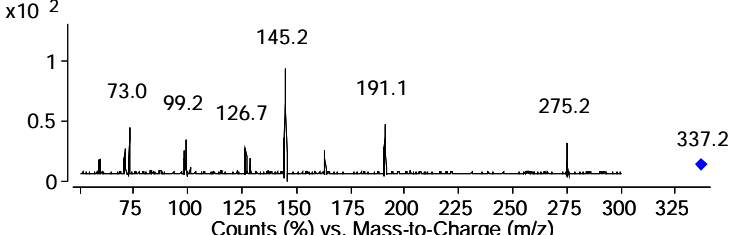
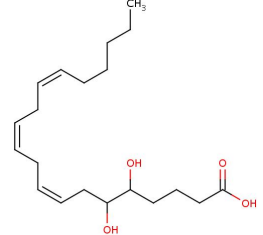
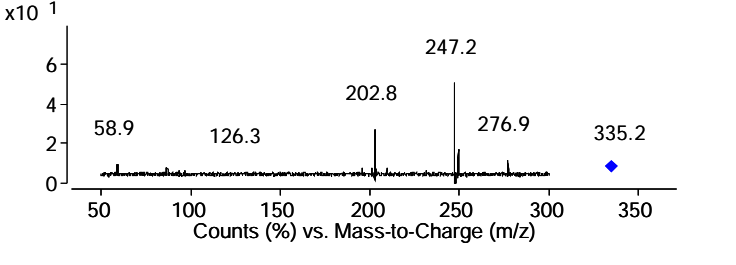
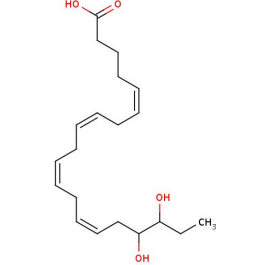
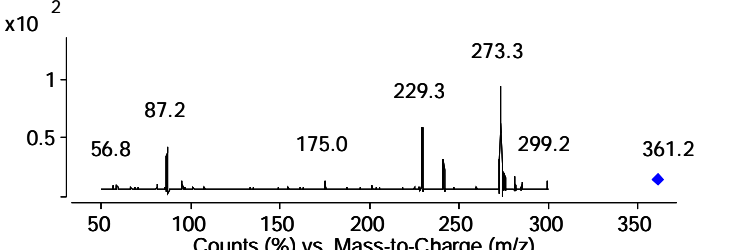
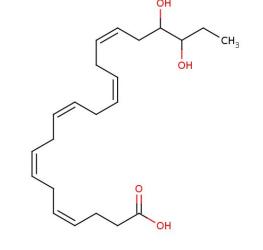


FIGURE S1 (supplemental): MS/MS spectra and MRM transition information of diols detected in piglet plasma

MS/MS Spectrum ^a	Compound and transition	Structure ^b
Precursor Linoleic Acid		
	<p>12,13-DiHOME m/z 313.1 → 183.2</p>	
	<p>9,10-DiHOME m/z 313.1 → 201.1</p>	
Precursor Arachidonic Acid		
	<p>14,15-DiHETrE m/z 337.2 → 207.1</p>	
	<p>11,12-DiHETrE m/z 337.2 → 167.1</p>	
	<p>8,9-DiHETrE m/z 337.2 → 127.0</p>	

 <p>Counts (%) vs. Mass-to-Charge (m/z)</p>	<p>5,6-DiHETrE m/z 337.2 → 145.2</p>	
<p>Precursor Eicosapentaenoic Acid</p>		
 <p>Counts (%) vs. Mass-to-Charge (m/z)</p>	<p>17,18-DiHETE m/z 335.2 → 247.2</p>	
<p>Precursor Docosahexaenoic Acid</p>		
 <p>Counts (%) vs. Mass-to-Charge (m/z)</p>	<p>19,20-DiHDPa m/z 361.2 → 273.3</p>	

^a MS/MS fragmentation was performed in neg. ion mode on a triple quadrupole mass spectrometer

(Agilent 6460), precursor ion = [M-H]⁻ (negative ion mode)

^b Structures were extracted from HMDB (71)

