

## Micro-Gas Chromatograph – SRA INSTRUMENTS

### Connection

#### Serie R, Serie N, Serie S

Your MicroGC communicate with 2 serial links  
These 2 x RS232 are connected to the computer with the provided cables

#### Serie PGC-3000

Your MicroGC communicate with 2 x RS485 serial links  
Refer to ATEX norm for proper connection

#### MyGC

Internal links are done & installed on COM 5 & 6



### Carrier Gas

Your MicroGC can use **He, H<sub>2</sub>, N<sub>2</sub>, Ar** as carrier gas  
Please check the correct carrier gas configuration  
A wrong configuration would damage your instrument

**Quality = 99,9995 % purity**  
**Pressure = 5,5 bars ± 0,2 bars (80 ± 2 PSI)**

Connection between pressure regulator and MicroGC :

- Use 1/8" stainless steel or clean copper tube and Swagelok fittings
- Purge all tube and dead volumes
- Don't use liquid leak detector

(GasType menu)  
User : **service**  
Pwd : **gasconfig**

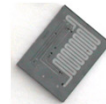


### Sample

You have to clean your sample and control the pressure if necessary :

**Max. Pressure = 2 bars (rel.)**  
**No liquid / No solid**

Use the in-line frit to protect your injector from particles



Use a "Genie filter" if your sample has a positive pressure

### Other recommendations...

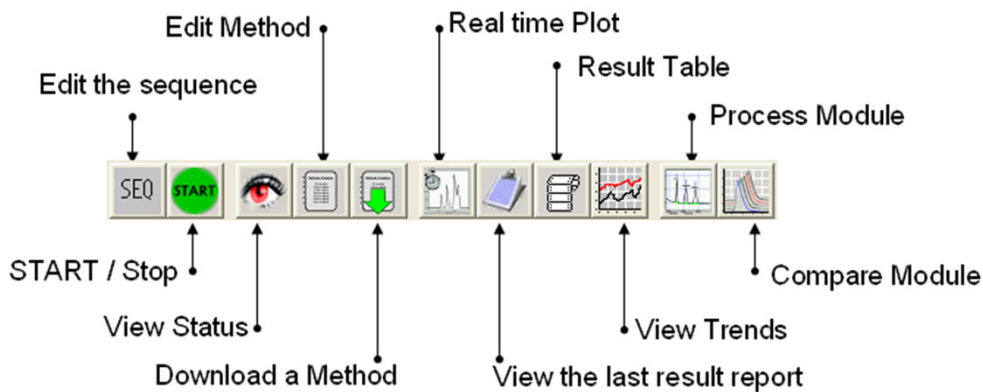
- When you start the MicroGC : download a « purge » method with all detectors OFF to purge columns for about 10 minutes
- You can bake out the column at max. temperature : bake out duration is between 24 and 48 hours
- Download a standby method with detectors OFF when MicroGC is not used



It's recommended to let carrier gas flow on the column all the time

## SOPRANE : the MicroGC software

### Menu – Soprane



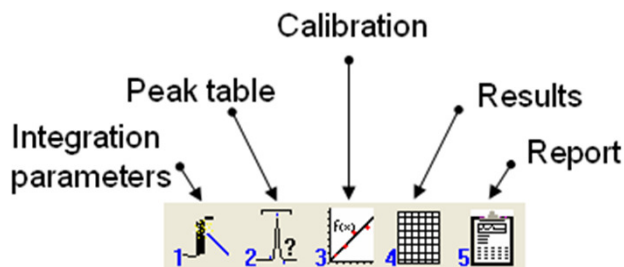
### Develop a method

0. Analytical parameters (Temp., Press., Time...)
1. Integration parameters
2. Peaks identification (Peak table)
3. Calibration / Quantification
4. Results
5. Report



When you develop a method for the first time, always proceed step by step and channel by channel.

Shortcuts :



### Mouse & Keyboard shortcuts

#### ZOOM (all the software) :

Unzoom chromatogram view : right double click

Keyboard : use arrows to move and CTRL + arrows to zoom



#### Identification (Process module) :

In the identification view, use CTRL + right click from a peak to add it in the table

#### Results (Soprane main screen) :

Export your results in Excel (\*.dif file) by clicking "Export to" in the results table

