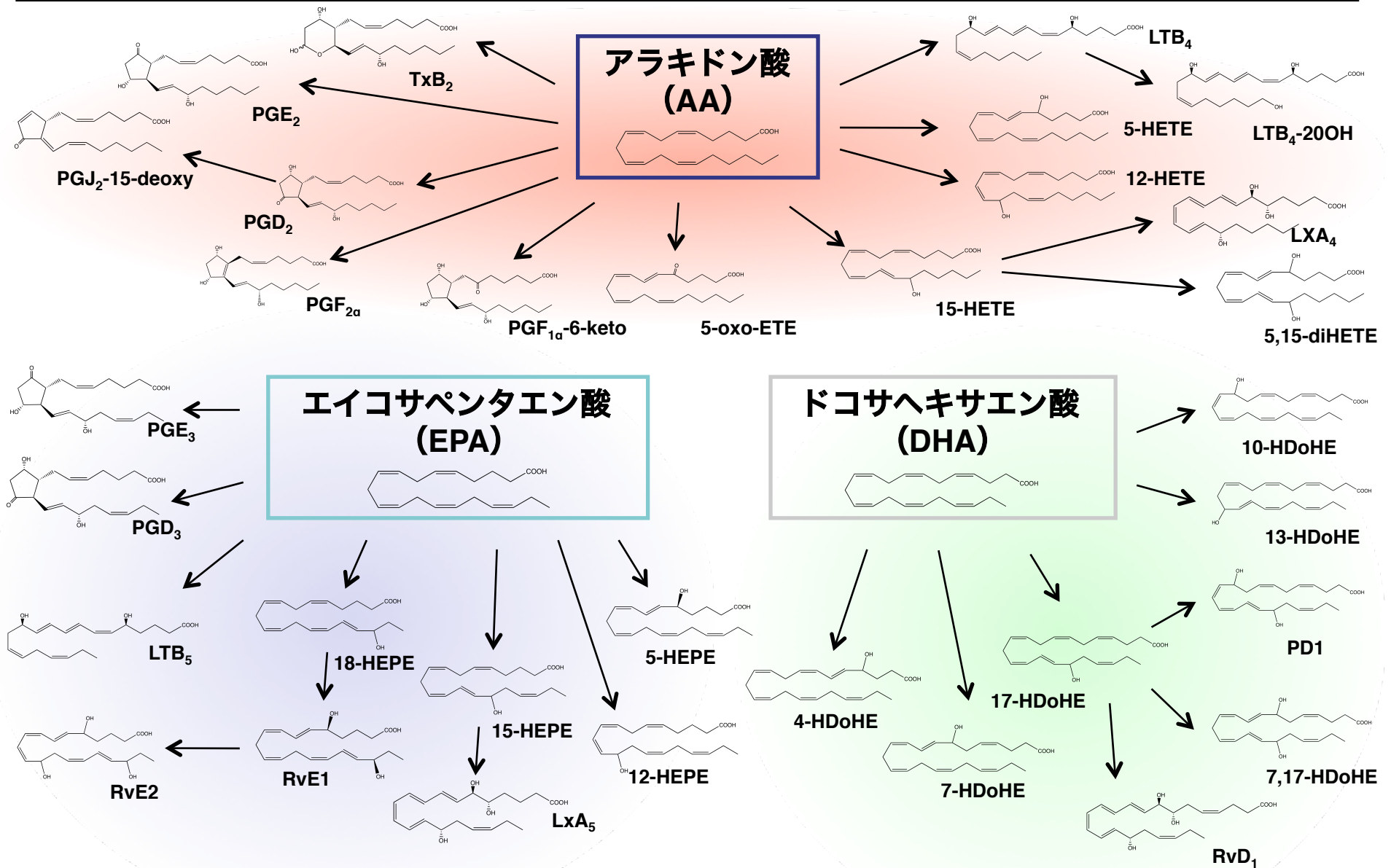
Eicosapentaenoic acid (EPA) → Resolvin Es
C20:5 ω 3



Docosahexaenoic acid (DHA) → Resolvin Ds
C22:6 ω 3
Protectins

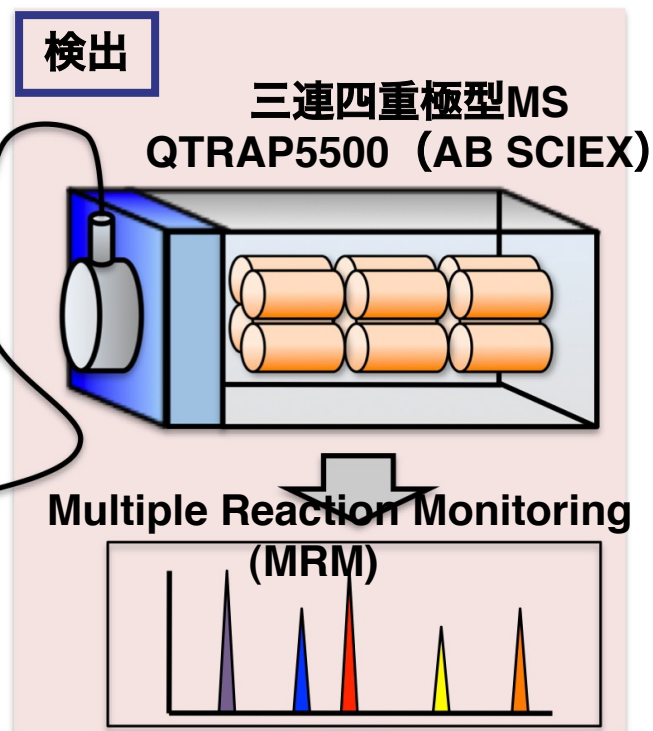
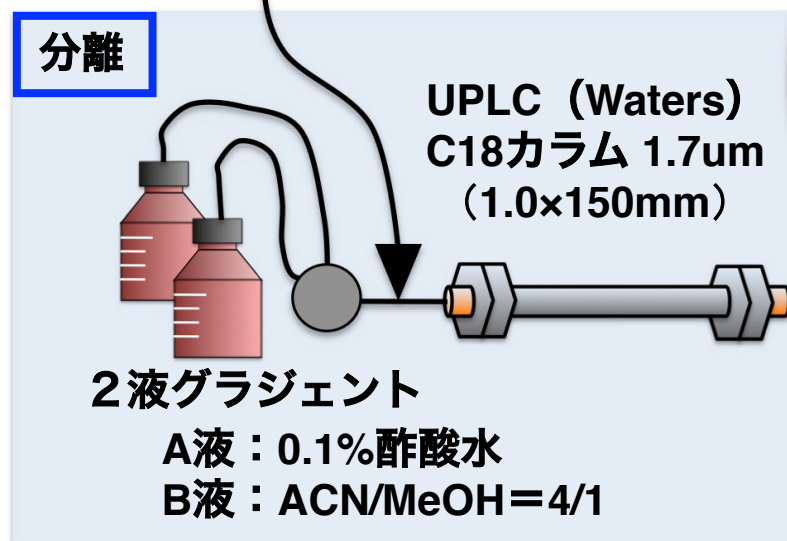
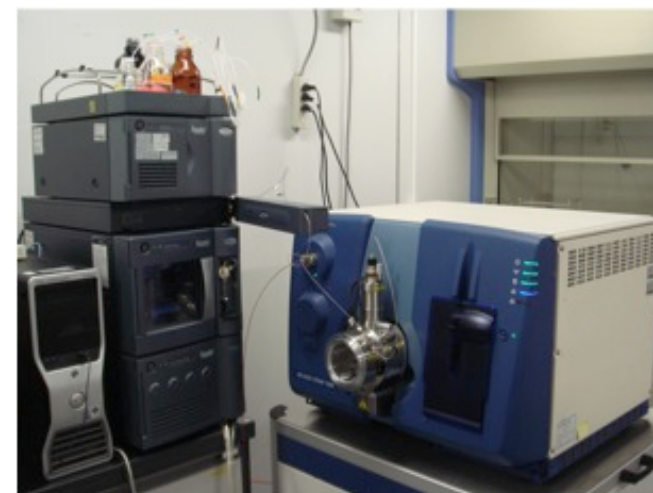
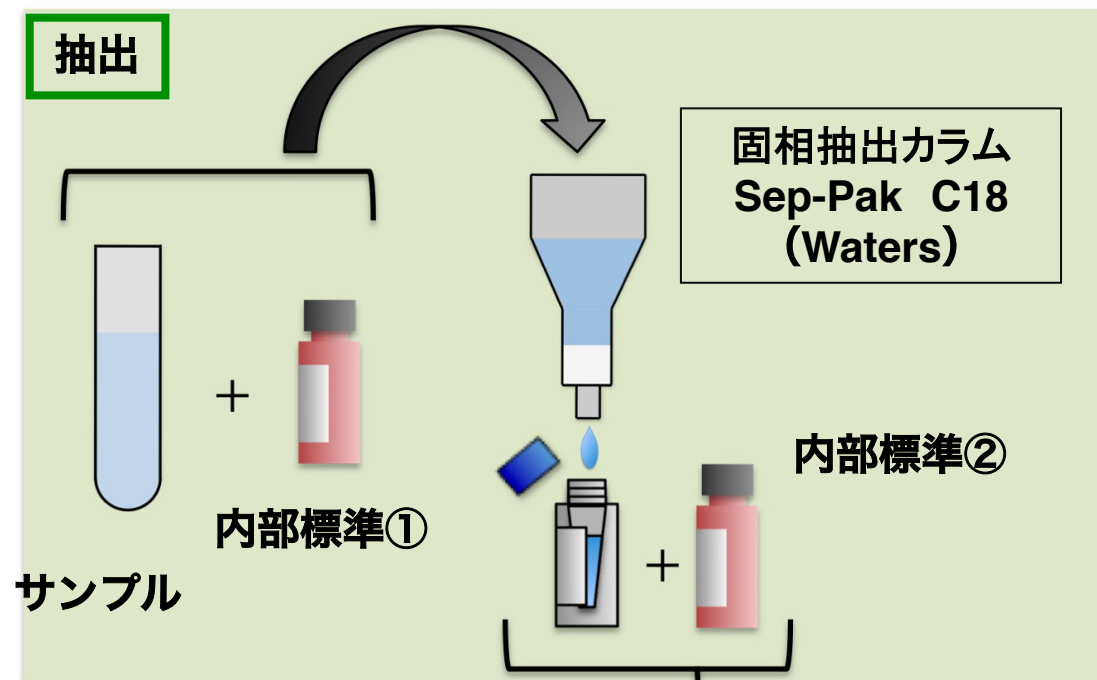


脂肪酸代謝マップ

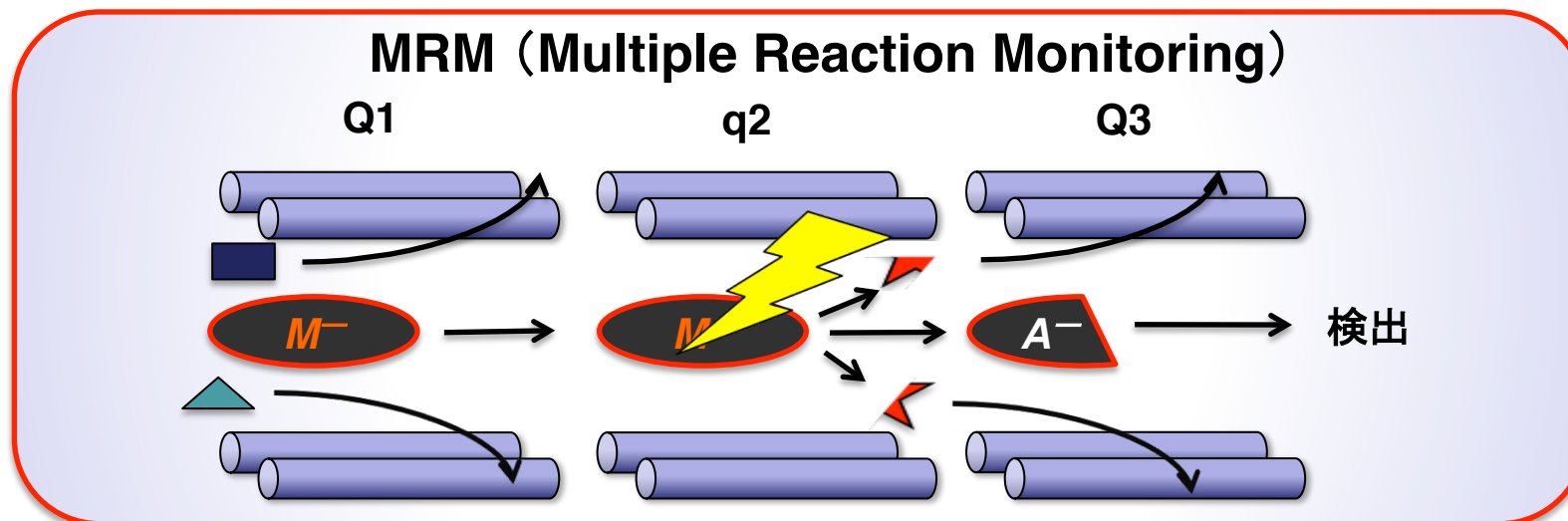


脂肪酸代謝フローの全体像を捉えることが重要

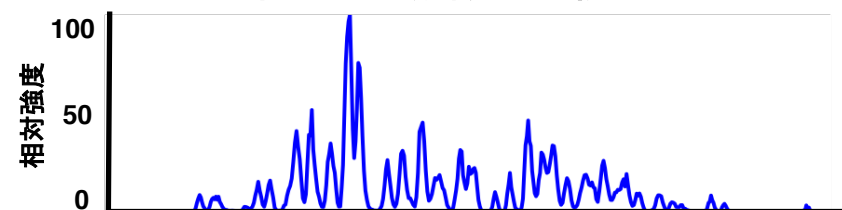
LC/MS/MSを用いた脂肪酸代謝物の包括的測定系



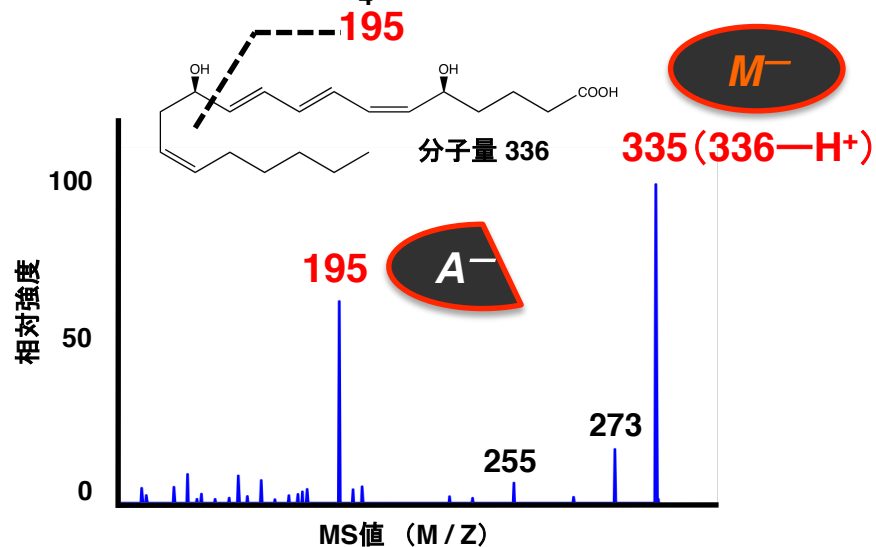
MRMシステムによる脂肪酸代謝物の測定



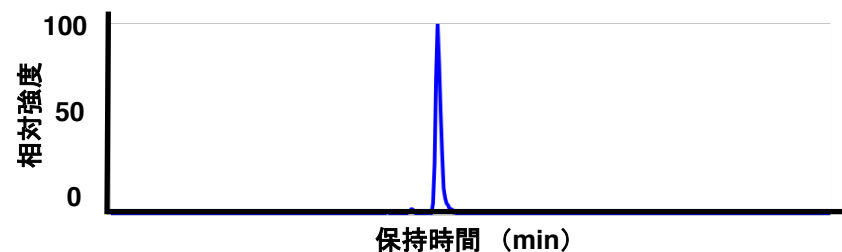
トータルイオンクロマトグラム (TIC)
様々な脂肪酸代謝物のmixture



ロイコトリエン B_4 のMS/MSスペクトル



MRM ロイコトリエン B_4 (335/195)



MRM総チャネル数 275チャネル

測定化合物総数 204種類

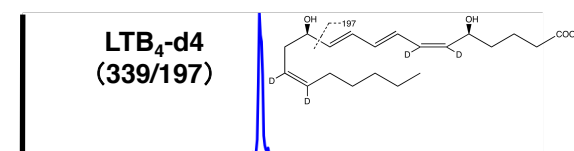
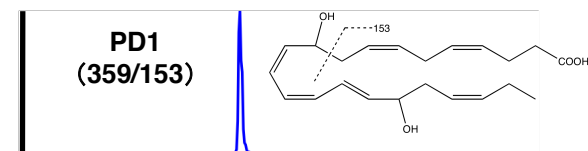
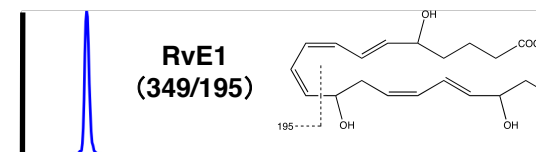
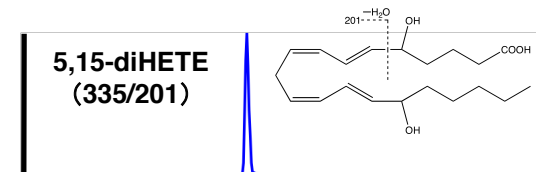
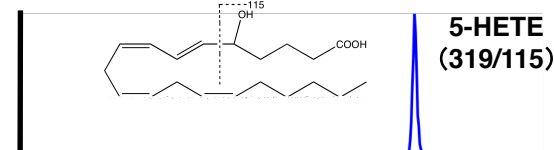
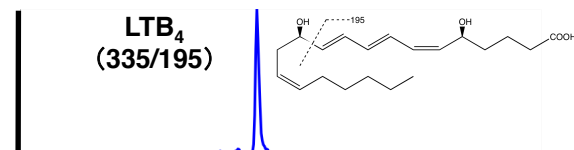
AA代謝物 47種類
EPA代謝物 55種類
DHA代謝物 56種類
DPA代謝物 45種類
その他 1種類
内部標準 6種類

酸化脂肪酸 MRMチャネル

Name	Q1	Q3	RT
001_PGE2	351.098	271.2	15.20
002_PGD2	350.933	271.1	15.50
003_PGD2-1	350.933	189.1	15.50
004_1S-keto-PGE2	349.019	235.0	15.70
005_PGE2-20OH	366.945	287.2	6.78
006_PGE2-20OH	366.945	189.1	6.78
007_LXA4	351.143	217.0	16.10
008_LXB4	351.120	221.1	15.30
009_PGF2a	353.137	193.2	14.90
010_PGF1a-6keto	369.187	163.0	12.60
011_TXB2	368.997	195.1	14.50
012_LTB4	335.111	195.0	19.10
013_LTB4-20OH	351.026	195.1	12.90
014_PGL2-15-deoxy	315.272	271.1	21.40
015_LTD4	495.085	177.1	17.40
016_LTE4	438.081	333.2	18.10
017_LTE4	438.081	351.1	18.10
018_5-HETE	319.010	114.9	25.00
019_5,6-EET	319.084	191.1	27.30
020_5,6-DHT	337.100	115.1	22.80
021_8-HETE	319.106	154.9	24.10
022_8,9-EET	319.102	154.9	26.80
023_8,9-DHT	337.100	127.1	22.30
024_9-HETE	319.079	123.1	24.61
025_11-HETE	319.005	167.0	23.62
026_12-HETE	319.107	179.1	24.00
027_11,12-EET	319.105	167.0	26.50
028_11,12-DHT	337.100	167.1	22.00
029_15-HETE	319.157	218.9	23.00
030_14,15-EET	319.095	219.2	25.60
031_14,15-DHT	337.100	207.1	21.10
032_16-HETE	319.111	233.1	22.20
033_16-HETE	319.111	189.0	22.20
034_17-HETE	319.106	247.1	21.90
035_17-HETE	319.106	203.2	21.90
036_18-HETE	319.102	261.1	21.80
037_18-HETE	319.102	217.1	21.80
038_19-HETE	319.098	275.1	21.40
039_19-HETE	319.098	231.1	21.40
040_20-HETE	319.106	289.1	21.10
041_AA+O	319.018	257.2	—
042_5,6-diHETE-1	335.234	219.0	21.40
043_5,6-diHETE-2	335.235	219.0	21.80
044_5,6-diHETE-3	335.236	219.0	22.20
045_5,6-diHETE-1	335.234	115.0	21.40
046_5,15-diHETE	335.111	201.1	18.70
047_8,15-diHETE	336.966	234.9	18.30
048_AA+2O	335.234	273.1	—
049_AA+3O	351.091	289.1	—
050_5-oxo-EET	317.041	203.2	26.40
051_12-oxo-EET	317.041	203.2	24.30
052_15-oxo-EET	317.040	113.2	23.50

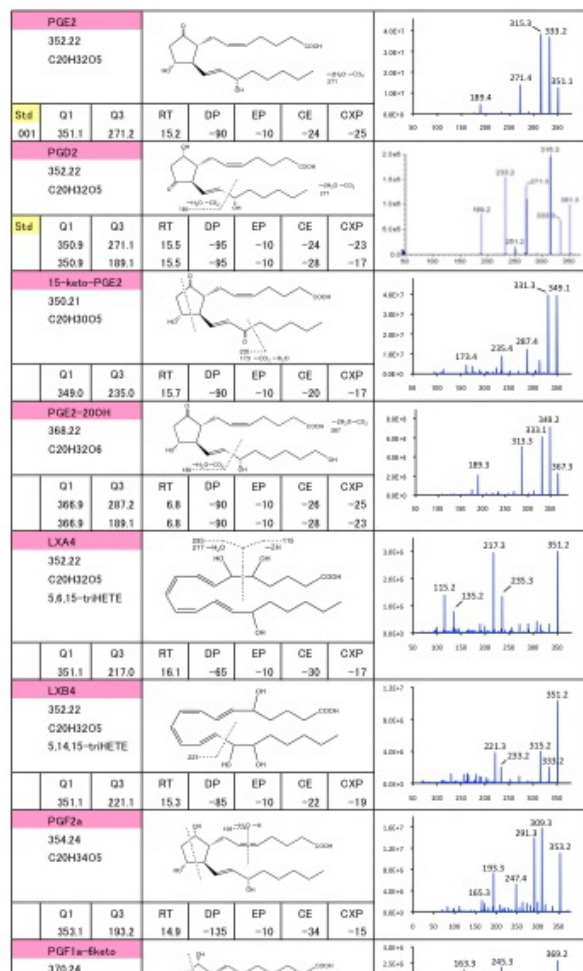
Name	Q1	Q3	RT
053_PGE3	349.013	233.2	14.10
054_PGE3	349.013	189.2	14.10
055_PGD3	349.112	188.9	14.40
056_LXA5	349.032	115.1	15.00
057_LXA5	349.010	215.0	15.00
058_LXB5	349.100	221.1	14.00
059_PGF3a	351.100	191.2	14.90
060_PGF1a-delta17-6-keto	367.100	163.0	12.60
061_TXB3	367.100	195.1	14.20
062_PGL3-15-deoxy	313.100	269.1	21.40
063_LTD5	493.100	177.1	17.40
064_LTE5	436.100	349.1	18.10
065_5-HEPE	317.011	114.9	22.40
066_8-HEPE	316.988	155.0	21.80
067_9-HEPE	316.962	149.1	22.10
068_9-HEPE	316.962	123.1	22.10
069_11-HEPE	317.096	167.0	21.50
070_11-HEPE	317.096	195.0	21.50
071_12-HEPE	317.127	179.0	21.80
072_12-HEPE	317.052	135.0	21.80
073_15-HEPE	316.961	219.1	21.40
074_15-HEPE	316.961	175.1	21.40
075_14,15-EpETE	316.997	207.2	23.70
076_14,15-diHETE	335.258	207.1	18.00
077_18-HEPE	316.992	209.1	21.00
078_18-HEPE	317.139	215.2	21.00
079_17,18-EpETE	317.090	259.1	23.10
080_17,18-diHETE	335.258	247.2	18.70
081_17,18-diHETE	335.258	203.2	18.70
082_20-HEPE	317.046	287.0	20.72
083_EPA+O	317.044	255.2	—
084_LTB5	332.946	185.1	17.70
085_RvE2	333.108	115.1	17.09
086_6,18-diHEPE	333.100	205.1	18.10
087_8,18-diHEPE	332.962	155.0	16.70
088_8,18-diHEPE	332.962	159.1	16.70
089_9,18-diHEPE	333.100	139.1	18.20
090_11,18-diHEPE-1	333.100	167.1	16.83
091_11,18-diHEPE-2	333.100	167.1	17.11
092_11,18-diHEPE-3	333.100	167.1	18.14
093_11,18-diHEPE-1	333.098	123.0	16.83
094_11,18-diHEPE-2	333.098	123.0	17.11
095_11,18-diHEPE-3	333.098	123.0	18.14
096_12,18-diHEPE	332.940	179.1	16.86
097_12,18-diHEPE	332.940	163.1	16.86
098_15,18-diHEPE	333.100	219.1	17.70
099_17,18-diHEPE-1	333.104	201.0	18.10
100_17,18-diHEPE-2	333.105	201.0	18.60
101_17,18-diHEPE-1	333.104	245.1	18.10
102_17,18-diHEPE-2	333.105	245.1	18.60
103_17,18-diHEPE-1	333.104	213.1	18.10
104_17,18-diHEPE-2	333.105	213.1	18.60

Name	Q1	Q3	RT
139_4-HDoHE	343.102	100.9	25.40
140_7-HDoHE	343.063	141.0	24.20
141_8-HDoHE	343.068	189.2	24.40
142_10-HDoHE	343.068	153.1	23.50
143_11-HDoHE	342.981	121.1	23.90
144_11-HDoHE	342.981	149.0	23.90
145_13-HDoHE	343.136	193.0	23.20
146_14-HDoHE	342.998	205.1	23.40
147_14-HDoHE	342.998	161.1	23.40
148_16-HDoHE	343.091	233.2	23.00
149_17-HDoHE	342.977	245.1	23.10
150_17-HDoHE	342.977	201.1	23.10
151_16,17-EpDPE	342.954	233.1	25.80
152_20-HDoHE	342.960	241.1	22.60
153_19,20-EpDPE	343.106	241.1	25.10
154_19,20-diHDoPE	361.044	229.2	20.00
155_21-HDoHE	343.077	255.1	22.10
156_22-HDoHE	342.960	313.1	22.50
157_DHA+O	343.094	281.2	—
158_4,14-diHDoHE	359.075	101.0	19.80
159_4,14-diHDoHE	359.107	187.1	19.80
160_7,14-diHDoHE	359.118	112.9	19.19
161_7,14-diHDoHE	359.118	177.0	19.19
162_7,17-diHDoHE	359.103	199.0	18.70
163_10,17-diHDoHE	359.107	153.0	18.60
164_4,20-diHDoHE	359.098	101.0	19.00
165_5,20-diHDoHE	359.100	113.1	19.10
166_7,20-diHDoHE	359.100	217.1	19.10
167_8,20-diHDoHE	359.100	125.2	19.20
168_10,20-diHDoHE	359.098	159.0	18.08
169_10,20-diHDoHE	359.035	239.3	18.08
170_11,20-diHDoHE	359.100	165.1	19.10
171_13,20-diHDoHE-1	358.913	193.1	17.90
172_13,20-diHDoHE-2	358.913	193.1	18.18
173_13,20-diHDoHE-3	358.913	193.1	18.51
174_14,20-diHDoHE	359.035	233.1	18.27
175_14,20-diHDoHE	359.032	205.0	18.27
176_19,20-diHDoHE-1	358.983	227.2	19.51
177_19,20-diHDoHE-2	358.983	227.2	19.97
178_20,22-diHDoHE	359.100	241.1	19.00
179_14,21-diHDoHE	359.047	253.1	17.60
180_4Hy-19,20-EpDPE	359.080	100.9	20.27
181_4Hy-19,20-EpDPE	359.080	245.1	20.27
182_7Hy-19,20-EpDPE	359.091	141.0	19.60
183_8Hy-19,20-EpDPE	359.128	125.2	19.66
184_10Hy-19,20-EpDPE	359.106	153.0	19.36
185_11Hy-19,20-EpDPE	359.100	165.2	19.42
186_11Hy-19,20-EpDPE	359.100	193.0	19.42
187_13Hy-19,20-EpDPE	359.092	193.0	19.34
188_14Hy-19,20-EpDPE	359.104	205.0	19.42
189_14Hy-19,20-EpDPE	359.104	161.2	19.42
190_16Hy-19,20-EpDPE	359.108	233.2	18.38

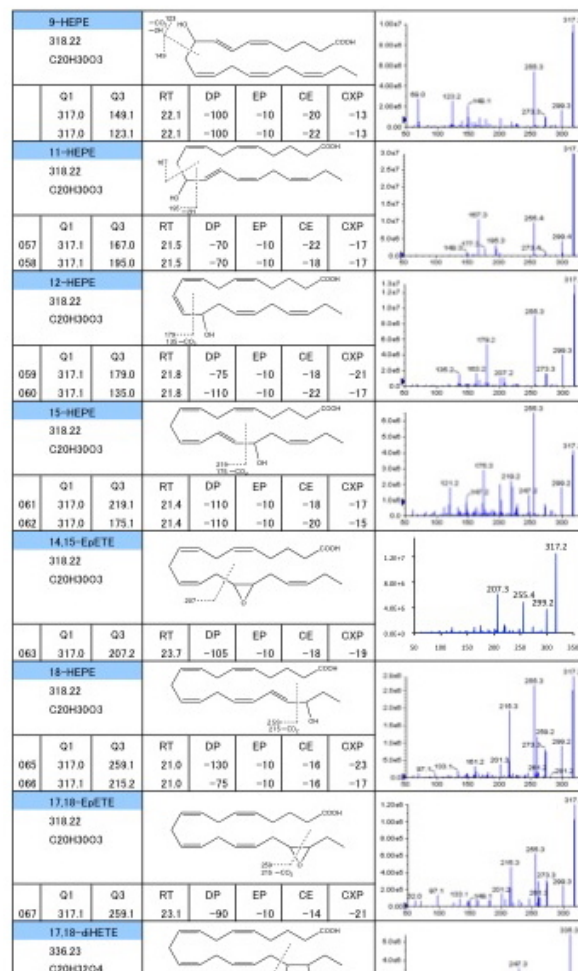


LC-MS/MS data and MRM conditions of PUFA metabolites

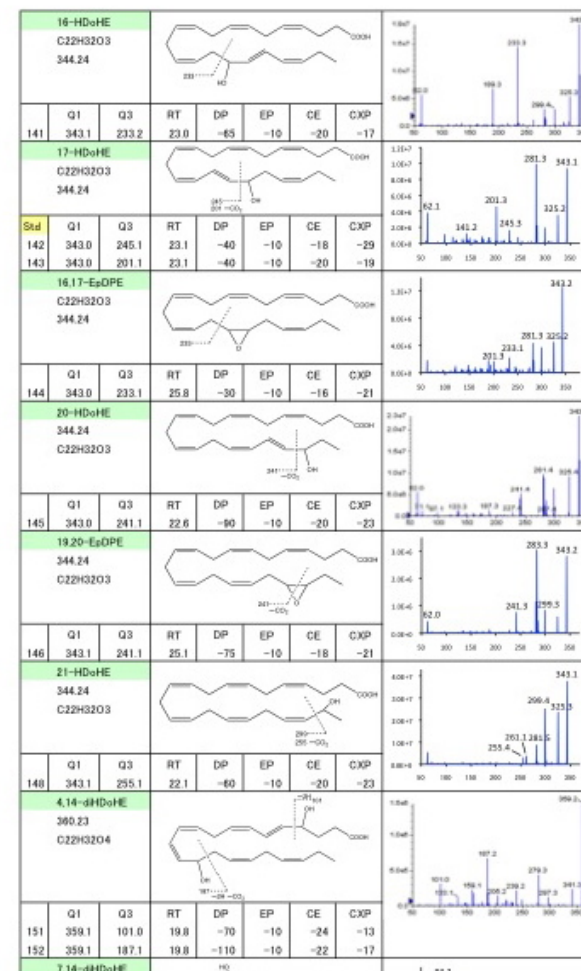
ArA-derived



EPA-derived



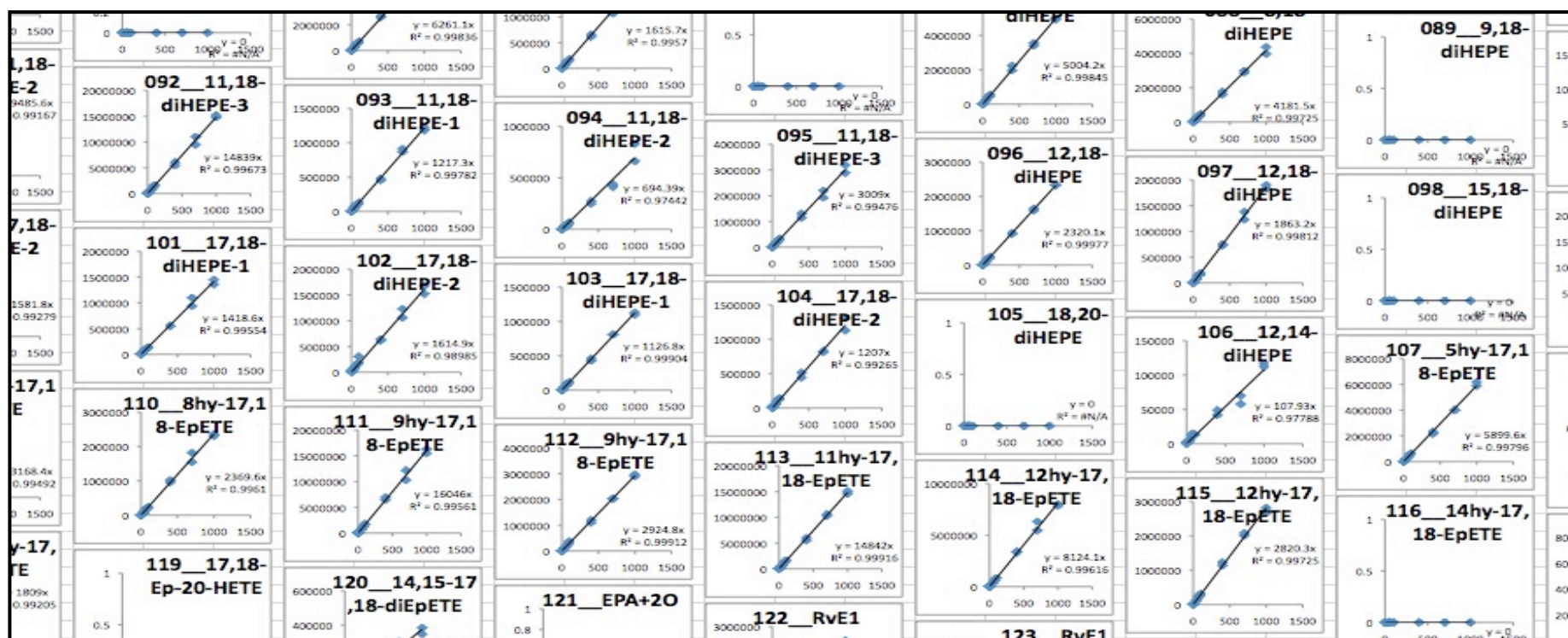
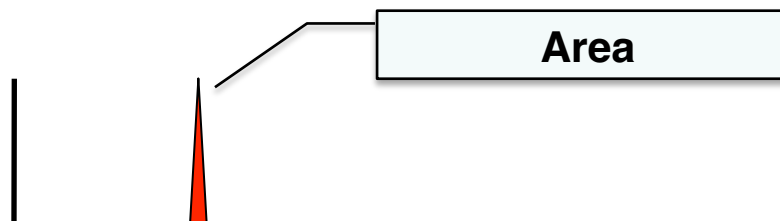
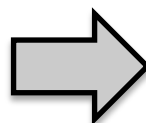
DHA-derived



Linear standard curves for the quantitative analyses

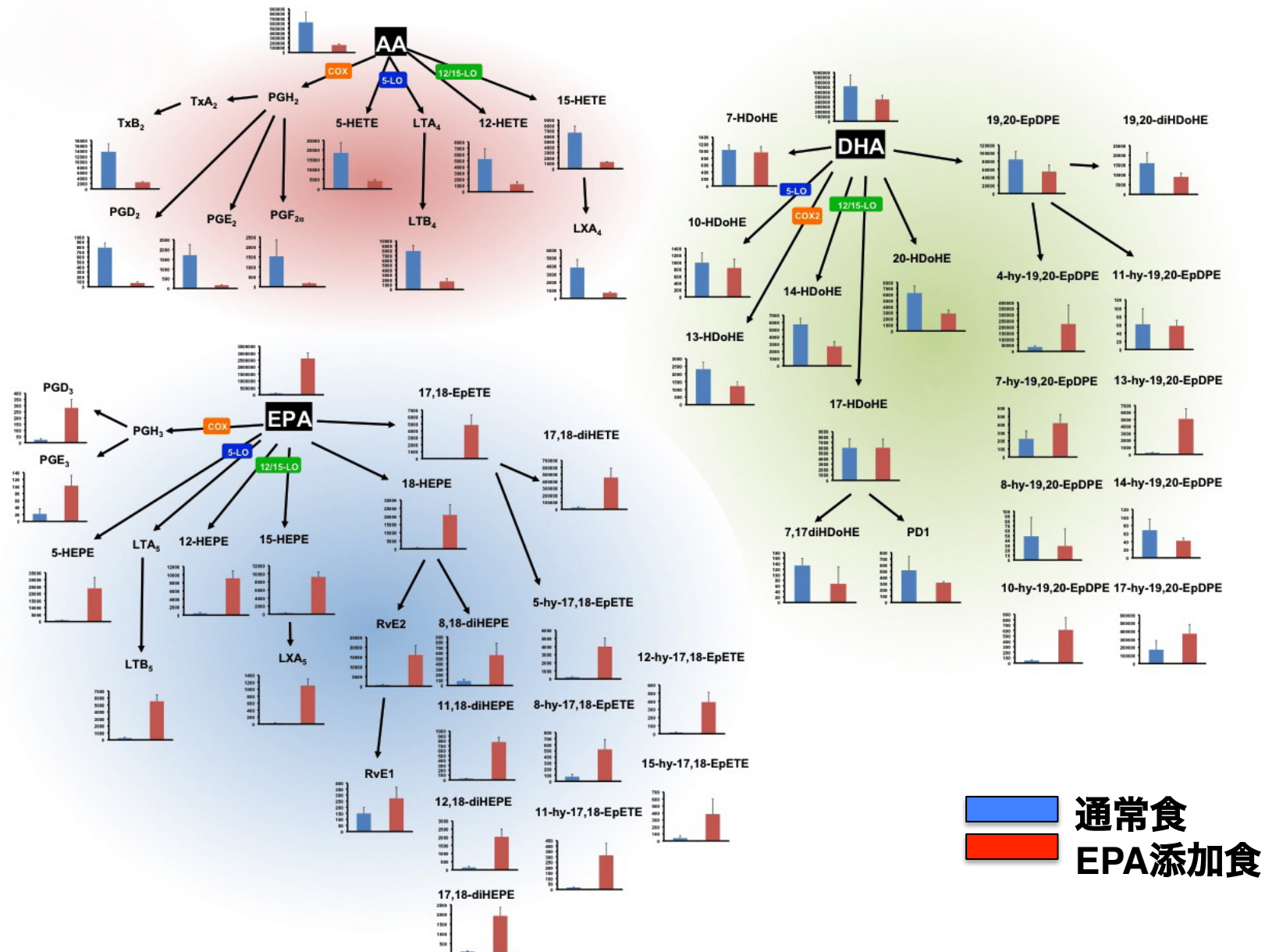


standard X
on column



Linear range: 1pg~1000pg on column

脂質メタボローム解析の具体例

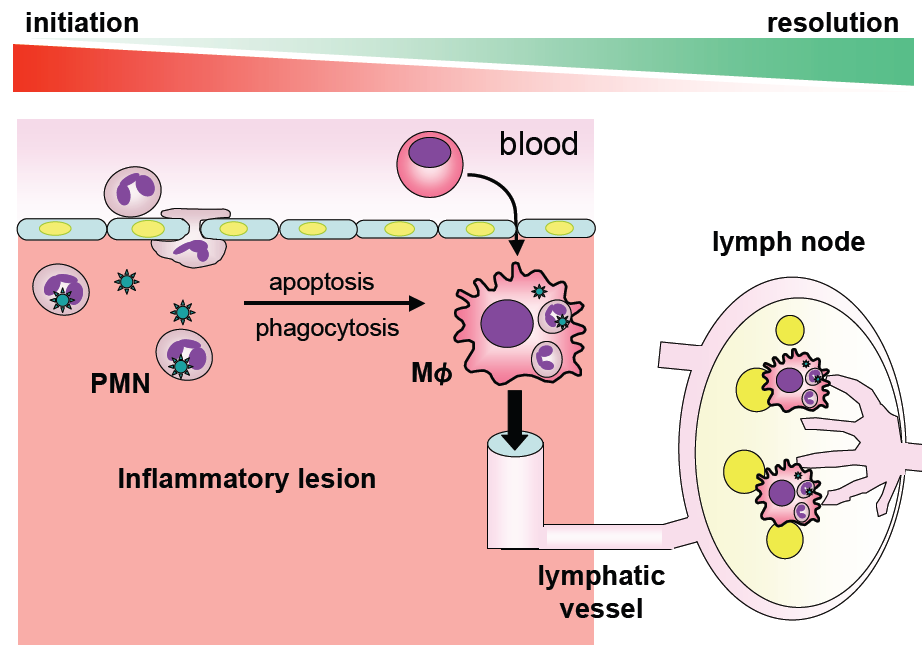
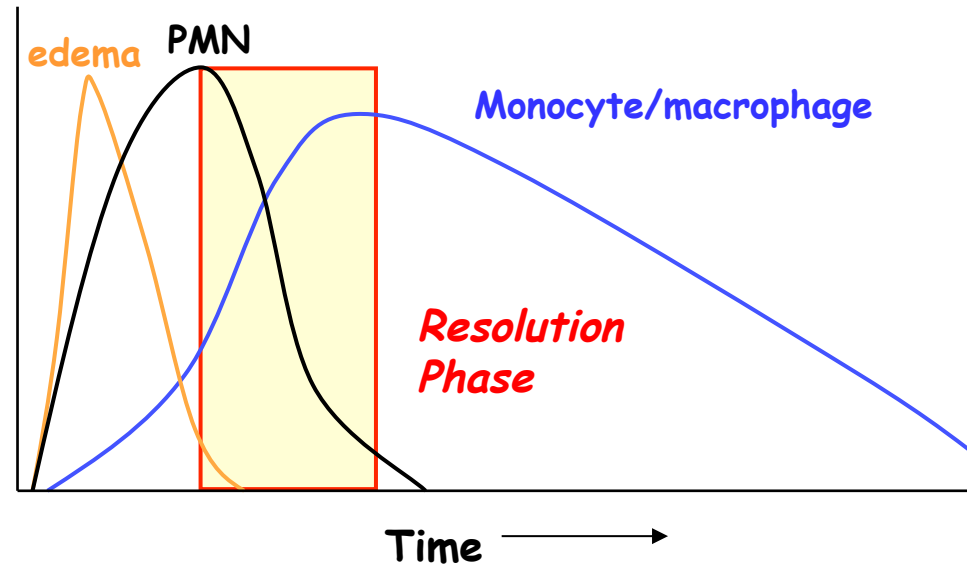


メタボロミクスを用いた

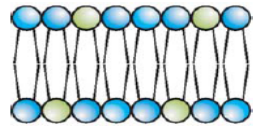
1. 炎症の収束機構の解析

2. n-3系脂肪酸由来の抗炎症性 代謝物の新規探索

炎症反応の開始(initiation)と収束(resolution)

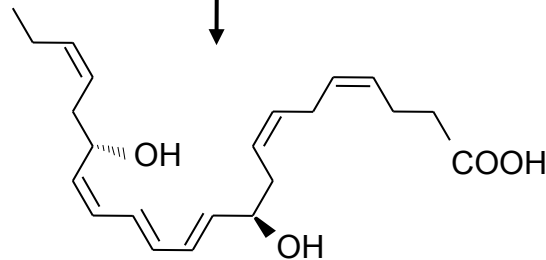


DHA由来の抗炎症性メディエーター: プロテクチンD1 (PD1)



↓
ドコサヘキサエン酸
(DHA; C22:6 n-3)

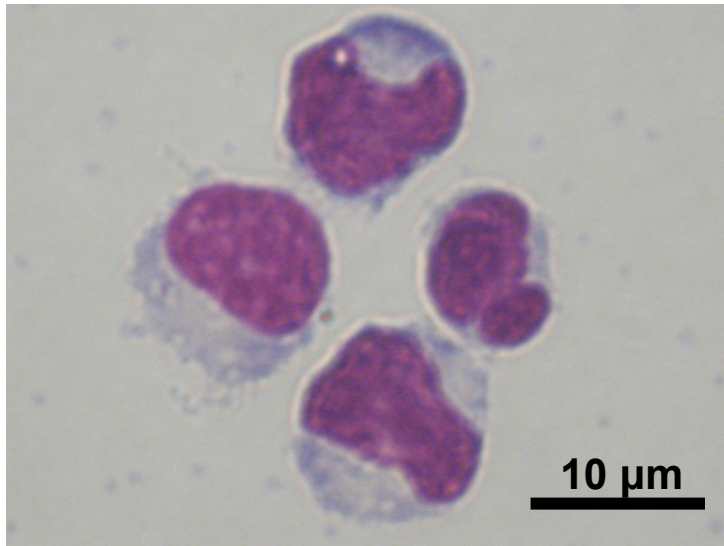
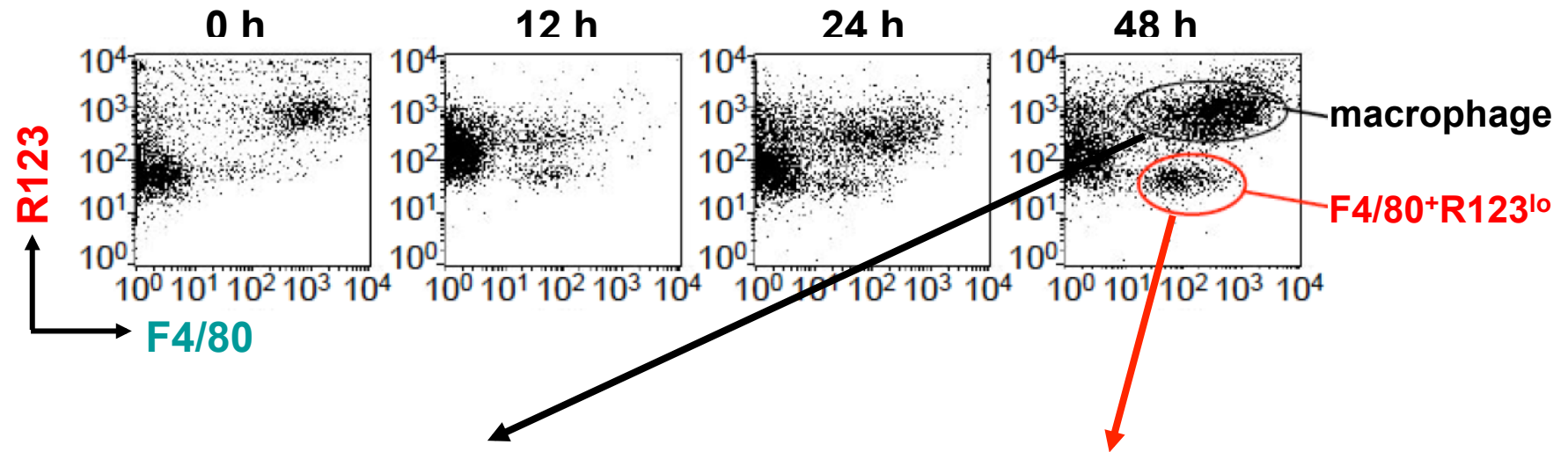
↓
12/15-LOX



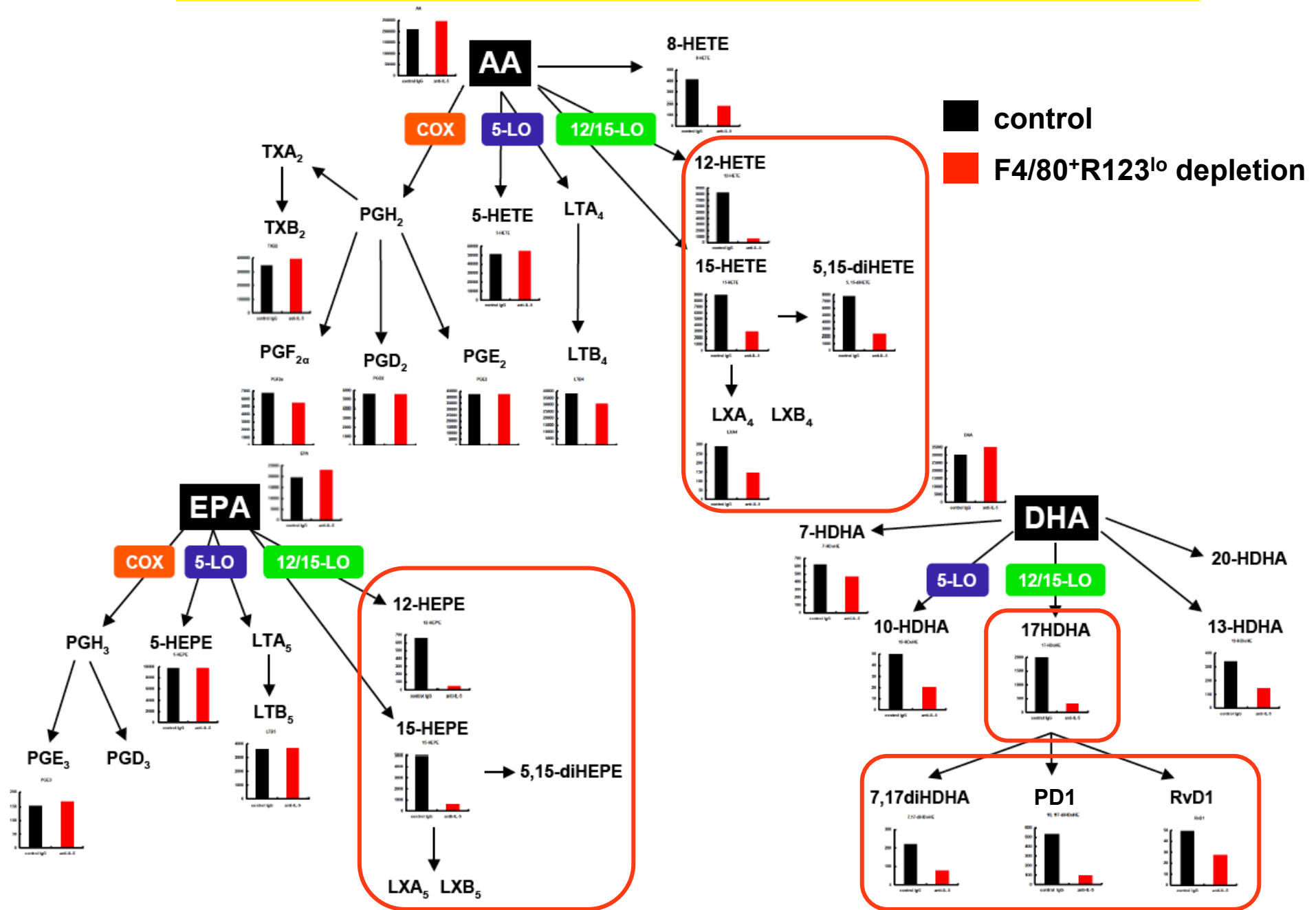
Protectin D1
(10,17S-diHDHA)

- ・急性炎症(好中球浸潤)の抑制
- ・マクロファージの貪食能促進
- ・神経細胞死の抑制
- ・上皮組織(角膜)の修復促進
- ・インスリン抵抗性の改善

炎症収束期に出現するF4/80⁺R123^{lo}細胞

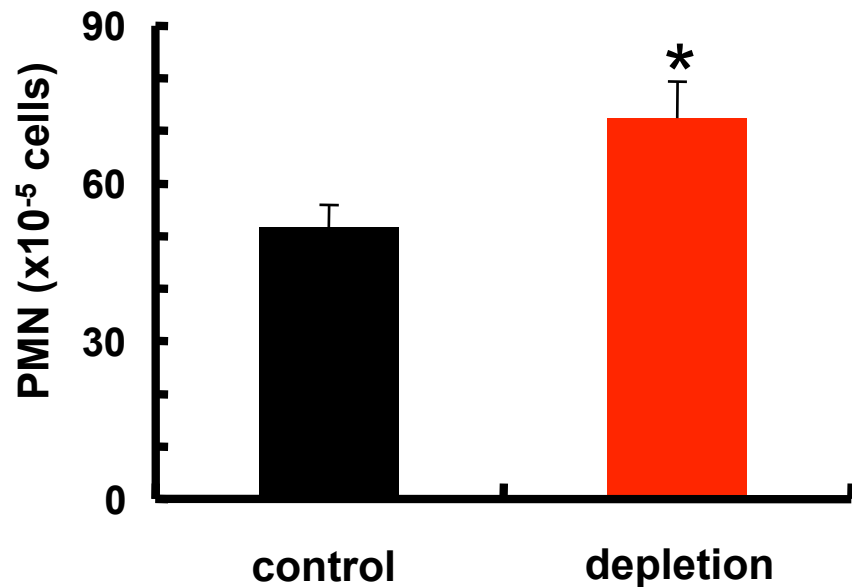


F4/80⁺R123^{lo} depletionによる脂肪酸代謝物の変動



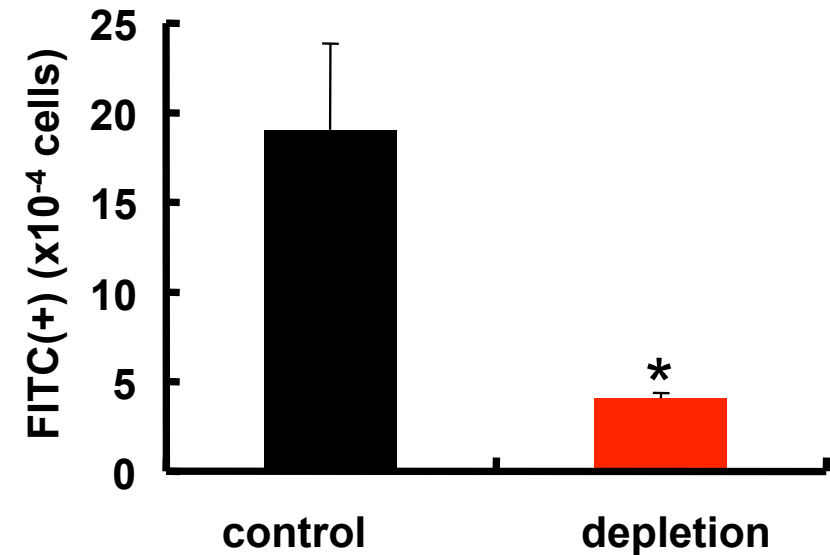
F4/80⁺R123^{lo} depletionにより炎症収束が遅延する

PMN numbers in exudates



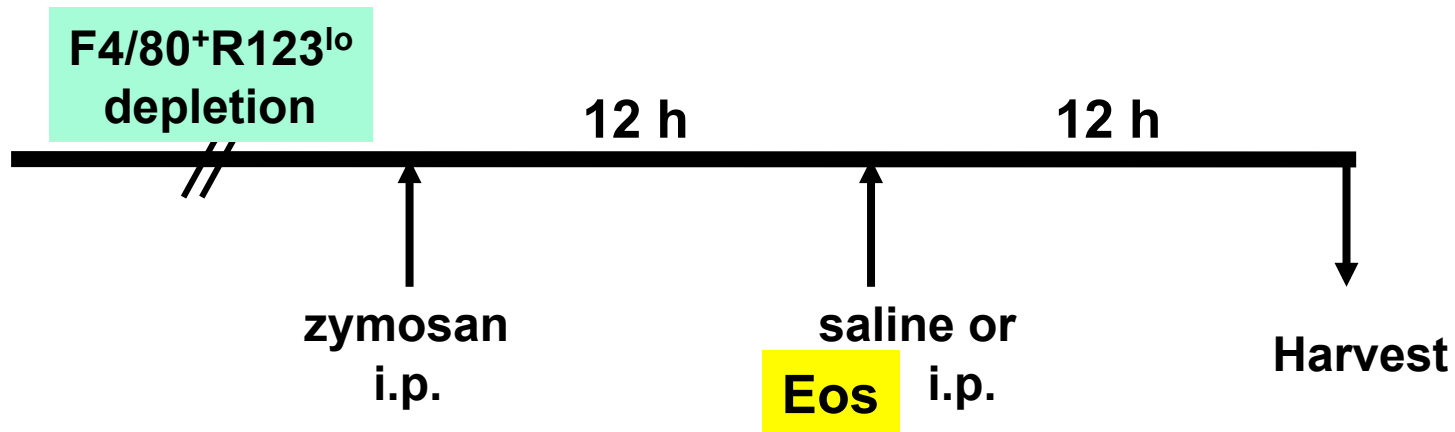
*p < 0.05, n = 4-6, mean ± s.e.m

Leukocytes carrying zymosan in parathymic lymph node



*p < 0.05 n = 5-6, mean ± s.e.m

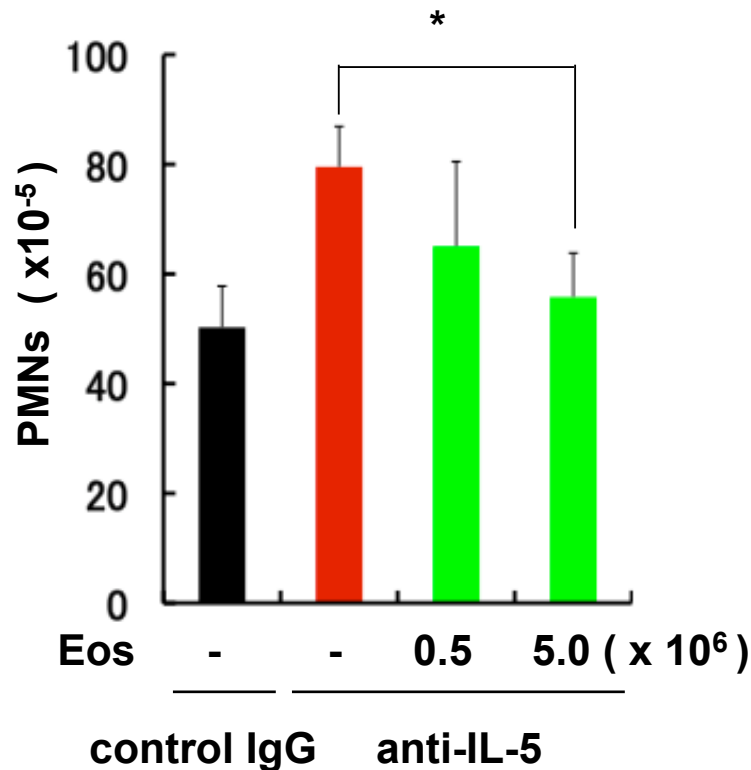
**F4/80⁺R123^{lo} depletionによる炎症収束の遅延は
好酸球を外から補うことで再び回復するか？**



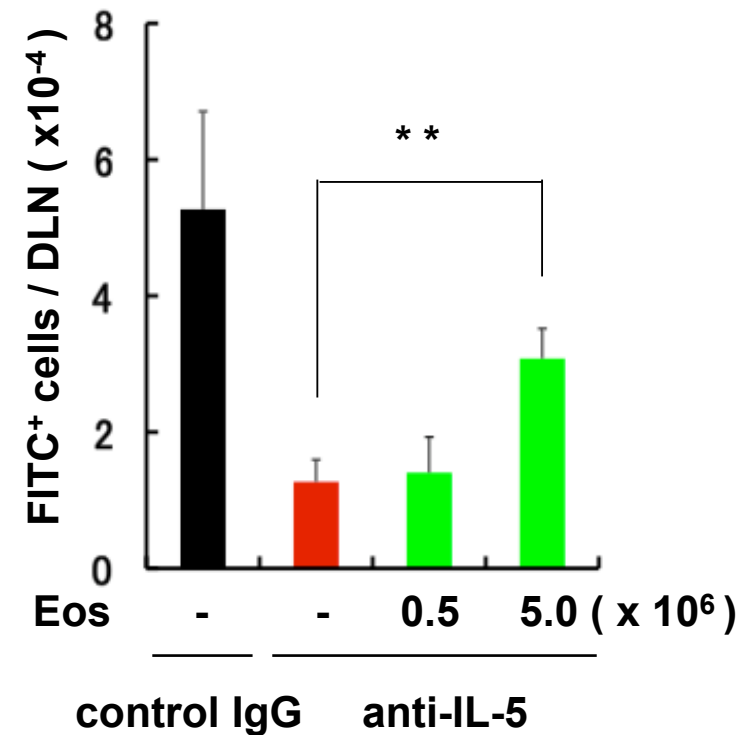
Eos (好酸球) : IL-5トランスジェニックマウスより単離

F4/80⁺R123^{lo} depletionによる炎症収束の遅延は 好酸球を外から補なうことで回復する

PMN numbers in exudates

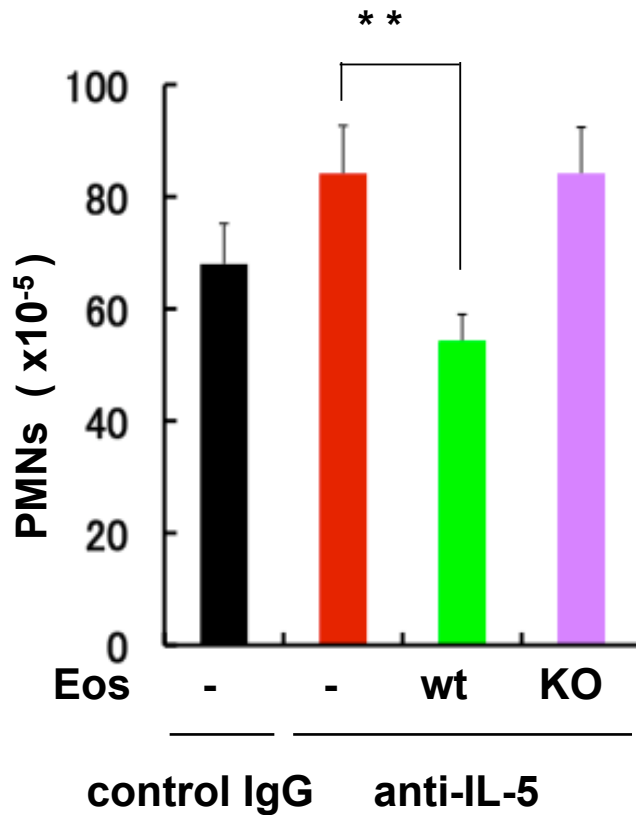


Leukocytes carrying zymosan in parathymic lymph node

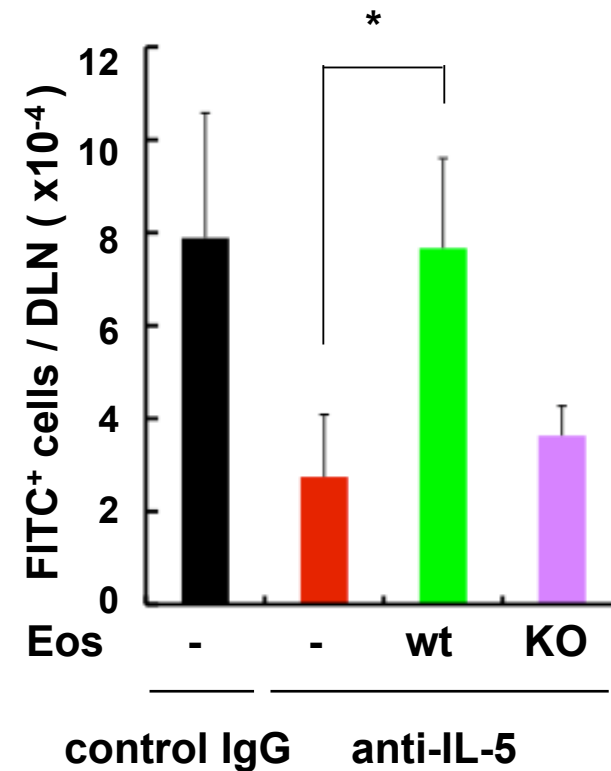


好酸球の炎症収束を促進する機能は 12/15-LOXを欠損した好酸球では認められない

PMN numbers in exudates



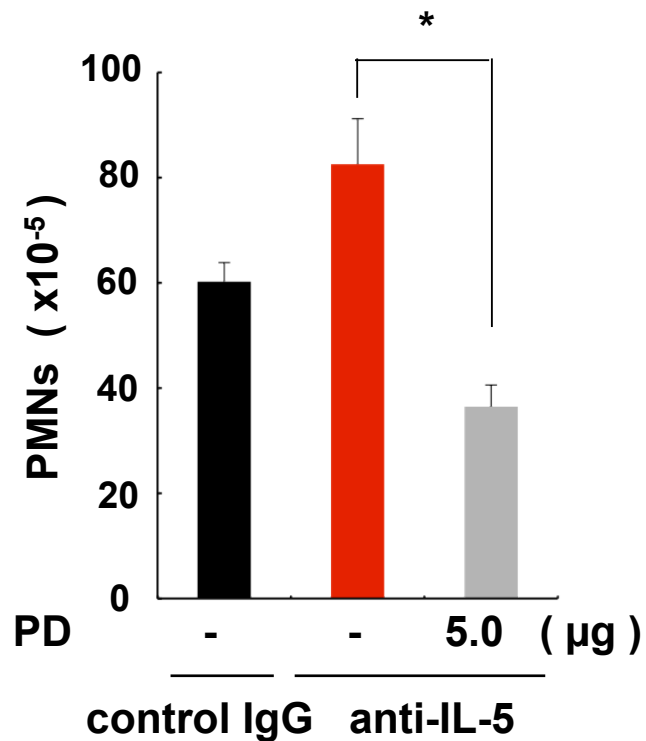
Leukocytes carrying zymosan
in parathymic lymph node



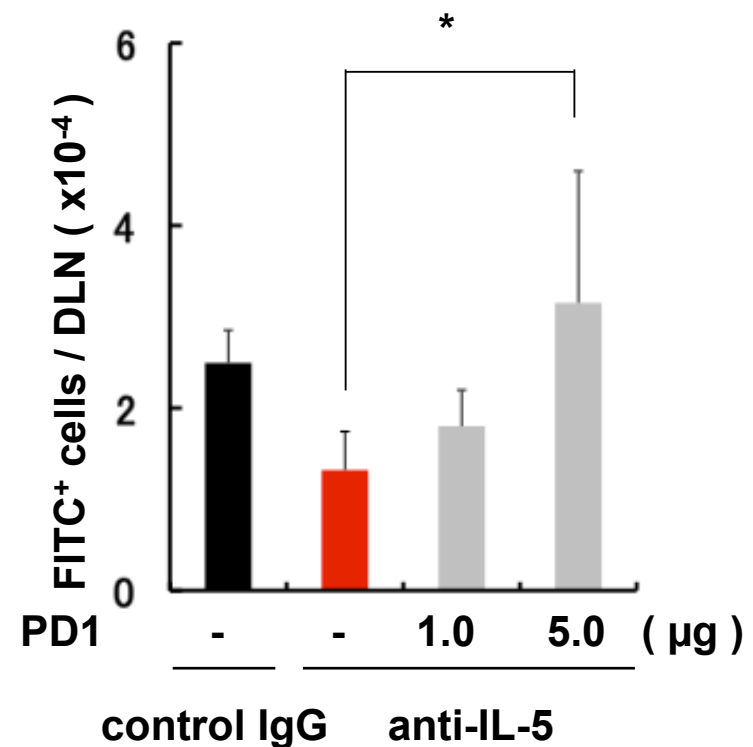
wt: wild type eosinophils
KO: 12/15-LOX-deficient eosinophils

F4/80⁺R123^{lo} depletionによる炎症収束の遅延は 12/15-LOX産物Protectin D1の投与で回復する

PMN numbers in exudates



Leukocytes carrying zymosan in parathymic lymph node



PD1: Protectin D1 *i.p.* injection

炎症収束における好酸球の役割

