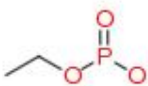


# Report

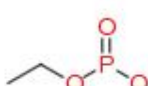
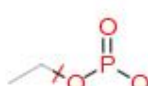
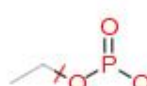
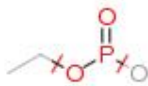
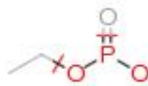
Input:

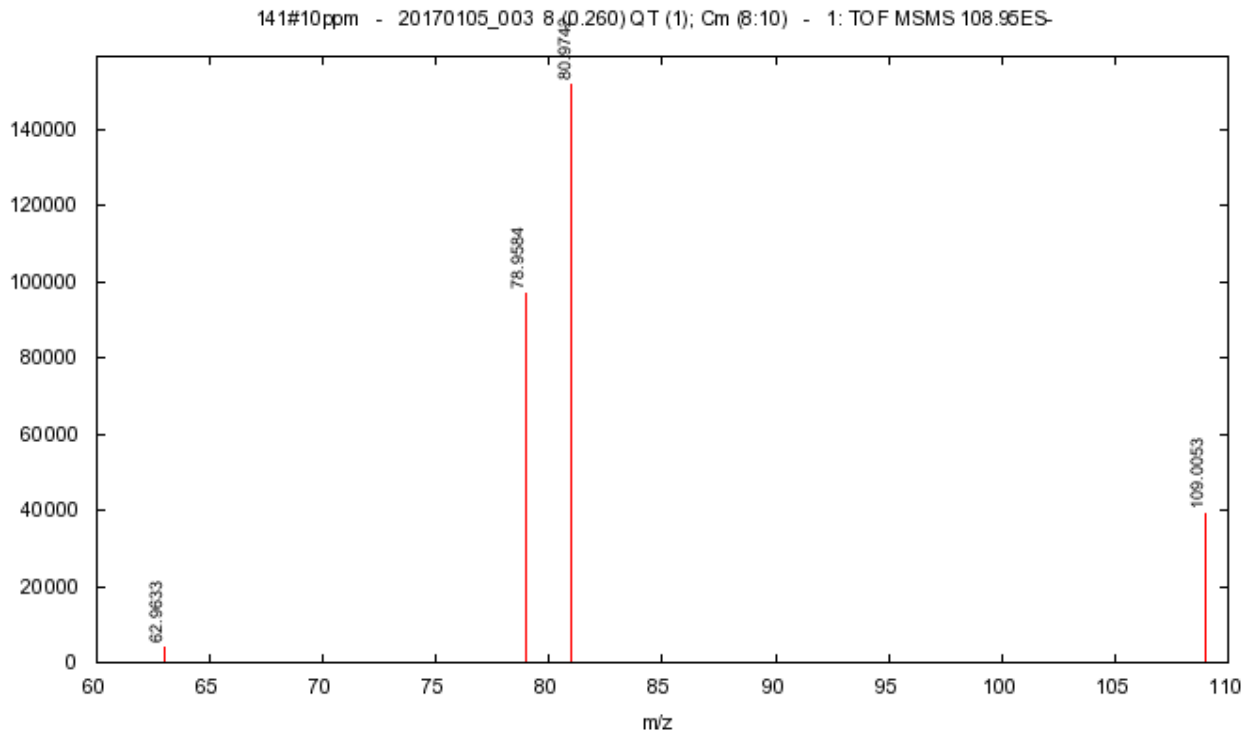
	<b>ID (job)</b>	91
	<b>Mass (Da)</b>	109.0055
	<b>Formula</b>	C <sub>2</sub> H <sub>6</sub> O <sub>3</sub> P
	<b>DBE</b>	1.5

Experiment:

<b>Product ion(s) (Da)</b>	109.0053 62.9633 78.9584 80.9742 +/- 0.01 in negative mode, structure filter 1
<b>DBE</b>	-10 to 50
<b>Electron count</b>	both
<b>Maximum H deficit</b>	6
<b>Fragment number of bonds</b>	4
<b>Scoring</b>	aromatic: 6, multiple: 4, ring: 2, phenyl: 8, other: 1 H-deficit: 0, hetero modifier: 0.5, max score: 16
<b>Order:</b>	mass
<b>Plot:</b>	show <input checked="" type="radio"/> hide <input type="radio"/>
<b>Files:</b>	<a href="#">CSV</a>

Results:

<p><b>109.0053</b>      <math>\bar{-}</math> (+2H)</p>  <p>109.0055 (-0.2.mDa) C<sub>2</sub>H<sub>6</sub>O<sub>3</sub>P (-none)</p>	<p><b>80.9742</b>      <math>\bar{-}</math> (+3H)</p>  <p>80.9742 (+0.0.mDa) (S:0.5, B:1) H<sub>2</sub>O<sub>3</sub>P (-C<sub>2</sub>H<sub>5</sub>)</p>	<p><b>78.9584</b>      <math>\bar{-}</math> (+1H)</p>  <p>78.9585 (-0.1.mDa) (S:0.5, B:1) O<sub>3</sub>P (-C<sub>2</sub>H<sub>5</sub>)</p>
<p><b>62.9633</b>      <math>\bar{-}</math> (+2H)</p>  <p>62.9636 (-0.3.mDa) (S:1.5, B:2) O<sub>2</sub>P (-C<sub>2</sub>H<sub>5</sub>O)</p>	<p><b>62.9633</b>      <math>\bar{-}</math> (+1H)</p>  <p>62.9636 (-0.3.mDa) (S:4.5, B:2) O<sub>2</sub>P (-C<sub>2</sub>H<sub>5</sub>O)</p>	



You are running version 3.0.w.013

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