

General MS Services

Itemized Fees (\$)

<p>ESI/MS (Electrospray Ionization/ Mass Spectrometry)</p> <p>Positive-ion ESI or negative-ion ESI; sample introduction by infusion</p>	<p>40.00 per sample or acquisition 15.00 per sample for Ziptip™ purification as required</p>
<p>ESI/MS/MS (Electrospray Ionization/ Mass Spectrometry/ Mass Spectrometry) Ion-trap, triple quad, or Q-ToF instruments</p> <p>For structure elucidation, confirmation of structure, and determining fragmentation pathways</p> <p>Samples introduced by infusion; Positive-ion or negative-ion ESI; MS/MS with CID; Daughter-ion, Parent-ion, or Neutral-loss analyses available</p>	<p>50.00 per sample 20.00 for each additional parent ion analyzed by MS/MS in the same sample 15.00 per sample for Ziptip™ purification as required</p>
<p>Nucleic Acids</p> <p>DNA oligomers, RNA oligomers, PCR products, other nucleic acids</p> <p>Samples introduced by infusion; Negative-ion ESI/MS analysis; MaxEnt™ deconvolution of data to generate a neutral molecular-mass spectrum</p>	<p>40.00 per sample or acquisition 35.00 per sample requiring precipitation purification prior to analysis</p>
<p>LC/MS and LC/MS/MS (Liquid Chromatography/ Mass Spectrometry) Ion-trap, triple quad, or Q-ToF instruments</p> <p>Positive-ion ESI; sample introduction by nanoLC or microLC</p>	<p>110.00 per LC/MS run 15.00 per sample for Ziptip™ purification as required</p> <p><i>Discount rates apply for high number of samples and depending on gradient or isocratic LC runs</i></p>
<p>MALDI/ ToF (Matrix-assisted Laser Desorption Ionization/ Time of Flight)</p> <p>Positive-ion or negative-ion MALDI; linear or reflectron mode</p>	<p>25.00 per sample or acquisition 45.00 per accurate-mass measurement (i.e. high-resolution measurement with internal standard) 15.00 per sample for Ziptip™ purification as required 8.00 per sample or acquisition if the user provides the samples pre-spotted on their own plate</p>
<p>Special Project/ Method Development</p>	<p>Inquire</p>