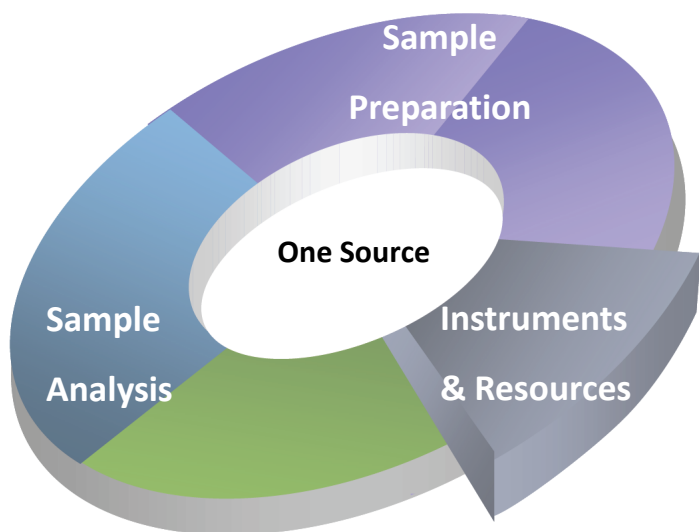
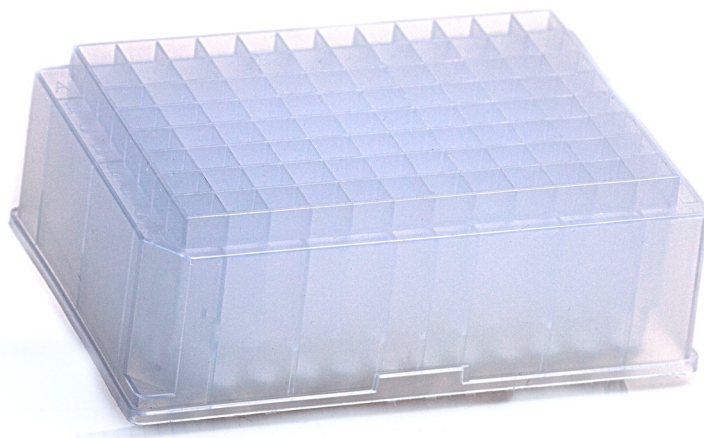


*Purity™* is the unique Phospholipid depletion device which individually integrates screen filtration with the targeted removal of phospholipids and proteins in plasma/serum. The technology utilizes Acidic particles which exhibit selective affinity towards phospholipids while remaining non-selective towards a range of basic, neutral and acidic compounds. The phospholipid retention mechanism is based on highly selective Lewis acid-base interaction between the ions (functionally bonded to the SPE stationary phase) and the phosphate moiety consistent with all phospholipids. It eliminates ion suppression through protein precipitation and phospholipid removal.



### Protocol 1:

1. Precipitate Proteins by adding 300  $\mu\text{L}$  of 1% formic acid in acetonitrile to the Phospholipid removal plate or spin column, followed by 100 $\mu\text{L}$  plasma or serum.
2. Mix by vortexing /shaking plate or by aspirating or dispensing with 0.5 -1mL pipette tip.
3. Apply Positive Pressure or Vacuum- The packed-bed filter / frit assembly acts as a depth filter for the concurrent physical removal of precipitated proteins and phospholipids. Small molecules (e.g. pharma compounds and metabolites) pass through unretained.

### Protocol 2:

1. Add plasma and crash solvent in desired ratio for optimal lipid and protein removal, 3:1 methanol to plasma ratio is recommended. Modification of pH can be made to ionize the analyte of interest. For basic compounds, we recommend 0.1% to 1.0% formic acid in methanol. For acidic compounds, we recommend 5mM to 10mM ammonium formate buffers at pH 9.
2. Mix by vortexing / shaking plate or by aspirating / dispensing with 0.5 - 1mL pipette tip.
3. Apply vacuum or Positive Pressure. The packed-bed filter/frit assembly acts as a depth filter for the concurrent physical removal of precipitated proteins and chemical removal phospholipids. Small molecules (e.g., pharmaceutical compounds and metabolites) pass through unretained.

