

# Automatic device for electrophoretic analysis

# User Manual





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# 1 SAIO: Preamble

SAIO performs the automatic electrophoresis of Serum Proteins, Hemoglobins, Lipoproteins, Isoenzymes, not concentrated Urine protein etc., and other methods that require incubation step, which will be performed in a semi-automatic way. Additional specific instructions for the methodologies will be describe in the provided kits.

The equipment of the machine consists of:

• A robotic unit that performs in a fully automated way all the phases of the process: the buffering of strips from 1 to 2, which can be made of cellulose acetate or other kind of supports, the drying of the strip, the samples deposit, migration, staining, destaining, reading with incorporated densitometer;

- A built-in Computer runs the robotic controllers ;
- A software for the machine functioning and management of the program.

#### 1.1 User Manual and Maintenance

SAIO was designed and developed in full compliance with all requirements of safety and healthy requirements. Please READ CAREFULLY every part of this guide before installing and use the machine. Here you will find all the information on how to use the machine and how to keep it efficient. Keep this booklet with care, in case you lose it, ask for another copy to the manufacturer. If you sell or give the machine to another person, do not forget to hand this booklet to the new user.

The manufacturer disclaims any liability for any damage that may be directly or indirectly resulting to persons, things and animals occurred as consequence of a wrong observance of the procedures given and explained in this user guide, especially the instructions regarding the installation, the right use and the maintenance of the machine.

#### <u>Index</u>

- 1. Safety Notes
- 2. Unpacking and Installation.
- 3. Operating modes.
- 4. Reagents and samples loading.
- 5. Placement of the membranes of cellulose acetate.
- 6. Running of Machine.
  - a) Serumprotein
  - b) Hemglobin

- 7. Unloading, cleaning and maintenance.
- 8. Guarantee Conditions.

# 1.2 Safety Notes

The chief of the laboratory must run all the working procedures in the correct way and he must adopt collective and individual protective measures from exposure: all the workplaces, where biological agents are handled, must be marked with the appropriate risk signal, and they should be provided adequate sanitation measures in case of accidental propagation and emergency procedures. During SAIO use the following measures have to be adopted:

- SAIO must be used strictly in accordance with the purpose for which it was designed and as specified in this guide. The machine must be used only by qualified persons, any other type of use is considered improper and therefore dangerous. The manufacturer will not assume any responsibility for damages arising out of misuse.
- Always remove the plug from the power outlet before cleaning or doing any maintenance.
- Any modification to the machine is extremely dangerous and it implies the end of the guarantee.
- In case you need an intervention to the electrical system, please contact only those persons who are qualified and authorized.
- To ensure the efficiency of the machine and its proper functioning is essential to follow the manufacturer instructions and call only qualified and authorized persons for the maintenance. It should be checked at least twice a year.
- Switch it off incase of failure and/or malfunctioning, do not touch it and contact the manufacturer and /or an authorized service center.
- When using the electrical equipment the following safety measures must be complied: Do not touch the appliance with wet or damp hands or feet; do not unplug by pulling on power cord. To unplug, grasp the plug not the power cord; do not expose the appliance to the weather (rain, sun, etc.); when replacing the power cord use only another one with the identical technical characteristics.
- When the machine is not used anymore, it is recommended to unplug the power cord with the hands.

- To ensure safe operations, such as the analysis, it is important to have an adequate lighting at a workstation.
- Eating, drinking, smoking, handling contact lenses, applying cosmetics, and storing food for human consumption must not be permitted in laboratory areas.
- Never pipet by mouth. Use of the automatic pipet.
- Do not forget to use DPI, which is any equipment that must be worn and kept on in order to protect the body from risks of security (laboratory clothing, disposable sterile gloves, and goggles).
- Decontaminate work surfaces before and after use, and immediately after spills (see point "1.6 Draining, cleaning and maintenance procedures");

Avoid unnecessary proximity to high EMF sources (e.g. radiology equipment and CT scanners), that may cause interference with the machine, thereby affecting its proper functioning.

# 1.3 Unpacking

Make sure the package is intact and has no obvious signs of damages, otherwise do not install the machine, but complaint to the carrier and inform our agent or the distributor. Open the box from the top, according instructions, and pay attention to take out the machine upright. (Picture 1.1)



**Picture 1.1 - Unpacking** 

Check the following contents of the box:

- Electrophoresis machine
- User Manual;
- Power Cord;
- External Sensor Cable;
- Reagent containers;
- Starter kit containing absorbent cardboards, buffer, staining, destaining, dry or wet acetate strips, smart card. This kit allows you to carry out electrophoresis tests;

#### 1.4 Installation

Put the machine on a table or on a work bench perfectly level.

Open the Plexiglas cover of the machine, remove any sponges or tape.

Unlock the cart by removing the lock located on the steel frame on the left side, it is sufficient to unscrew anticlockwise the pommel on the PVC. That lock must be stored; it can be useful in case of relocation of the machine it must be reapplied in order to prevent any movement of the cart.

The instrument contains from 1 to 2 strips holder and 1 sample holder base, but the sera applicator will be in another provided container. On the left side it is clearly visible an opening with all computer connectors to be linked to external devices.

On the same panel there is also the master power switch, the electric switch and two and 1.6 A fuses. A smart card reader is placed vertically on the left side of the optical reader, that is located on the worktable of the machine.

On the front side of the machine, on the right, there is the switch button that becomes illuminated when the machine is on.

The machine equipment includes also some tanks containing reagent which load it the reagent in the both chambers (Imbibition and staining / Destaining) Migration Chamber.

There are 6 tanks of reagents: Staining, Destaining, Serumprotein Buffer, Hemoglobin Buffer, Distilled Water, Waste.

The kit will contain the reagents required for SAIO processes of electrophoresis with all the concerned methods. All machines are supplied with sensor cable, which connect the machine with the external tanks. Pay attention when filling the tanks with the appropriate reagents, as indicated on the caps and on the tanks, and pay attention when cleaning them.

In Picture 1.2 there are junctions of tubes connected to the pipes that go to the reagent tanks and liquid discharge. On each socket it is indicated the predisposed liquid.



Picture 1. 2 - Connections of probes on the external level

On the same plate there is also a black female connector which must be connected to the male connector, which is linked to the sensors of the tanks. The male connector can be inserted snugly into the receptacle, the female connector, then turn it clockwise to ensure a reliable physical and electrical connection.

Before going on with the electrical connections, it is necessary to verify that the main voltage of the network corresponds to the values shown on the rating plate located on the lateral side of the machine. Moreover it is important to check sure that the residential electrical system is adequate for the maximum power absorbed by the unit, which is indicated on the rating plate.

The electrical safety of the machine is ensured only if it is properly connected to an efficient grounding system in compliance with all electrical safety standards.

The manufacturer cannot be held responsible for any damage caused by the absence of the grounding system. Also make sure the connection is made at a plant and outlets of adequate capacity to power absorption.

In general, the use of adapters, multiple sockets and/or extensions is not recommended. If they must be used, utilise only those single or multiple adapters and extensions which comply with the current safety regulations. However, make sure not to exceed the current rating marked on the single adapters and extensions, or the maximum power marked on the multiple adapters.

Plug the power cord into the socket with switch and with fuse tubes and connect it to 220V outlet.

At this point SAIO machine can be switched on using the switch located on the socket group, and then pushing the power button on the front side of the machine. Once the machine is switched on, it will not move till the start command will be issued. After the first switching on, it is recommended to run the RESET command following the instructions of the software.

(Control Machine  $\rightarrow$  users service)

The delicate software manages every movements of the machine and all its sensors. At the end of the electrophoretic process the software acquires the reading data from the densitometer, than it analyses them and creates a densitometric report with all the data related to the examined sample.

#### 1.5 Operating Procedures

#### **Loading of Reagent**

This user guide contains instructions about the loading related to the electrophoresis of serum proteins. Other analysis procedures can be found described in the kits, with everything needed to run them .



Picture 1. 3 – Saio Complete Plan

The interior of the unit is accessed by opening the cover (the power to the electrical circuits and the motors will be consequently automatically interrupted).

- 1. Robotic arm (pliers)
- 2. SmartCard reader
- 3. Positioning of the lid of the migration chamber
- 4. Densitometer reader
- 5. Applicator with an automatic wash distilled water
- 6. Serum holder with 16 basin where inject serum (25-30 micro liters)
- 7. Migration chamber with bridges (papers holder) with bibulous paper 70x20mm

that the absorbing the buffer become electrically conductive and keep adequately wet the strip during migration.

- 8. Cover of the migration chamber
- 9. Positioning of strip drying
- 10. Imbibition strip tank
- 11. Tank for staining and destaining of the

#### Necessary content to run 400 serum proteins tests

- 2 x 25 pieces of strips supported on Dry Plates
- 1 x 100 ml buffer Tris hippurate
- 1 x 200 ml Staining Ponceau S ready to use
- 2 x 250 ml Destaining Concentrate 20 X
- 20 Applicator papers (70 x 13) mm
- 6 Strip drying papers (92 x 47) mm
- 4 Migration chamber floaters (70 x 20) mm

Product identification and intended use

The reagents and accessories for the clinical electrophoresis diagnosis of serum proteins are intended to be used to research of disproteinemie and to quantify albumin, alpha-1, alpha2, Beta and gamma globulins. The reagents should be used only for the purpose that has been defined above and by a qualified laboratory.

Use fresh serum, as specified in laboratory procedures and in conformity with the guidelines. Avoid the use of plasma, because the fraction of fibrinogen may disturb the interpretation of the gamma area. Do not use samples hemolyzed or containing cells.

#### **Reagents preparation**

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Concentrate buffer: Pour about 80 ml of distilled water in the bottle

#### Ponceau S destaining ready to use

Concentrate destaining 20 X Pour about 5 liters of distilled water in the bottle

Keep strip and reagents to a ambient temperature  $+15^{\circ}/+30^{\circ}$ C.

The best-before dates are indicated on the labels.

Once opened the reagents are stable and can be used until expiration date, if closed immediately after use and protected from direct sunlight avoiding the evaporation.

#### Warning:

If black spots or mold inside the bottle will occur it will be necessary to remove the buffer If black spots or mold inside the bottle will occur it will be necessary to remove the destaining

The products are intended for professional use in laboratories. The left over products, the waste resulting from their use and the containers must be removed in accordance with current legislation

#### Protocol

Insert the absorbing paper (floating) in the migration chamber.

Insert the paper for the drying of the strip, that is located on ton one side of the migration chamber.

Supported strips on Mylar, that can be found in the kit, can be applied on special strips holder frames. It is necessary to hook these strips matching the holes with the pins present on the strips holder, after having lifted the safety lock, making a slight pressure on the central part of strip holder. Place the strips holder provided with strips in the chamber containing the electrophoresis buffer. If you have to run more than 8 electrophoresis, load also other strips on the additional strips holder.

#### Serum holder

Remove the serum holder frame by sliding outward.

Pipette from 25 to 30 ml of each sample, starting from the basin with number 1 (first row and the first basin on the operator side), and put the serum in the order indicated (see Picture 1. 3) Insert the 70 x 14 mm paper buffer into the proper strips holder. (See Picture 1.4)



**Picture 1.4 – Insert samples** 



Picture 1.5 - Insert serum holder

Close the lid of the machine.

Choose the method and start the electrophoretic process.

#### Scanning of the electrophoretic traces.

At the end of the process of staining and drying, the strips take automatically into the reading position in the scanning compartment.

Afterwards it is possible to proceed with the analysis of the performed readings and make any corrections. To include any demographic data and other actions on the charts and on the printers, consult the user guide in the section "2. Introduction to Software"

After turning off the machine it is necessary to empty the serum holder and clean all other components

#### 1.6 Draining, cleaning and maintenance of the machine

Follow these guidelines to perform the unloading and cleaning operations.

- Turn Off the instrument
- Open the cover of the Instrument
- Remove the strips holder frame, release the membranes,
- Wash the frames with running water and let them dry.
- Remove the absorbing buffers from the strips holder and from the dripping support of the frame. Throw them into the special waste.
- Extract the serum Holder and remove the serum from the wells and wash it with distilled water
- Put all components in the proper places in the machine.
- Close the lid of the machine.

#### Maintenance operation

The Maintenance must be performed only by qualified personnel, during repairs remove the plug from the socket.

Disassemble the back panel by unscrewing the fixing screws. At this stage pay attention when unplugging the cable used for grounding system.

The manufacturer will provide appropriate instructions and wiring diagrams to the authorized personnel for the maintenance and/or repairs. The customer should always contact authorized service centres, or, when possible, send the machine to the head office.

#### 1.7 Warranty Conditions

SAIO is guaranteed for 12 months from the purchase date, which is shown on a valid receipt given by the seller.

The warranty includes free repair or replacement of component parts of the machine that are

imperfect because of manufacturing defects.

Warranty does not cover damage or defects due to:

- Negligence or careless use (failure to follow the safety instructions),

- Improper installation or maintenance operated by non-qualified personnel,

- Damage caused by transportation or any circumstances which cannot clearly be traced back to defects resulting from errors in the manufacturing process.

The warranty does not cover any rubber tubing / silicone, neither the non-return valves, the filters for the reagents or for washing water, because they can be subject to natural wear or natural contamination during normal operations of the machine.

The warranty does not cover improper installation with not in conformance with the maintenance procedures as specified in this user guide.

The warranty does not cover any damages caused to the unit by improper use, customer neglect, or repairs performed by an unauthorized technician.

In case of repair at an Authorized Service Center, all transport risks and any costs will be borne by the purchaser. It is in any case excluding the extension of the guarantee as a result of any failure.

#### 1.8 Procedures for the disposal of the machine.

To dispose the machine SAIO is necessary to contact specialized companies as permitted or required by applicable law, they will be responsible to recycle and dispose the machine.

#### 1.9 Technical Specifications

Analysis	Electrophoresis	of	Serum	Protein,	Hemoglobin,	(Urine	Protein,

	Lipoprotein)
Electrophoretic Media	Dry cellulose acetate plates (available in 2 sizes)
Analytical Capacity of	Up to 16 samples per cycle on 2 plates
standard plate	
Analytical Capacity of mini	Up to 8 samples per cycle on 2 plates
plate	
Analytical Throughput -	8 micro samples in about 30 minutes
standard plate	16 micro samples in about 50 minutes
Analytical Throughput -	4 micro samples in about 30 minutes
mini plate	8 micro samples in about 50 minutes
Sample Plate	Removable, with 16 wells
Robotic Arm	Electronic check integrated for presence of objects
Applicator Washing	Automatic by circulation of deionized water
Migration Chamber	Automatic loading of buffer solution
Staining and Destaining	Automatic loading of solutions
Trays	
Liquid Control	By infrared sensor and by time control
Interface	USB, LAN
Power Supply	220 – 240 V ac 50 – 60 Hz
Dimensions	50x45x45 cm
Weight	25 Kg
Operative System	Windows 7
User Interface	Keyboard and mouse (not included) by USB or PS/2
Monitor (not included)	VGA connection
Computer	Integrated into the instrument
Reading System	8 indipendent channel
Bright Source	8 Led ultrabright
Densitometer Linearity	0 to 2.8 D.O.
Software	Dedicated for instrument and densitometer management

# 2 Introduction to Software

SAIO has a program that exploits all the possibilities that are offered by the intuitive Windows lite. In addition to a quantitative, reliable and fast electrophoretic analysis, SAIO can record all the data in a standard database, which is easily and rapidly reachable.

The program is structured with windows that help the user in the sequence of operations to be performed.

On the first screen you can start clicking with the mouse on the menu, choosing the type of operation.

The program is very flexible, both during the performing of the electrophoretic process, and after reading the samples.

During the execution electrophoresis, in order to speed up the operations, it is possible to modify the traces of an earlier scanning and to keep the data archive.

The scan results are automatically presented on the video, and it is possible to correct them and include any comments. Within the program there is an interactive curves editor, which can also modify the trace.

The editor shows the image of migration, the resulting curve and the numerical results with the identification of normality and with the personal data of the patient. It is possible to change the position of the minimum, the names of the fractions and to add comments by simply clicking the mouse on the corresponding button that is present on the window. Using the 'Previous' and 'Next' key, you can scroll through the results of the various patients.

An interesting feature of SAIO is the printing: it produces a report that, apart from the graph of the curve and the numerical results, also contains the image of migration, the personal data and comments. The resulting printing can then be delivered directly to the patient, with no need to attach other documents such as reports or images of migration

### 2.1 Tutorial

Before the detailed description of the operations offered by SAIO, the user may consult the following tutorial for an overview of the operational program.

a) Start the machine double clicking on SAIO icon at the center of the Desktop.

**b**) When initializing SAIO a splash page will appear (see Picture 2.1).



Picture 2. 1 – Splash page

**a)** After initialising stage the "Today's Data" will appear as first page. (*see Picture 2.2*).

L.:Ma	r 13 2012 22:20:29)			
File Features ?	serum bolder	acks		
Archive				
Archive seru	m holder today	Current Method: Sieroproteine	Job Data: 27/04/2012	^
		Patient N.: Port serum 1 0 0	N.: Slot N.: Patient Type: 255 Human	~
Port serum		Surname:	Name:	
	State Empty	Code Acceptance:	Total Protein: Age: Sex	le 🗸
	N. Port serum 00	Date Acceptance: 27.04.2012	Birth Date:	
	Notes	Data Lists:		42 <sup>4</sup>
00				
	Slot n. 1			
Control Machine	N. Retention Stru	r's data 🔍 Search Exams 💽	Archive	V
Ready		Sieroproteine	27.04.20	12 - 20:31:55

Picture 2. 2 – Today's Data

SAIO utilises a programme with two menus, the first one appears on the top of the application and is a dropdown menu (see Picture 2.3);

In the dropdown menu you can select some useful features:

• From the "File" menu you can select the "Select language" command to access to the multilanguage program;

- From the menu "Features" you can select the commands:
  - ✓ "Reset Layout File" to restore size and positions of objects on the screen;
  - ✓ "Turning off the alarm" to turn off the alarm after the completion of an electrophoretic process.
  - ✓ "Loading Kit" to load the kit providing the serial number, the unlock code, the serial machine inventories.

✓ "Alarm Kit Remainder" to indicate the minimum threshold number of strips.



Picture 2. 3 – Dropdown menu

The second menu is composed of two levels:

The first level (see Picture 2. 4), below the application, shows all the functions exported by SAIO. The second level (see Picture 2.5), up to the application, shows the subdivisions of feature selected on the first level.

The program also includes a status Bar (see Picture2.6) that provides the user with further information. In particular the bar, starting from the left, is divided into three fields:

- *State*, to indicate a process running on the machine
- *Selected Method*, to indicate the reference method during the electrophoretic process of the samples.
- *Date and time*, date and hour of the day.



Ready Sieroproteine 09.05.2012 - 10:15:56 Picture 2. 6 - Status Bar

- **b**) Before starting the electrophoretic process, please ensure that you have selected the appropriate method:
- If you have selected the appropriate method with the correct setting of the parameters go to **step e**).
- In the case the method has not been correctly selected or it is the first use of the software or any need to modify the parameters, see **Chapter 3 Methods** and then go to step e)
- c) In "Current data" click on the yellow icon to add a new sample holder
- **d**) Digit the number of patients that must be added to serum holder and click OK (see Picture 2.7).

Port serum		
	State Preparation	Aries Enter the number of patients to be added to port serum (Min.1-Max.24)
	N. Port serum 01	8
	Notes Serum holde 27/04/2012 - J	er of 20:47:08
	Slot n. 1 N. Retention	Str0

Picture 2. 7 – Insert number Patient

e) When inserting the number of patients (from a minimum of 1 to a maximum of 16) on the image of the serum holder, on the left, you will see the same number of wells

becoming red. The green well indicates that it is possible to insert the data of the patient by writing in the tab highlighted in yellow. (see Picture 2.8).

File Features ?									
Management	serum holder 📋 🛅 Change Trac	ks 📄 Print							
Archive	m holder today								
= = Steroprot	enne 2012 46 (1)	Current Meth Sieroprotein	nod: Ie			Job Data: 27/04/2012			^
		Patient N.: 1	Port serum N. 1	.: Slot N.: 1	Patient Ty Human	pe:		~	and the
Dert corum		Surname:		-	Name:				
	State	Code Accept	tance:	Tota	l Protein:	Age:	Sex:		-
	Preparation					<b>*</b>	Male	~	
000	N. Port serum 01	Date Accepta 27.04.2012	ance:	~	Birth Date 27.04.2	2012		×	
ŎŎŎ	Notes	Data Lists:							
	Serum holder of 27/04/2012 - 20:46:07		×						
	Slot n. 1								
Control Machine	e Methods Moder's	data 🖸 Searc	th Exams 🛛 💽 A	Archive					~
Ready	Terra succession Terra	Sier	oproteine			2	7.04.2012 - 2	0:46:44	

Picture 2.8 - Managment serum holder

**f)** UAfter the insertion of all the data of the first patient, (see Picture 2.9) click the yellow icon with a *floppy disk* symbol to save data and confirm by pressing OK

Current Met Sieroproteir	hod: Ie			Job Dat 27/04/20	ta: )12	1	
Patient N.:	Port serum N.	: Slot N.:	Patient T	ype:		_D	
1	1	Human				*	
Surname:			Name:				
James			Camero	n			
Code Accep	tance:	Tota	I Protein:	Age:		Sex:	
52986		7.6		0	*	Male	*
Date Accept	ance:		Birth Date	e:			
27.04.2012		~	27.04.	2012			×
Data Lists:							
	<b>v</b>						

Picture 2. 9 – Save Data

**g)** Once the data of the patients have been entered and saved, the SAIO automatic process can be started by clicking on the key that lights up with yellow icon (see Picture 2.10).

**NOTE**: <u>it is possible to insert the data after the start of the automatic process of electrophoresis</u>



Picture 2. 10 – Start Electrophoresis Process

**h**) When running the machine for the samples reading, on the screen will appear "Machine Control" (see Picture 2.11), which illustrates the operational functions.

The image shows the correct placement of objects inside the machine.

The left column shows the status of the liquid levels.

At the center of the field called "Migration", check the values of voltage, the current migration and the remaining time for migration.

The "Machine Status" indicates the status of the machine.

The "List of services" allows the user to perform routine maintenance of the machine.

Aries (Rel.:Sep 17	2013 20:2	:39)	X-
File Funzionalita'	?		
Cruscotto Con	nandi		
Livelli			
Livelli Esterni	Stato		
Scarico Colorante Decolorante H2O Tampone	Vuoto Vuoto Vuoto Vuoto Vuoto	Migrazione Volts: 90 mA: 2 0 Tempo restante Migrazione: 00:00:00	
Livelli Interni	Stato		
Colorante Colorante Imbibizione	Pieno Pieno Pieno Pieno	Accesa	I A A
		Lista Servizi	
		N. Servizi Utente	
Valvole	Posi		
	255		
Controllo Maco	china [	Metodiche 🔀 Dati Odierni 🔍 Ricerca Esami 💽 Archivio	
Pronto		Sieroproteìne	26.09.2013 - 21:37:28

Picture 2. 11 – Control Machine

*i*) At the end of the electrophoretic process after the densitometric reading of the strip, you can see the traces going back to: "Current data" → "Edit traces" (see Picture 2.12)

😨 Aries (Rel	.:Mar 13	2012 22	2:20:29)					
File Features	?							
Manageme	ent serum l	nolder	Change	Tracks	Print			
Archive				6				
🕒 🗁 02	/02/2012	1		2		9		
	(1) Jam (2) Gari (3) D'Au	ies Cam ibaldi Gi	eron iuseppe Gabriele	C S	Current Meth Sieroprotein	od:	Job Data: 27/04/2012	^
	(4)	inunzio -	Janliele	F	atient N.:	Port serum N.: Slot N	.: Patient Type:	
	(5)			1		1 1	Human	<ul> <li>Image: A set of the set of the</li></ul>
	(6)			5	urname:		Name:	interest of
	(7)			100	James		Cameron	
	(8)						A	
Valori Frazioni						Grafico		8
Siero umar	10 da 5					×		
Frazioni	%	Fat	Int.Rif.%	g/dL	Int.Rif.g/o	L		
Albumina	56,9	1,00	52,0-65,0	0,00	3,12-5,2	0		
Alfa1	2,7	1,00	2,5-5,0	0,00	0,18-0,4	0		
Alfa2	11,5	1,00	7,0-10,0	0,00	0,42-0,8	0		
Beta	14,4	1,00	10,0-15,0	0,00	0.84.1.6			
Gamma	14,5	1,00	14,0-20,0	0,00	0,04-1,0			
Ratio A/G			Proteine Tot	ali g/d	Ē			
1,32 (Rif.: 1	,00-2,10)	6	0,00 (Rif.: 5,0	0-8,50)				
Control Ma	achine	🔼 Metho	ods 🛛 🛃 Toda	ay's da	ta 🛛 💽 Searcl	Exams SArchive		
Ready					Sier	proteine	29.04.2012	- 10:45:19

Picture 2. 12 – Change Tracks

On the "Edit Traces" screen both the graph and the relative values of the fractions for each sample are illustrated. At the top left there is also the list of the analyzed samples.

On the same screen, as in the previous one, you can enter or edit data of each sample.

On the graph window it is possible to insert any necessary changes, such as: inserting, moving and deleting minimum, changing the base line, and zooming by clicking on the squared button on top right of the graph window (see Picture 2.13)



Picture 2. 13 - enlarge image

**NOTE**: For more details on the changes effectively the tracks, you can refer to Section 4.2 "Edit Traces".

*j)* Once made final changes of the histograms can be traced to the press of selecting the "Print" window at the top, next to the "Change Tracks" (see Picture 2.14)

Featur	es ?							
Manag	ement serum holder	Chan	ige Tracks	Print				
		9 🥥						
ų.	Data Acc.	Cod. Acc.	Metodica	Cognome	Nome	Eta'	Sesso	Tipo Paziente
1	03/02/2012		Sieroproteine			0	M	Umano
2	03/02/2012		Sieroproteine			0	М	Umano
3	03/02/2012		Sieroproteine			0	М	Umano
4	03/02/2012		Sieroproteine			0	М	Umano
5	03/02/2012		Sieroproteine			0	M	Umano
6	03/02/2012		Sieroproteine			0	М	Umano
7	03/02/2012		Sieroproteine			0	М	Umano
3	03/02/2012		Sieroproteine			0	М	Umano
9	03/02/2012		Sieroproteine			0	M	Umano
10	03/02/2012		Sieroproteine			0	М	Umano
11	03/02/2012		Sieroproteine			0	М	Umano
12	03/02/2012		Sieroproteine			0	М	Umano
13	03/02/2012		Sieroproteine			0	М	Umano
14	03/02/2012		Sieroproteine			0	М	Umano
15	03/02/2012		Sieroproteine			0	М	Umano
16	03/02/2012		Sieroproteine			0	M	Umano
Contro	l Machine 🛛 🚺 🛛	1ethods 🛛 🛃 1	oday's data 🛛 🧧	Search Exams	Archive			
-		Neg 1			August 11			

**Pictrure 2. 14 – Print Reports** 

You can print the report by clicking the right mouse button on the row and you will see a check mark appearing on the left of the numbered line, otherwise you can choose to select all the read samples by clicking on the symbol with a tick (see Picture 2.15)

		2					
Ν.	Data Acc. Cod	Acc. Metodica	Cognome	Nome	Eta'	Sesso	Tipo Paziente
√1	03/02/2012	Sieroproteine			0	М	Umano
√2	03/02/2012	Sieroproteìne			0	М	Umano
√3	03/02/2012	Sieroproteine			0	М	Umano
<b>√</b> 4	03/02/2012	Sieroproteine			0	М	Umano
√ 5	03/02/2012	Sieroproteine			0	М	Umano
<b>√</b> 6	03/02/2012	Sieroproteine			0	М	Umano
√7	03/02/2012	Sieroproteine			0	М	Umano
√8	03/02/2012	Sieroproteine			0	М	Umano

Picture 2. 15 – Print all selected samples

**k**) Selected the samples to be printed by clicking on the yellow Printing icon (5), than a Print Preview window will appear.



**Picture 2. 16 – Print Preview** 

**I)** Print the report clicking on the report icon

# 3 Methods

🗾 Control Machine 🛛 🚺 Methods 🛛 🔀 Today's data 🛛 🔍 Search Exams 🛛 💽 Archive

In this section of the program the user can choose the method for the analysis of samples. Clicking on the menu "*Methods*" to access to "*Configuration Parameters*", "Definition Fractions" and "Definition Comments" these windows allow the following operations:

- Choosing a method from the default ones by selecting it with the mouse, in the window "List of methods" section on the left side of the screen and setting it as *current method* for analysis of samples using the yellow button "*Set the current method of work*" .
   This operation updates the second field of the status bar with the current descriptions of the methodology. It is also possible to set *Configuration Parameters* for each method in the default system on the right part of the screen.
- Defining a group of fractions for the current method, both concentration value and absolute value, for predefined patients in the system.
- Managing comments for the current method.

# 3.1 Configuration Parameter

Configuration Parameters	Definition Fractions	Definition Comments	



In the column "List methods" on the left side, you can see the different types of methods that can be performed (Picture 3.1).

	Configuration Parameter
List me	ethods
N.	Method
1	Hemoglobin
2	Lipoproteins
3	Urinary protein
√4	Serum proteins

Picture 3. 1 – List Method

To select a method, just clicking on it and a yellow check mark icon will light up . Clicking on this icon it will display a message saying that the method was successfully selected and that the program will restart according to the chosen method.

On the "Configuration Parameters" screen, on the right in front of blue window you can make all the possible changes of parameters:

• Voltage: you can vary the tension moving the vertical arrows, or you can vary the current deselecting first in the indicator box next to "volts"

• **Applied Time** (sec-min): you can vary the migration times, drying times of the strip, times of imbibition, staining and destaining (see Picture 3.2)

Picture 3. 2 – Applied Time

- *Outside normality Symbol*: in the printing reports you can decide whether to highlight in red any abnormal results or to use symbols that indicate out of range.
- Unit of measure concentration: you can choose the unit of the desired concentration unit.
- *Representation*: you can choose to put in the report the average absolute values , normal values of concentration, concentration, reference of concentration by selecting the respective check box with the mouse.
- *Viewing and Intensity color (1-255) Migration*: it includes the ability to put together the histogram and the real electrophoretic trace of the patient. You can change the width and the position of the trace (see Picture 3.3)



Picture 3. 3 – Visualizing option of the trace in print

- Areas ratio: Shows the areas relation and allows the inclusion of any comments.
- *Print*: it allows you to insert the name of the method on the report before printing.

# **3.2Definition Fractions**

File Features ?			
Configuration Parameters	Definition Fractions	Definition Comments	

Create fractions group connected to the method, in particular choose the sample category between human and veterinary (Picture 3.4)

s rea	atures /									
Co	nfiguration Parameters	Definitio	n Fraction	s 📄 Defir	nition Comme	ents				
		2								
Fract	ions Value	Sample Ca	tegories				Su	m of frac	ctions (eg i	2 +3 +4 +5)
) Co	ncentration (g/dL)	Patient type	<b>::</b>	Human			✓ Divi	idend:	1	
🕑 Ab	solute (%)	fractions gr	oups:	Human	Serum fro	m 5	- Divi	isor:	2+3+4+5	
N.	Fraction name	_	Men Min.	Max,	Wom Min.	ian Max.	Baby Min.	Max.	-	$\bigcirc$
N.	Fractions	Factor	Min.	Max.	Min.	Max.	Min.	Max	Grid	
1	Albumin	1.00	52.0	65.0	52.0	65.0	52.0	65.0	D No	
2	Alpha1	1.00	2.0	5.0	2.0	5.0	2.0	5.0	) No	
3	Alpha2	1.00	7.0	13.0	7.0	13.0	7.0	13.0	D No	× ×
4	Beta	1.00	8.0	15.0	8.0	15.0	8.0	15.0	D No	
5	Gamma	1.00	12.0	21.0	12.0	21.0	12.0	21.0	D No	
	Ratio A/G		1.00	2.10	1.00	2.10	1.00	2.10	,	

**Picture 3. 4 – Definition Fraction** 

In the Picture 3.4 two buttons are highlighted in yellow 🕃 🥃 , they allow you to delete or add a group of fractions. SAIO machine usually is delivered and installed with all the parameters of

the methods already set correctly, but in case you want to make any changes to the methods, please follow the below procedure.

To add a group of fractions click the icon and enter a new group of fractions from a minimum of 1 to a maximum of 12 and confirm with OK. A message will appear asking to enter the name of the new group of fractions (see Picture 3.5).

Aries	Aries 🛛 🔀
Do you want to add a new group of fractions? (Min.1-Max.12)	Enter the name of the new group of fractions
OK Cancel	OK Cancel

Picture 3. 5 – Group of Fraction

To change the normal minimum and maximum values of the fractions, select with the mouse the fraction and on the left will appear a checkmark and simultaneously also the values of fraction in the fields above (see Picture 3.6)

	Fraction name Albumin		Men		Woman		Baby		
N. 1			Min. 52.0	Max. 65.0	Min. 52.0	Max. 65.0	Min. 51.0	Max. 65.0	
N.	Fractions	Factor	Min.	Max.	Min.	Max.	Min.	Max.	Grid
1	Albumin	1.00	52.0	65.0	52.0	65.0	52.0	65.0	No
2	Alpha1	1.00	2.0	5.0	2.0	5.0	2.0	5.0	No
3	Alpha2	1.00	7.0	13.0	7.0	13.0	7.0	13.0	No
4	Beta	1.00	8.0	15.0	8.0	15.0	8.0	15.0	No
5	Gamma	1.00	12.0	21.0	12.0	21.0	12.0	21.0	No
-	Ratio A/G		1.00	2.10	1.00	2.10	1.00	2.10	

Picture 3. 6 – Normal Values Modification

Change the value of the fraction in the appropriate field, a checkmark will appear in yellow. Clicking this button the value will update. Save the changes by clicking on the saving button.(see Figure 3.7)
Frac	tions Value	Sample Ca	tegories				Su	m of frac	tions (eg 2	+3 +4 -
<ul> <li>Concentration (g/dL)</li> <li>Absolute (%)</li> </ul>		Patient type		Human 👻 Human Serum from 5 👻				idend:	1 2+3+4+5	
		fractions gr	oups:					isor:		
N.	Fraction name	_	Men Min.	Max.	Wom Min.	an Max.	Baby Min.	Max.		
N.	Fractions	Factor	Min.	Max.	Min.	Max.	Min.	Max.	Grid	REF
1	Albumin	1.00	52.00	65.00	52.00	65.00	51.00	65.00	No	
2	Alpha1	1.00	2.0	5.0	2.0	5.0	2.0	5.0	No	
3	Alpha2	1.00	7.0	13.0	7.0	13.0	7.0	13.0	No	
4	Beta	1.00	8.0	15.0	8.0	15.0	8.0	15.0	No	
5	Gamma	1.00	12.0	21.0	12.0	21.0	12.0	21.0	No	
	Ratio A/G		1.00	2.10	1.00	2.10	1.00	2.10	Pi	

Picture 3.7 – Save the updated fraction

An important parameter is the **factor**, placed in the first right column containing the names of the fractions. The factor can increase or decrease the numeric read value for each fraction. This operation is performed essentially in the calibration phase of the machine by using a known sample or a control sample.

Fractions Values: values can be expressed in percentage or in g/dL

**Samples Category**: the types of patients can be chosen in the adjacent field (human or animal), or it is possible to choose the group fractions (see Picture 3.8)

Sample Categories			Sample Categorie	S	
Patient type:	Human	~	Patient type:	Human	~
fractions groups:	Dog Horse		fractions groups:	Human Serum from 5	~
Mei	Cat Cow Human		Me	Human Serum from 5 n Human serum from 6	

Picture 3. 8 – Sample Category

**Sum of Fractions**: it determines the fractions on the basis of the calculation of the ratio A /G, in particular to the dividend is assigned the fraction Albumin and to the divisor the Globulin.

You can color the area of fractions of report by clicking on the button which allows two operations (see Figure 3.9):

• choose a color for the selected fraction on the list;

• select a grid for the selected fraction on the list;

Co	nfiguration Par	ameters 📄 🗎 Definition Fract	ions 🛛 🗎 Defi	nition Comme	ents				
		Select the color of the f 🤶							
Frac	tions Value	Colori di base:	_			Su	n of frac	tions (eg 2	+3 +4
0 Co	ncentratio		🔲 uman		1	🖌 Divi	dend:	1	
⊙ Ab	solute (%)		📕 luman	Serum fro	m 5	🖌 Divi	sor:	2+3+4+5	
				Wom	an	Baby			
N.	Fraction		lax.	Min.	Max.	Min.	Max.		
3	Alpha2	Calari norosu slimesti	3.0	7.0	13.0	7.0	13.0		
N.	Fractions		Max.	Min.	Max.	Min.	Max.	Grid	
1	Albumin		65.00	52.00	65.00	51.00	65.00	No	6
2	Alpha1		5.0	2.0	5.0	2.0	5.0	No	
13	Alpha2	Definite at a start and a strengtherest as	13.0	7.0	13.0	7.0	13.0	No	
4	Beta	Definisci colori personalizzati >>	15.0	8.0	15.0	8.0	15.0	No	
5	Gamma	OK Annulla	21.0	12.0	21.0	12.0	21.0	No	C
	Ratio A/G		2.10	1.00	2.10	1.00	2.10		

Picture 3.9 – Fractions Color

### **3.3Definition Comments**



In *Definition Comments* is possible to insert default comments that can be used in case of pathologies reported by densitometric reading of the traces.

To enter a new comment, select the blank field at the top side and then add to it to the comment list

by clicking on the icon **O**. A confirmation message will appear in order to validate the successful saving the list of comments of the current method.

To remove comments select the comment line in the list and click the icon

File Features	?		27-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1
Configurat	ion Parameters	Definition Fractions	Definition Comments
	0		
Comment:			
N. 0	Comments		
1 A	nomalous patter	n in zone beta	
2 B	isalbuminemia		
3 C	hronic Infection		
4 Ir	nflammation		
5 P	resence of mono	clonal pattern	

Picture 3. 10 – Definition Comments

## 4 Today's Data

2	Control Machine	Methods	🔀 Today's data 📗	Search Exams	C Archive	
-		The second se				

Picture 4. 1 – Today's Data Window

The Today's Data section contains all the information related to the samples acquired during the day. It is possible to start the process of electrophoresis, to access and to modify the results, to print out reports.

Click on the menu "*Today's data*" to access the "*Management Serum holder*" window, "*Change Traces*" and "*Print*" that allow you to perform, respectively, the following operations:

- If needed, insert a new serum holder using the "*Add*" new serum holder to the current method key and enter patient data in each empty slot of serum holder. The "Previous serum holder details" and "*Next serum holder details*" keys allow scrolling among the patients of the day. If no additional sample has to be inserted in serum holder, you can start the electrophoretic process pressing the button is "*Start machine the samples reading*"
- It offers a general overview of the samples collected during the day by visualizing the patient data, which are modifiable by the user; the image of migration and the graph of the curve, which can be changed by the user with interaction of the mouse with the graph; the numerical data of the fractions, which can be automatically updated according to the curve changes. In this screen there is also a small window, underneath the fractions, dedicated to comments where it is possible to enter a new comment, or edit an existing one. To access the details of a new sample, use "*Previous patient details*" and "*Next patient details*" keys



or click directly on a control sample from the "Archive" on the left side of the window.

• Print one or more samples reports selecting them from the list with the right button of the mouse or selecting all of them pressing the "*Select Patient*" and "*Print patient*" keys.

### 4.1 Managment Serum Holder



The operations that can be made in this screen are: inserting a new serum holder of the day; refilling each slot with patient data and starting the process of reading samples. The application can be configured with a 16 slots serum holder (See Figure 4.2).



Picture 4. 2 – Serum Holder 16 samples

Please find below description for each particular field on this screen:

Field Name	Description
	The "Previous serum holder details" and "Next serum holder
	details" are used to scroll the list of serum holder in the "Archive".
	The "Add" key used to insert a new serum holder of the day. It is
Ŧ	possible to insert only one serum holder before starting the reading
	process

	The "Sava patient and note serum holder" key to save any changes
	of the patient data or to add a note to the serum holder.
	The "Clean" button reset the parameters of the screen.
	The "Start machine for the sample reading" button runs the
	electrophoresis process and the sample reading.
Archive	This is the tree control on the left that collect the serum holders of
Archivio Archivio Portasieri Sieroproteine 01/02/2012 20:17 (5)	the day.
Serum Holder	This control is located in the central part on the left side of the
	screen, it gathers 24/48 slots for the patient samples.
Portasiero	Each slot of the serum holder can assume one of the following
	<ul> <li>WHITE: empty slot. Select with the mouse an empty slot to clear it and to disable the data of the patient. Then new data can be entered.</li> </ul>
	Salo 🛛 🔀
	Enter the number of patients to be added to port serum ${f max} \ 16$
	8
	OK Cancel
	Picture 4.3 - Number Patient
	• YELLOW: free slot. It is possible to start inserting one or more patients in the serum holder. By double-clicking on the slot a new screen will appear with the data of the

patients that can be added.

The right side of the screen relating to patient data is enabled for entering patient data.

- GREEN: indicates the slot selected by the mouse with the corresponding data of patients included (or to be included).
- RED: slot occupied by a patient. Selecting a slot with the left button of the mouse it is possible to update patient data in the slot. Selecting an occupied slot with the right button of the mouse, a drop-down menu will appear and it will be possible to remove patient from the slot clicking "Delete Patient".
- BLACK: Invalid slot. Selecting a non-valid slot with the left button of the mouse it is possible to clear and disable the data of the patient to the right of the screen. Selecting a non-valid slot with the right button of the mouse it is possible open a drop-down menu with which you can select the "Restore Patient" to restore the patient in the deleted slot.

The serum holder contains also the following labels:

• Status: indicates the status of current serum holder.

The possible values: EMPTYING , PREPARATION, READING, PROCESSED and CANCELLED

- N. serum holder: indicates the number of serum holders.
- Note: additional note that can be associated to the serum holder.
- Number of Slots: The number of selected slots.
- Remainder: indicates the number of strips still present in the Smart Card

Slot n. 1	To facilitate the scrolling of the serum holder and consequently the patient detail linked to the selected slot.
Patient Data	It is situated on the right side of the screen and it is used to insert, to modify the data of the patient.

Table 1: Management Serum Holder

## 4.2 Change Tracks

File Features ?		
📄 Management serum holder	Change Track	s 📄 Print
Archive		
🛅 Tracks Archive today		
		Current Method:
		Sieroproteine

In the window below it is possible to run the following operations: changing the patient, attaching a comment to the patient, changing the image of migration, changing the factor, changing the graph of the curve and the numerical data of the fractions with the interaction of the mouse with graph (see Picture 4.4).

😨 Aries (Rel	.:Mar 13	2012 22	: 20: 29)								
File Features	?										
Manageme	ent serum l	nolder	Change	Tracks	5 Print						
Archive	/02/2012 (1) Jam	es Cam	eron	-	Current Meti	Ø				Job Data:	~
	(2) Gari	baldi Gi	useppe Sabriola		Sieroprotein	е				27/04/2012	
	(3) D Ar (4)	inunzio (	apriele		Patient N.:	Port	serum N.	: Slot N.:	Patient 7	Гуре:	
	(5)					1		1	Human		× =
	(6)				Surname:				Name:		
	(8)			~	James				Camero	on	
Valori Frazioni						R	Grafico				٥
Siero umar	io da 5					~					
Frazioni	%	Fat	Int.Rif.%	g/dl	Int.Rif.g/	dL					
Albumina	56,9	1,00	52,0-65,0	0,0	3,12-5,	20		A			
Alfa1	2,7	1,00	2,5-5,0	0,0	0,18-0,	40					
Alfa2	11,5	1,00	7,0-10,0	0,0	0,42-0,	80		1			
Beta	14,4	1,00	10,0-15,0	0,0	0,60-1,	20					
Ratio A/G 1,32 (Rif.: 1	,00-2,10)	.,	Proteine Tot ),00 (Rif.: 5,0(	ali g/a	1L				<u> </u>		
Control Ma	achine	🔼 Metho	ds 🔰 🛃 Toda	ay's da	ata 🛛 💽 Searc	h Exan	ns 🚺 🖸 A	rchive			
Ready					Sier	oprot	eine			29.04.201	2 - 10:45:19

Picture 4. 4 – Change Tracks Window

**Factor**: The factor is a parameter located in the first column to the right side of the names of the fractions, it is used to increase or decrease the numerical read value of each fraction (Picture 4.5)

Valori Frazioni					
		$\frown$			*
Frazioni	%	Fat	Int.Rif.%	g/dL	Int.Rif.g/dL
Albumina	56,9	1,00	52,0 - 65,0	0,00	3,12-5,20
Alfa1	2,7	1,00	2,5 - 5,0	0,00	0,18-0,40
Alfa2	11,5	1,00	7,0 - 10,0	0,00	0,42-0,80
Beta	14,4	1,00	10,0 - 15,0	0,00	0,60-1,20
Gamma	14,5	1,00	14,0 - 20,0	0,00	0,84-1,60
		$\bigcirc$			

**Picture 4. 5 – Factor Parameter** 

To modify the Factor of the desired fraction it is necessary to click on it with the left button of the mouse and the following window will display asking to insert the number between 0.7 and 1.5(see Picture 4.6)

Changes factor Pattern 1					
insert a number between 0.70 and 1.50					
1.00					
	OK Cancel				

Picture 4. 6 - Modify Pattern Factor

Please find below description for each particular field on this screen:

Field Name	Description			
	The "Previous patient details" and the "Following patient details" buttons scroll the list of patient in the "Archive".			
	Aries Inserire il nome o l'indirizzo IP della macchina dove esportare i risultati dei pazienti Decelhoss OK Cancel Picture 4. 7 - Export Insert name machine IP address			
	The "Export patient result" key exports to an external database the results of the reading of a selected patient in the tree-control on the left side of the screen. By clicking on the button a screen will display to show the machine where to export the selected patient data. Usually it is necessary insert the name and the IP address.			

( <del>+</del> )	The "Add a new fraction group to the current method" button will
	add a new fraction group to the current method of work accordingly
	to data inserted in "Fraction Values"
	The "Save patient and graph" save a data modification.
	The "Clean" button reset the screen parameters.
A	The "Associate a comment to a patient" associate a comment to the
$\sim$	"comments" list of the patient.
	The "Disassociate patient comment" disassociates a comment from
	a patient.
	The "Print patient report" key visualizes the print preview of the
	report
	The "Changeable notiont date" key can modify the nationt date
$\bigcirc$	Once pressing this button, it will not be possible to change the detail
	once pressing this button, it will not be possible to change the detail
$\mathbf{C}$	The "No changeable patient data" key indicates that it is not possible
	to change the data, since the report has been already delivered. It is
	only possible to read the data.
Archive	This is the tree-command the collect all the patient of the day.
Fraction Values	
	This command, located in the middle left part of the screen, collects
	the fraction values into a numerical format. It will be automatically
	entered when you modify the graphs of the reading.
	The fraction list becomes more editable with the "Factor" column
	that can change the fraction value.
	The list of the fractions will be editable for the "Fractions,

	"Int.Rif.%", "Int.Rif", "Ratio A/G" and "Total Proteins" only					
	when the number of the fractions of the graph will be different from					
	the default number of the current method for the selected patient.					
	Otherwise you can add a new fraction group for the current method					
	clicking the button: "Add a new fraction group to the current					
	method".					
	It is not possible to edit the fraction list and it is fully completed in					
	all its fields. The combination of the fraction groups will have the					
	name of the default fraction groups with the current methodology.					
Patient Data	To insert, modify the patient data.					
	·					

Table 2: Change Tracks

## 4.2.1 Graph



Picture 4. 8 – Graph

The chart associated with the patient is the result of the reading performed by the scanning process of the samples. The graph displays the minimum values through the vertical lines.

These minimum levels can be moved using the mouse with a drag-and-drop operation on the graph. The minimum lines can be deleted or added on the graph by double clicking the mouse.

It is possible to open a drop down menu with the right mouse button on the graph, in order to make relevant changes. (see Picture 4.9):



Picture 4.9 – Drop Down Menu Graph

• *Color Patterns*: the patterns of the graph can have the color set up in the definition of the fractions for the current method of work and can take the dotted curve, or be transparent (see Picture 4.10)



Picture 4. 10 – Color Pattern

- *Subtract Baseline*: to modify the baseline of the graph
- *Minimum Compute* : it allows automatic setting of the minimum value on the graph
- *Export*: to export the fractions values of the graph to a file. CSV or. TXT

- Gamma Correction: it is confidential and may only be changed by the expert staff.
- *Zoom slider*: zoom the graph for a better consultation; enlarges the area in which you placed the mouse on the graph. (See Picture 4.11)



Picture 4. 11 – Zoom Sider

- *Displaying the grid of the fraction values:* a banner allows the flow of comments related to the graph.
- *List*: the possibility to enter a legend inside the chart (see Picture 4.12)



Picture 4. 12 – Insert Legend

• Zooming of the Image: enlarge the entire image of the graph clicking on "zoom" on the right of the graph window



Picture 4. 13 – Zooming Of the Image

### 4.3 Print



**Picture 4. 14 – Print Window** 

In this screen you can select and print one or more patients of the day or export or capture patient data.

File Funzional	lita' ?								
📄 Managemer	nt serum holder	Change	Tracks 📄 Print	1					
		00	<b>)</b>						
N.	Date Acc.	Acc. Code	Method	Surname	Name	Age	Sex	Patient Tiente	^
<li>✓1</li>	27/09/2009		Sieroproteine	Cognome1	Nome1	0	M	Human	
3	27/09/2009		Sieroproteine	Cognome3	Nome3	0	Μ	Human	
									~

Picture 4. 15 – Printing Mode

Please find below description for each particular field on this screen.

Field Name	Description
	- coord Protein

	"Previous patient details" and "Following patient details" keys scroll the patient list.
	"Select patient" key in order to select the patients. Clicking twice you deselect all the patients of the list.
	<ul> <li>"Export patient results" key exports to an external database the reading results of the selected patients. Clicking the key you will see the following screen:</li> <li>Aries Inserire il nome o l'indirizzo IP della macchina dove esportare i risultati dei pazienti Inserire il nome o l'indirizzo IP della macchina dove esportare i risultati dei pazienti Insertre 4. 16 - Export – Insert name of machine and IP address In order to show where to export the patient data, usually you need to enter the name and the IP address of the machine</li></ul>
	"Clean" key to reseat the screen parameters.
9	The "Print patient" button displays the print preview with the selected reports.
	"Changeable patient data" to put in read only the selected patient data.
Patient List	The list contains all the patient of the day. To print one of the patient of the list click with the right button of the mouse on it.

Table 3: Print

#### 4.3.1 Print Preview

With this operation it is possible to print reports of patients.

On the header of the report there is the name of the laboratory where the tests have been performed "Personal Data/Laboratory", in the central part of the report appear some patient data, the graph and the image of the migration, on the footer the comments and the signature of the analyst. Please find below description for each particular field on this screen (See Picture 4.17)



Picture 4. 17- Print Preview

Please find below description for each particular field on this screen.

Field Name	Description
Close	
	"Close" key to close the "Print preview"
Print	
	"Print" key to start the print of the patient reports.
[<< <> >> >>]	These keys scroll the pages of the report print previews.
Goto	
	"Goto" key to access to a specific print preview page.
65% 🛩	
	The combo is used to set the zoom percentage of the page before
	printing. Any modification will appear only on the monitor.

Table 4: Print Preview

## 5 Control Machine

🗾 Control Machine 🛛 🖸 Methods 🛛 🖉 Today's data 🔄 Search Exams 🛛 💽 Archive

This section of the program allows you to connect SAIO (electronics and software) to the physical machine (the robotics, sensors) and to the electrophoresis operations. Clicking on the menu " Control Machine" to access the "Dashboard Control" that allows you to: turn on and off the machine; to modify the voltage of the chamber during a migration; to run a selected process from the list of the services, to pause and resume a process already running. If an error occurs it is possible to go to a next step or stop the process permanently. The error of the process will be outlined in a separate control at the bottom of the screen.

The window also has status indicators (full or empty) for the levels of the machine that are classified as external levels such as Staining, Destaining, H2O, buffer, buffer HB and waste and as internal levels such as staining, destaining, Imbibition, buffer in Migration Chamber.

"Turn on " and "Turn off " keys to switch on/off the machine for the electrophoretic process or to perform other operations selected by the user.
"Process Pause" and "Restart Process" keys to pause or to restart the process.
"Start Process" key to start a process selected by the command "Process List"
"Continue Process" key can be clicked when the process stops because an error occurred. Pressing this button the error is deleted and the process goes on.
"Stop the process" key can be clicked when a blocking error occurs. In this case the button stops the process.



"Alarm" key lights when the process ends correctly or in case of error. Clicking on this button the alarm turns off.

The Control Machine window appears as shown in the Picture 5.1. On the top there is a power off button O that means the machine is turned off because the mechanics is not supplied. Click on button to switch on the machine.(see Picture 5.1).



Picture 5. 1 – Control Machine

- The picture shows the correct placement of objects inside the machine.
- The left column shows the status of the inside and outside liquid levels. The liquid levels are highlighted in red or in green on the basis on their status
- In the center of the field called "*Migration*" you can find the values of voltage, the current migration and the remaining time of the migration.
- The "State Machine" is the status of the machine.
- The "Services list" in order toper form routine maintenance of the machine.

#### Maintenance service button

The "Service Maintenance" button opens a drop-down menu (see Picture 5.2)



Picture 5. 2 – service Maintenance

Two options can be selected: the first allows the final user to load the cleaning process list (see Picture 5.3)

List Servi	ices	
Ν.	User services	<b>^</b>
1	RESET	
2	LOAD SERUM BUFFER IN CHAMBER 1	
3	LOAD SERUM BUFFER IN CHAMBER 2	
4	LOAD SERUM BUFFER IN IMBIBITION	
5	LOAD HEMOGLOBIN BUFFER IN CHAMBER 1	
	LOAD HEMOGLORIN RHEEED IN CHAMRED 2	<b>`</b>
<		>

Picture 5. 3 – User services

The second option allows you to perform maintenance tasks that are intended only for qualified.

A password is required in order to proceed the operation and as displayed in the following window:

Immissione Passwor	d	
Immettere la password o	dell'operatore	
	ОК	Cancel

Picture 5. 4 – Password Maintenance

# 6 Cancellation



SAIO keeps stored the database and authorize the cancellation of samples or whole serum holders from the archive. Clicking on the menu "Cancellation" you access the windows of "Delete Exams" and "Delete Serum holder" and proceed the following operations:

- Clicking on the "Search patient in the archive" new window will display where the user can enter the time interval for the research and he decide the type of desired search, for example by last name, first name, doctor, sector, patient type, sex or method. After selecting the desired type, the research will start by clicking with the mouse on the key "Search Start". The result appears in the list. Delete one or more samples from the list by selecting them using the right button of the mouse or selecting them all clicking "Select Patient" and then the "Delete Patients".
- Clicking on the key "*Search serum holder in the archive*" the last entered serum holder will be searched. The result will be displayed in the "*Cancellation*" in the top left part of the screen. In order to delete the serum holder press the "*Delete serum holder*".

le Featur	es ?				
📄 Delete	e Exams 🛛 📄 Delete serum holder				
<u>۔</u> ۷.	Date Acc. Acc. Code Method	Surname	Name	Age Sex	Patient Type
				0.42	50

Picture 6. 1 – Delete Exams

## 6.1 Cancella Esami

Delete Exams	Delete serum holder	
--------------	---------------------	--

Clicking the yellow key ("Search patient in the archive") vou can find the patients in the database and then delete them. During the search the "Search Filter" window is displayed (see Picture 6.2). The search result is displayed in the list. Delete one or more samples from the list by selecting them using the right button of the mouse or selecting them all clicking "Select Patient" and then the "Delete Patients". (see Picture 6.3).

File Features ?							
Delete Exams	💱 Filter Search						
	Setting Filters						
N. Date	ti						
	From:	To:					
	18.02.2014	18.02.2014	~				
	Surname:	Name:					
	Doctor:	Department:					
	Patient Type:		Sex:				
		*	~				
	Method:						
	Serum proteins		~				

**Picture 6. 2 – Setting Filters** 

File Feat	ures ?							
Dele	e <b>te Exams 📄</b> Dele	ete serum holder						
		3						
N.	Data Acc.	Cod. Acc.	Metodica	Cognome	Nome	Eta'	Sesso	Tipo Paziente
1	02-02-2012		Sieroproteine			0	М	Umano
2	02-02-2012		Sieroproteine			0	М	Umano
3	02-02-2012		Sieroproteine			0	M	Umano
4	02-02-2012		Sieroproteine			0	М	Umano
5	02-02-2012		Sieroproteine			0	M	Umano
6	02-02-2012		Sieroproteine			0	M	Umano
7	02-02-2012		Sieroproteine			0	M	Umano
8	02-02-2012		Sieroproteine			0	М	Umano
9	02-02-2012		Sieroproteine			0	М	Umano

#### Picture 6. 3 – Filters Delete Patients

	"Select Patient" key allows selecting all the patient of the list. Double clicking deselects them all.
$\Theta$	"Delete Patients" key deletes the patient from the database.
	The " <i>Clean</i> " key resets the screen parameters.
S	<i>"Search patient in the archive"</i> key displays the window for the search of the patient in the archive.
Lista Pazienti	The list contains the patients of the day. To print a patient from the list click on it with the right button of the mouse. The selected patient will be marked with the check, click again to deselect it.

Table – delete Exams Patient

Delete Exams	Delete serum holder
Archive	
Archive seru	ms holder
The second s	
😑 🔄 Sieroprote	eine
Sieroprot	eine 012

Picture 6. 4 – Delete Serum Holder

Clicking on the "Search serum holder in the archive" button the last entered serum holder will be searched and then you can delete it. (See Picture 6.4).

Delete Exams	Delete serum holder							
Archive								
Archive serums holder Sieroproteine 28/03/2012 19:11 (6)								
		Current Method: Sieroproteine					Job Data: 28/03/2012	
		Patient N.:	Port serum N.:	Slot N.:	Patient Ty	/pe:		
		41	6	1	Human			
		Surname:			Name:			
Port serum		1			14 -			
	State	Code Accep	tance:	Tota	I Protein:	Age:	39. 72	Sex:
	Done						4	Male
	N Dort sorum	Date Accept	ance:		Birth Date	0		
	06	28.03.2012		*	29.04.	2012		
		Data Lists:						
	Notes							
	28/03/2012 - 19:11:54							

Picture 6.5 – Delete Serum Holder in the Archive

# 7 Archive

		100	
Control Machine	Methods	🔀 Today's data	🔄 Archive

SAIO machine allows searching in the database both the samples that must be tested and the ones already tested

### 7.1 Advanced Search

Current Met	hod:			Job Data:	
Serum prote	ins			//	
Patient N.:	Port serum N.:	Slot N.:	Patient T	ype:	
0	0	0			
Surname:			Name:		
	Current Metl Serum prote Patient N.: 0 Surname:	Current Method: Serum proteins Patient N.: Port serum N.: 0 0 Surname:	Current Method: Serum proteins Patient N.: Port serum N.: Slot N.: 0 0 0 Surname:	Current Method: Serum proteins Patient N.: Port serum N.: Slot N.: Patient T 0 0 0 Surname: Name:	Current Method:       Job Data:         Serum proteins      //         Patient N.:       Port serum N.: Slot N.:       Patient Type:         0       0       0         Surname:       Name:

Picture 7. 1- Advanced Search

If clicking "Search patient in the archive" button a new window will display where the user can enter the time interval for the research and he decide the type of desired search, for example by last name, first name, doctor, sector, patient type, sex or method. (See Picture 7.2)

Advanced Sea	r 😵 Filter Search		
Archive	Setting Filters		
Exams Archive			
	From:	To:	
	18.02.2014	18.02.2014	~
	Surname:	Name:	
Fractions Value	Doctor:	Department:	
Tractions	Patient Type:	Sex:	
Fractions		<u>×</u>	
	Method:		
	Serum proteins	~	

Picture 7. 2 – Setting Filters

After selecting the desired type, the research will start by clicking with the mouse on the button "Start Search". The result will be displayed in the "Archive".

For more information about curves and patients, it is necessary select an item from the "Archive".

### 7.2 Exam Archive Print



Print one or more reports clicking on the right button of the mouse or selecting them through "*Select Patient*" button and press the "*Print Patient*" (see Picture 7.3). In this screen it is possible to print, export and block one or more samples selecting them from the list with the mouse clicking or select them with the "*Select Patient*" button.

File Featu	ires ?								
Advar	nced Search 🛛 📔	Print							
Ν.	Date Acc.	Acc. Cod	e Method	Surname	Name	Age	Sex	Patient Type	-
1	17/09/2012		Sieroproteine			0	М	Umano	
2	17/09/2012		Sieroproteine			0	М	Umano	
3	17/09/2012		Sieroproteine			0	М	Umano	
4	17/09/2012		Sieroproteine			0	М	Umano	
5	17/09/2012		Sieroproteine			0	M	Umano	
6	17/09/2012		Sieroproteine			0	М	Umano	
7	17/09/2012		Sieroproteine			0	M	Umano	
8	17/09/2012		Sieroproteine			0	М	Umano	
1	08/08/2012		Sieroproteine			0	M	Umano	
2	08/08/2012		Sieroproteine			0	М	Umano	
3	08/08/2012		Sieroproteine			0	М	Umano	
4	08/08/2012		Sieroproteine			0	М	Umano	
5	08/08/2012		Sieroproteine			0	M	Umano	
6	08/08/2012		Sieroproteine			0	М	Umano	_
7	08/08/2012		Sieroproteine			0	M	Umano	
8	08/08/2012		Sieroproteine			0	М	Umano	
9	08/08/2012		Sieroproteine			0	М	Umano	
10	08/08/2012		Sieroproteine			0	М	Umano	
11	08/08/2012		Sieroproteine			0	M	Umano	
12	08/08/2012		Sieroproteine			0	М	Umano	
13	08/08/2012		Sieroproteine			0	M	Umano	
14	08/08/2012		Sieroproteine	and service and services	Sector States	0	М	Umano	
1	07/08/2012	4888	Sieroproteine	GIGLIOTTI	GERMANA	0	F	Umano	
		1000000000					10000	1.2.2.1	1920

Picture 7.3 – Print Patient

#### 7.2.1 Registries

It is possible to create a database with information regarding doctors, sectors, diseases and laboratories. Clicking on the "Registries" button you access the windows of "Laboratory", "Doctors", "Sectors", "Diseases" that allow the following operations:

- Management of headers for the report printings of the laboratories;
- Management of the patient's pathology;
- Management of the doctor's data;
- Management of the sectors.

### 7.2.2 Quality Control

It is possible to create a database with information regarding the quality control of the examined samples. Clicking on the "Quality Control" key you access the windows of "Configuration",

"Results" and "Graphs" that allow the following operations:

- -Management of the quality control of the examined samples;
- Report of Quality Control in tabular format.
- Report of the quality control in a graphical format

## 8 Registries

It is possible to create a database with all the information related to doctors, sectors, disease and laboratories.



### 8.1 Laboratory

In this screen it is possible enter all the information about the report.

The field concerning the description of the "Electrophoresis of serum proteins" is available only for the reading, and it is set up with the method name displayed on the *Method/Parameters configuration* screen. (see Picture 8.1)

File Features ?				
Laboratory Disea	ses	Doctor	Department	
			Name of the laboratory on printing	
Title 1 (Pos. X-Y):	135	40	GAMMA DIAGNOSTIC CENTER	
Title 2 (Pos. X):	175		MEDEA	
Title 3 (Pos. X):	120		Marmolar Street 20 - Tel/Fax 00408632905	
Title 4 (Pos. X):	125		QUADRO PROTEICO ELETTROFORETICO	
FirmaX:	500		The Doctor	Mario Rossi
Pos.X-Y Text Line	1		Pos.X.Li Text Line 2	Pos.X.Li Text Line 3
✔ 135;40 GAMMA D	IAGNO	STIC	175 MEDEA	120 Marmolar Street 20 - T

Picture 8. 1 – Changing the header

Please find below description for each particular field on this screen.

Field Name	Description
	"Previous laboratory detail" and "Following laboratory detail" keys
	scroll the list of the laboratories names
	"Set the laboratory as printing heading" key sets the selected header
	as a referential one.
Ð	"Add new laboratory" key adds a new heading into the list.
	"Remove laboratory" keys deletes the selected heading of the laboratory.
	"Save the laboratory" key saves the selected heading of the laboratory.

"Clean" key resets the screen parameters.
"Printer Properties" displays a drop-down menu:
Laboratory Diseases Doctor Department
Select Font
Font in Italic Name of the I Underline Font
Title 1 (Pos. X
Title 2 (Pos. X) MEDEA
This menu manages the type of font, of character
(italics/bolt/underlined), of color and the print preview.

Table 5: Laboratory

## 8.2 Diseases

Features ?	
Laboratory Diseases Doctor Department	
isease Description:	
Jiseases List	
Varicella Morbillo	

Picture 8. 2 – Disease

Please find below description for each particular field on this screen:

Field Name	Description
	"Previous diseases detail" and "Following diseases detail" keys scroll the list of diseases.
Ŧ	"Add new disease" key enters a new disease in the list.
$\Theta$	"Remove Disease" key deletes the selected disease.
	"Save disease" key saves the selected disease in the list.
	"Clean" key resets the parameters of the screen.

## 8.3 Doctor

In this screen it is possible to manage the doctors' data:

File Features ?			
Laboratory Diseases Doctor Department			
Surname:			
Name:			
Specialization:			
Surname	Name	Specialization	
Pinto	Alfredo	Pediatra	
Coppola	Pasquale	Urologo	

Picture 8. 3 – Doctor Windows

Please find below description for each particular field on this screen

Field Name	Description
	"Previous doctor detail" and "Following doctor detail" keys scroll the list of the doctors.
Ŧ	"Add new doctor" keys enters a new doctor in the list.
Θ	"Remove doctor" key deletes the selected doctor from the list.
	"Save doctor" key saves the doctor selected in the list.
	"Clean" key resets the parameters of the screen.

Table 7: Doctor

### 8.4 Sector

In the screen it is possible to manage the sector data.

File Features ?
Laboratory Diseases Doctor Department
Department Name:
Departments List
Urologia
Pediatria
Ostetricia

Picture 8.4 - Sector

Please find below description for each particular field on this screen

Field Name	Description
	"Previous sector detail" and "Following sector detail" keys scroll the sector list.
Ŧ	"Add new sector" key enters a new sector in the list
$\bigcirc$	"Remove sector" key deletes the selected sector from the list.
	"Save sector" key saves the sector selected in the list


"Clean" key resets the parameters of the screen.

Table 8: Sector

# 9 Quality Control



In the following screen it is possible to create a database with information regarding the quality control of the examined samples.

### 9.1 Configuration

In the following screen it is possible to insert, modify and delete references for the quality control of the samples

File Features ?			
Configuration Results	graphics		
Check:	Lot:	expiry:	
17	17	30.09.2009	~
Patient type:	fractions groups:		
Human	<ul> <li>Human serum 5 fraction</li> </ul>	<b>∽</b>	
Fraction	Expected Values Range +/-		
albumin 🗧 1) Albumin	0.00 0.00		
Check	Lot	Expiration Date	_
21	21	12/04/2010	
√ 17	17	30/09/2009	
16	16	21/09/2009	



Please find below description for each particular field on this screen

Field Name	Description
	"Previous control detail" and "Following control detail" keys scroll the sector list.
Ð	"Add new control" key enters a new sector in the list
Θ	"Remove control" key deletes the selected sector from the list.
	"Save control" key saves the sector selected in the list.
	"Clean" key resets the parameters of the screen.



Table 9: Configuration

### 9.2 Results

In the following screen it is possible to: manage quality control reports in tabular format.

File Features	?					
Configural	tion 📄 Result	s 📄 graphics				
Check:			Lot:		expiry:	
17			17		30.09.2009	<u>×</u>
Patient typ	e:		fractions groups	s:		
Human		×	Human serum	5 fraction	×	
	Data Samples	taken from t	o 18/02/2014			
Total	Albumin	Alpha 1	Alpha 2	Beta	Gamma	
0						
Max:	Albumin	Alpha 1	Alpha 2	Beta	Gamma	
Min:						
Media: Var:						
CV%:						

Picture 9.2 - Results

Please find below description for each particular field on this screen.

Field Name	Description
	"Previous control detail" and "Following control detail" keys scroll

	the list of the doctors.
$\mathbf{e}$	The "Print" key displays the print preview with the results of the quality control reports in tabular format.

Table 10: Results

## 9.2.1 Print Preview

In this screen it is possible to print the tabulate with the details of the control and the heading of the laboratory, which displays in the "Personal Data/Laboratory" screen.

🕱 Anteprima di Stampa	
Close Print  << << >> >>  Goto 65%	
Close         Print         I<	
Page 2 of 2	

Picture 9.3 - Print Preview

## 9.3 Graphs

Features ?		
Configuration	<b>jraphics</b>	
ieck:	Lot:	expiry:
7	17	30.09.2009 ~
itient type:	fractions groups:	
uman	Human serum 5 fraction	<u>~</u>
Fraction	Expected Values Range +/-	
1) Albumin	0.00 0.00 Da	ata Samples taken from
Mary		
Rif.	my sand and and and and and and and and and	Sarange and a sarange and a sarange
	ayaanat talaata ayaa ayaa ayaa ayaa ayaa ayaa	

In the following screen it is possible to: manage quality control reports in graphical format.

Picture 9. 4 - Graphs

Please find below description for each particular field on this screen.

Field Name	Description
	"Previous control detail" and "Following control detail" keys scroll the list of the doctors. "Previous control detail" and "Following
	control detail" keys scroll the list of the doctors.
$\Theta$	The "Print" button displays the print preview with the results of the quality control reports in tabular format

Table 11: Graphs

### 9.3.1 Print Preview

In this screen it is possible to print the graph with the details of the control and the heading of the laboratory, which displays in the "Personal Data/Laboratory" screen



Picture 9. 5 - Print Preview Graphs

### **10** Data exchange with third-party application

The import/export operations of the data include information exchange between SAIO applications and third-party. This exchange happen thorough the protocol TCP/IP with the communication channel SOCKET\_IMP\_EXP\_PORT (3300).

### **10.1** Import Patient

The import of patient data from an external database is performed in the feature "Current Data / Serumholder management". The start of this feature requires the following steps:

1) Enter the IP address or the name of the machine which contains the needed patient data.

2) Insert the number of patients which data are required. It is possible to import data for 24/48 patients, because the serum holder provides these number of locations for the samples to be tested.

3) The connection with the machine will start via the communication channel SOCKET\_IMP\_EXP\_PORT (3300).

4) The request is sent through the composition of the following buffer data:

unsigned char cRequest[2];

cRequest[0] = IMPORT\_PATIENTS; //codice di identificazione per l'operazione di import dati

cRequest[1] = NUM\_PAZIENTI; //numero pazienti da importare;

unsigned char cSize[4];

cSize[0..3] = sizeof(cRequest);

The buffer of request that will be sent on the communication channelwill be the result of the connections of the buffers CSize + cRequest. On top of the buffer will be copied the actual number of bytes that identifies the data import request.

If the machine specified in the paragraph 1 receives a buffer data with a length different from sizeof (CSizecRequest +) or if it receives a buffer data with the same length and with identification code different from IMPORT\_PATIENTS, it must discard the communication.

 In the positive case, the machine specified in paragraph 1, will dial a buffer data with the following format/

unsigned char cResponse[sizeof(unsigned char) + (NUM\_PAZIENTI\*SIZE\_DATAPATIENT)];

[0] = NUM\_PAZIENTI; //numero pazienti da inviare;

[1..n] = DATAPATIENT[NUM\_PAZIENTI]; //strutture dati dei pazienti da inviare;

in the 0 position will be insert the number of sent patients and from position1 will be insert the data structure, such as DATAPATIENT in a sequence similar to the number of patients

unsigned char cSize[4];

cSize[0..3] = sizeof(cResponse);

- The buffer response that will be sent to the SAIO communication channel and it will be the result of the connections of the buffers CSize + cResponse. On top of the buffer will be copied the actual number of bytes that identifies the data import response.
- Use the file specified in Appendix "A" to implement the communication buffer between the SAIO and the third parties.

### 10.2 Export Patient

The export of patient data to an external database is performed in the feature "Current data / Change Trace", "Current data / Print", "Search test / Advanced Search" and "Search Test / Print".

The start of this feature requires the following steps:

1. Enter the IP address or the machine name to which send patient data.

2. The connection with the machine will start via the communication channel SOCKET\_IMP\_EXP\_PORT (3300)

#### 3. The request is sent through the composition of the following buffer data:

unsigned char cSend[2 + (NUM\_PAZIENTI \* SIZE\_RESULTPATIENT)];

cSend[0] = EXPORT\_PATIENTS; //codice di identificazione per l'operazione di export dati cSend[1] = NUM\_PAZIENTI; //numero pazienti da esportare; cSend[2..n] = RESULTPATIENT[NUM\_PAZIENTI]; //strutture dati dei pazienti da esportare; unsigned char cSize[4]; cSize[0..3] = sizeof(cSend);

The export data buffer that will be sent on the communication channel to the third-party applications will be the result of the connections of the buffers CSize + cSend.

On top of the buffer will be copied the actual number of bytes that identifies the data import request.

In the case in which the machine specified in point 1 receives a data buffer with length different from sizeof (CSizecSend +) or on the same length receives an identification code different from EXPORT\_PATIENTS must discard the communication.

If the machine specified in the paragraph 1 receives a buffer data with a length different from sizeof (CSizecRequest +) or if it receives a buffer data with the same length and with identification code different from IMPORT\_PATIENTS, it must discard the communication.

In the case in which the machine specified in point 1 receives correctly the results from SAIO, then it must respond to the sender with a buffer that indicates the result of the export operation. It will respond with the following buffer:

unsigned char cResult[sizeof(unsigned long)];

cResult[0..3] = 1; //1 in caso di successo; 0 in caso di errore

unsigned char cSize[4];

cSize[0..3] = sizeof(cResult);

- The data response buffer that will be sent to the SAIO communication channel and it will be the result of the connections of the buffers cSize + cResult.
- Use the file specified in Appendix "A" to implement the communication buffer between the SAIO and the third parties.

# "A" Appendix

### Include per import/export data

#if !defined(DScanSocketData\_h)

#define DScanSocketData\_h

//numero porta del socket per la comunicazione con enti esterni

#define SOCKET\_IMP\_EXP\_PORT 3300

//codice per identificare un'operazione di import dei dati

#define IMPORT\_PATIENTS 0xBE

//codice per identificare un'operazione di export dei dati

#define EXPORT\_PATIENTS 0xCE

//struttura dati del paziente

typedef struct

{

unsigned long	m_ImportID;	//ID external - assigned to the patient
long	m_PatientType_Id;	/Tipo paziente: 0=indefinito, 1=umano, 2=cane, 3= gatto, //4=cavallo, 5=mucca.
char	m_LastName[64];	/Cognome
char	m_Name[64];	//Nome
char	m_AcceptanceCode[64];	//Codice Accettazione
double	m_TotProtein;	//ProteineTotali
byte	m_Age;	//Eta
byte	m_Sex;	//Sesso
unsigned long	m_AcceptanceDate; //Data Acc	cettazione
unsigned long	m_BirthDate;	//Data Nascita

#### } DATAPATIENT, \*LPDATAPATIENT;

#### #define SIZE\_DATAPATIENT sizeof(DATAPATIENT)

//single fraction structure

typedefstruct

{

	byte	m_FractionPos;	//Fraction numeration (1, 2,, n)
	char	m_FractionName[32];	//Name
	unsigned long	m_FractionStart;	/start position on the graph
	unsigned long	m_FractionEnd;	//End position on the graph
	double	m_FractionRatio;	/ Relation value A/G
	double	m_RatioNormMin;	/Relation Value A/G min
	double	m_RatioNormMax;	/ Relation Value A/G max
	double	m_FractionVal;	// Relation Value
	double	m_ValueNormMin;	/absolute value Min
	double	m_ValueNormMax;	/absolute value Max }
DATAF	RACTION, *LPDATAFRACTIO	ON;	
#define S	SIZE_DATAFRACTION	sizeof(DATAFRACTION)	
//struttur	a dati bitmap grafico		
typedef s	struct		
{			
	Int	m_Size;	//size file bitmap of the graph
	char	m_FileData[12288];	//data file (max 12 KB)
} DATA	GRAPH, *LPDATAGRAPH;		
#define S	SIZE_DATAGRAPH	sizeof(DATAGRAPH)	
//struttur	a dati bitmap strip		
typedef s	struct		
{			
	int	m_Size;	//size file bitmap of the graph
	char	m_FileData[3072];	//data file (max 3 KB)}

DATASTRIP, \*LPDATASTRIP;

#define SIZE\_DATASTRIP

sizeof(DATASTRIP)

#### //struttura dati del risultato del paziente

#### typedef struct

<sup>{</sup> 

DATAPATIENT	m_Patient; //dati paz	iente
unsigned long	m_AcquisitionID;	/Id lettura
unsigned long	m_StripPos;	//position number of slot in the serum holder
char	m_Comment[256];	/comment on the paragraph
double	m_NormValRifAG;	/Relation Value A/G
double	m_NormValRifAG_Min;	//Relation Value A/G Min
double	m_NormValRifAG_Max;	//Relation Value A/G Max
char	m_UnitOfMeasureAG[32];	//unit of measure of the relation A/G
double	m_NormValProtTot_Min;	//Proteins value total min
double	m_NormValProtTot_Max;	//Proteins value total max
unsigned long	m_StripLen;	//length of the graph
unsigned char	m_MinTags;	//minimum on the graph
DATAFRACTION	m_Fractions[12];	/fraction of the graph
DATAGRAPH	m_GraphBitmap;	//bitmap of the graph
DATASTRIP	m_StripBitmap;	//bitmap of the strip

} RESULTPATIENT, \*LPRESULTPATIENT;

#define SIZE\_RESULTPATIENT

sizeof(RESULTPATIENT)

#endif //DScanSocketData\_h

## Notes

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