## buffers

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```
[1]: import pandas as pd
[2]: Tris_con = 121.14 # g/mol
     NaCl\_con = 58.44 \# g/mol
     Imidazole_con = 68.08 # g/mol
     TCEP_{con} = 286.65 \# g/mol
     HEPES_{con} = 238.31 \# g/mol
     KCl\_con = 74.55 \# g/mol
     Tris_mol = {
         "Lysis_buffer" : 20/1000,
         "Elute buffer" : 20/1000,
         "Low-salt buffer" : 20/1000,
         "High-salt buffer" : 20/1000,
         "Dilute buffer" : 20/1000,
         "SEC buffer" : 0
     }
     NaCl_mol = {
         "Lysis_buffer" : 500/1000,
         "Elute buffer" : 500/1000,
         "Low-salt buffer" : 100/1000,
         "High-salt buffer" : 1000/1000,
         "Dilute buffer" : 0,
         "SEC buffer" : 0
     }
     Imidazole_mol = {
         "Lysis_buffer" : 20/1000,
         "Elute buffer" : 400/1000,
         "Low-salt buffer" : 0/1000,
         "High-salt buffer" : 0/1000,
         "Dilute buffer" : 0,
         "SEC buffer" : 0
     }
     glycerol = {
```

```
"Lysis_buffer" : 5/100,
   "Elute buffer" : 5/100,
   "Low-salt buffer" : 5/100,
    "High-salt buffer" : 5/100,
   "Dilute buffer" : 5/100,
   "SEC buffer" : 0
}
TCEP_mol = {
   "Lysis_buffer" : 1/1000,
   "Elute buffer" : 1/1000,
   "Low-salt buffer" : 1/1000,
   "High-salt buffer" : 1/1000,
   "Dilute buffer" : 1/1000,
   "SEC buffer" : 1/1000
}
HEPES_mol = {
   "Lysis_buffer" : 0/1000,
   "Elute buffer" : 0/1000,
   "Low-salt buffer" : 0/1000,
   "High-salt buffer" : 0/1000,
   "Dilute buffer" : 0,
   "SEC buffer" : 20/1000
}
KCl_mol = {
   "Lysis_buffer" : 0/1000,
   "Elute buffer" : 0/1000,
   "Low-salt buffer" : 0/1000,
   "High-salt buffer" : 0/1000,
   "Dilute buffer" : 0,
   "SEC buffer" : 250/1000
}
Volume = {
   "Lysis_buffer" : 1,
   "Elute buffer" : 0.5,
   "Low-salt buffer" : 1,
   "High-salt buffer": 0.5,
   "Dilute buffer" : 0.5,
   "SEC buffer" : 1
}
concentration = {
   "Tris" : Tris_con,
   "NaCl" : NaCl_con,
```

```
"Imidazole": Imidazole_con,

"TCEP": TCEP_con,

"HEPES": HEPES_con,

"KC1": KC1_con
}

buffer_list = ["Lysis buffer", "Elute buffer", "Low-salt buffer", "High-salt_

$\times \text{buffer}$ buffer", "SEC buffer"]

## other way around is more effective: a dictionary for every buffer
```

```
[3]: # Molar masses (g/mol)
     concentration = {
         "Tris": 121.14,
         "NaCl": 58.44,
         "Imidazole": 68.08,
         "TCEP": 286.65,
         "HEPES": 238.31,
         "KC1": 74.55
     }
     \# Buffers with molar concentrations (mol/L) and glycerol in \%
     buffers = {
         "Lysis buffer": {
             "Tris": 20/1000.
             "NaCl": 500/1000,
             "Imidazole": 20/1000,
             "TCEP": 1/1000,
             "HEPES": 0,
             "KC1": 0,
             "Glycerol": 5/100,
             "Volume": 1
         },
         "Elute buffer": {
             "Tris": 20/1000,
             "NaCl": 500/1000,
             "Imidazole": 400/1000,
             "TCEP": 1/1000,
             "HEPES": 0,
             "KC1": 0,
             "Glycerol": 5/100,
             "Volume": 0.5
         },
         "Low-salt buffer": {
             "Tris": 20/1000,
             "NaCl": 100/1000,
             "Imidazole": 0,
             "TCEP": 1/1000,
```

```
"HEPES": 0,
        "KC1": 0,
        "Glycerol": 5/100,
        "Volume": 1
    },
    "High-salt buffer": {
        "Tris": 20/1000,
        "NaCl": 1000/1000,
        "Imidazole": 0,
        "TCEP": 1/1000,
        "HEPES": 0,
        "KC1": 0,
        "Glycerol": 5/100,
        "Volume": 0.5
    },
    "Dilute buffer": {
        "Tris": 20/1000,
        "NaCl": 0,
        "Imidazole": 0,
        "TCEP": 1/1000,
        "HEPES": 0,
        "KC1": 0,
        "Glycerol": 5/100,
        "Volume": 0.5
   },
    "SEC buffer": {
        "Tris": 0,
        "NaCl": 0,
        "Imidazole": 0,
        "TCEP": 1/1000,
        "HEPES": 20/1000,
        "KC1": 250/1000,
        "Glycerol": 0,
        "Volume": 1
    }
}
```

```
for buffer_name, components in buffers.items():
    volume = components["Volume"] # in liters
    row = {"Buffer": buffer_name}
    for compound in concentration:
        mol_conc = components.get(compound, 0)
        mass = mol_conc * concentration[compound] * volume # g = mol/L * g/mol_
        ** L
        row[compound] = mass
```

```
# Glycerol as \% v/v, approximated as g/mL ~ mL (density 1.26 g/mL, but _{f U}
      ⇔assuming ~1 g/mL here)
         glycerol_percent = components.get("Glycerol", 0)
         glycerol_mass = glycerol_percent * volume * 1000 # convert L to mL
         row["Glycerol"] = glycerol_mass
         mass data.append(row)
     # Create DataFrame
     df_mass = pd.DataFrame(mass_data)
     df_mass.set_index("Buffer", inplace=True)
     df_mass.columns = [f"{col} [g]" for col in df_mass.columns]
     df_mass.round(3) # Round for display
[4]:
                       Tris [g] NaCl [g] Imidazole [g] TCEP [g] HEPES [g] \
    Buffer
    Lysis buffer
                          2.423
                                   29.220
                                                    1.362
                                                              0.287
                                                                         0.000
    Elute buffer
                          1.211
                                   14.610
                                                   13.616
                                                              0.143
                                                                         0.000
    Low-salt buffer
                          2.423
                                   5.844
                                                    0.000
                                                              0.287
                                                                         0.000
    High-salt buffer
                          1.211
                                   29.220
                                                   0.000
                                                              0.143
                                                                         0.000
    Dilute buffer
                          1.211
                                    0.000
                                                   0.000
                                                              0.143
                                                                         0.000
     SEC buffer
                          0.000
                                    0.000
                                                    0.000
                                                              0.287
                                                                         4.766
                       KCl [g] Glycerol [g]
    Buffer
    Lysis buffer
                         0.000
                                        50.0
    Elute buffer
                         0.000
                                        25.0
    Low-salt buffer
                         0.000
                                        50.0
    High-salt buffer
                         0.000
                                        25.0
    Dilute buffer
                         0.000
                                        25.0
    SEC buffer
                        18.638
                                         0.0
[5]: # Molar masses (q/mol)
     concentration = {
         "Tris": 121.14,
         "NaCl": 58.44,
         "Imidazole": 68.08,
         "TCEP": 286.65,
         "HEPES": 238.31,
         "KC1": 74.55
     }
     # Stock concentrations (mol/L or %)
     stock_concentrations = {
         "Tris": 1.0,
         "NaCl": 5.0,
         "Imidazole": 1.0,
         "TCEP": 0.5,
```

```
"HEPES": 1.0,
    "KC1": 3.0,
    "Glycerol": 100.0 # % v/v
}
# Volume of stock to prepare in liters
volume_L = 0.1 # 100 mL
# Calculate mass
for compound, stock_conc in stock_concentrations.items():
    if compound == "Glycerol":
       print(f"{compound}: 100 mL of pure glycerol (used as 100%)")
   else:
       molar_mass = concentration[compound]
       moles = stock_conc * volume_L
       mass = molar_mass * moles
       print(f"{compound}: {mass:.2f} g in 100 mL for a {stock_conc} M_L
 ⇔solution")
```

Tris: 12.11 g in 100 mL for a 1.0 M solution NaCl: 29.22 g in 100 mL for a 5.0 M solution Imidazole: 6.81 g in 100 mL for a 1.0 M solution TCEP: 14.33 g in 100 mL for a 0.5 M solution HEPES: 23.83 g in 100 mL for a 1.0 M solution KCl: 22.37 g in 100 mL for a 3.0 M solution Glycerol: 100 mL of pure glycerol (used as 100%)

```
[6]: # Define standard stock concentrations in mol/L or % for qlycerol
    stock_concentrations = {
                         # M
        "Tris": 1.0,
        "NaCl": 5.0,
                          # M
        "Imidazole": 1.0, # M
        "TCEP": 0.5,
                          # M
        "HEPES": 1.0,
                          # M
        "KC1": 3.0,
                          # M
        "Glycerol": 100.0 # %
    }
    # Calculate stock volumes needed for each buffer (in mL)
    stock_volumes_data = []
    for buffer_name, components in buffers.items():
        volume = components["Volume"]
        row = {"Buffer": buffer_name}
        for compound, stock_conc in stock_concentrations.items():
            final_conc = components.get(compound, 0)
```

```
if compound == "Glycerol":
                 # For qlycerol: use % directly
                 stock_volume_ml = (final_conc / stock_conc) * volume * 1000
      ⇔convert to mL
             else:
                 # For molar compounds
                 stock_volume_ml = (final_conc * volume) / stock_conc * 1000 #_
      ⇔convert to mL
             row[f"{compound} stock [mL]"] = stock_volume_ml
         stock_volumes_data.append(row)
     # Create DataFrame
     df_stock_volumes = pd.DataFrame(stock_volumes_data)
     df_stock_volumes.set_index("Buffer", inplace=True)
     df_stock_volumes.loc["Total"] = df_stock_volumes.sum()
     df_stock_volumes.round(2)
[6]:
                       Tris stock [mL] NaCl stock [mL]
                                                          Imidazole stock [mL] \
    Buffer
                                  20.0
                                                  100.0
                                                                          20.0
    Lysis buffer
    Elute buffer
                                  10.0
                                                   50.0
                                                                         200.0
    Low-salt buffer
                                                   20.0
                                  20.0
                                                                           0.0
    High-salt buffer
                                  10.0
                                                  100.0
                                                                           0.0
    Dilute buffer
                                  10.0
                                                     0.0
                                                                           0.0
    SEC buffer
                                   0.0
                                                     0.0
                                                                           0.0
     Total
                                  70.0
                                                  270.0
                                                                         220.0
                       TCEP stock [mL] HEPES stock [mL] KCl stock [mL] \
    Buffer
                                                      0.0
                                                                     0.00
    Lysis buffer
                                   2.0
                                                                     0.00
     Elute buffer
                                   1.0
                                                      0.0
    Low-salt buffer
                                                                     0.00
                                   2.0
                                                      0.0
    High-salt buffer
                                   1.0
                                                     0.0
                                                                     0.00
     Dilute buffer
                                   1.0
                                                     0.0
                                                                     0.00
     SEC buffer
                                                                    83.33
                                   2.0
                                                     20.0
     Total
                                                                    83.33
                                   9.0
                                                     20.0
                       Glycerol stock [mL]
    Buffer
    Lysis buffer
                                      0.50
    Elute buffer
                                      0.25
    Low-salt buffer
                                      0.50
    High-salt buffer
                                      0.25
    Dilute buffer
                                      0.25
     SEC buffer
                                      0.00
     Total
                                      1.75
```