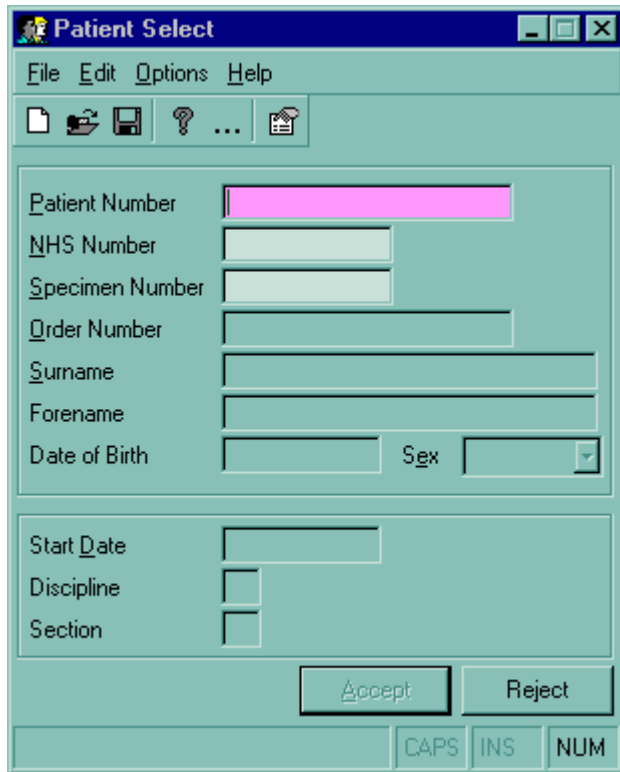


iLab GUI – guide to WRNQ

Remember :- PRESS TAB BETWEEN FIELDS.

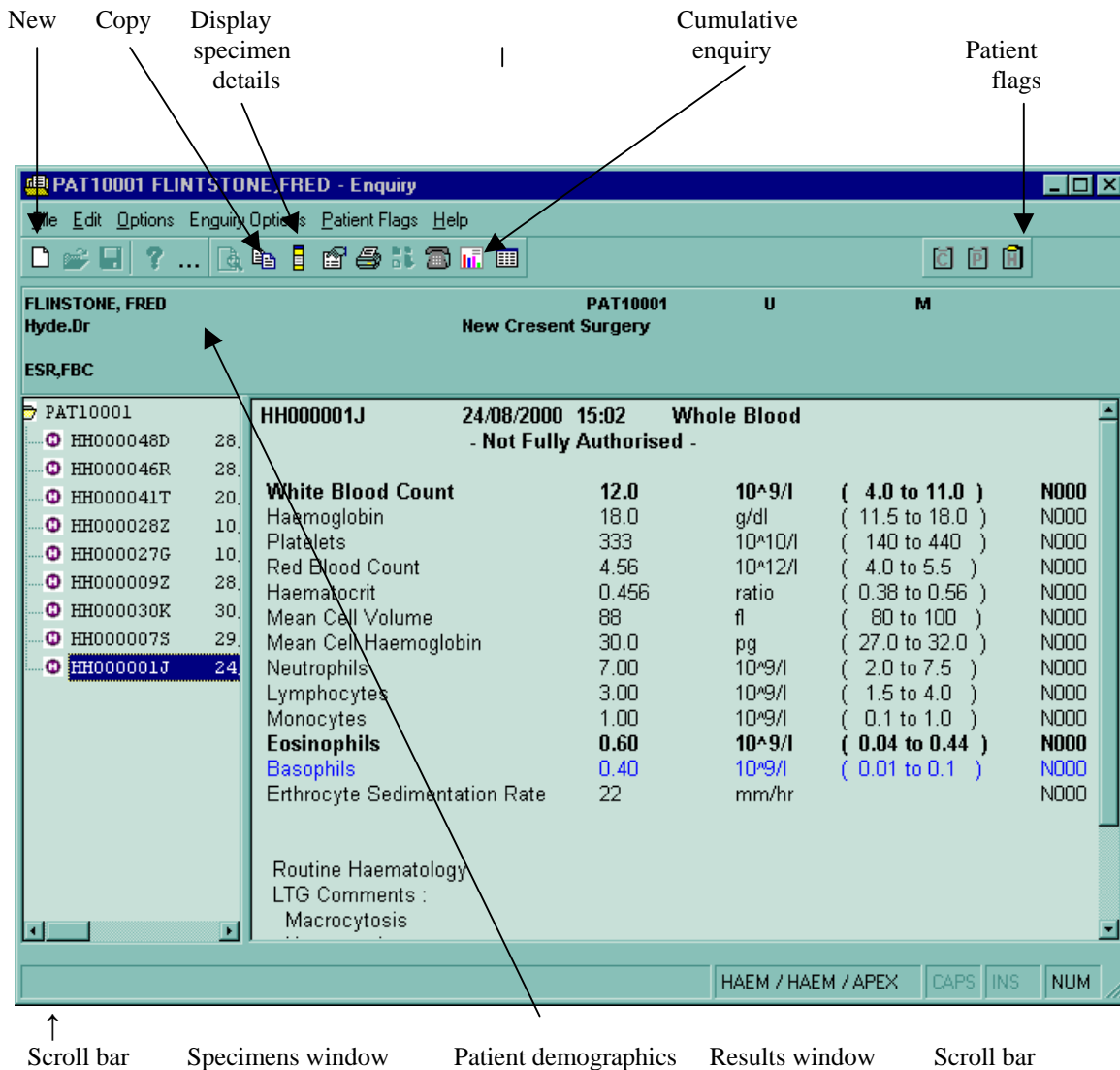
1. **To log on** - double click on the iLaboratory logon icon, type in your user name, press Tab, type in your password then press Enter.
2. **To run WRNQ from the menu** - double click on the WRNQ icon on the GUI menu or type WRNQ into the Option field and press Enter.
3. **Patient number** – type the Patient Number (Hospital number including 0s ie 0172679 or 0045781 you must have 7 numbers) into the appropriate field and **press tab**. Enter the 1st two characters of the patient's surname and **press tab**. Click on Accept or press Enter.



The screenshot shows a 'Patient Select' dialog box with the following fields and controls:

- File Edit Options Help (Menu)
- File, Edit, Options, Help (Toolbar icons)
- Patient Number (Text field, highlighted in pink)
- NHS Number (Text field)
- Specimen Number (Text field)
- Order Number (Text field)
- Surname (Text field)
- Forename (Text field)
- Date of Birth (Text field)
- Sex (Dropdown menu)
- Start Date (Text field)
- Discipline (Text field)
- Section (Text field)
- Accept (Button)
- Reject (Button)
- CAPS, INS, NUM (Buttons)

4. **Patient Details screen** – The first screen loaded displays the patient details as the main window on the right, with the specimen numbers for that patient on the left. Moving the cursor over a specimen number displays the specimen date & time, tissue type and result availability as tooltips. This information can be obtained for all the specimens by dragging the separator bar over to the right.
5. **Results screen** - Select the specimen required by clicking on the specimen number. You can move up and down through the available specimens using Cursor Up/Down keys, or by clicking on the specimen numbers.
If the report text is too long to be seen in entirety in the window press tab and use Page Up/Down (previous/next) keys to view the report (or use the scroll bar to move up and down). Pressing tab again moves the cursor back to the specimen number on the left side of the screen.



Results Enquiry window

- BB = Biochemistry
- HH = Haematology
- II = Immunology and worked referred to other Laboratories
- MM = Microbiology
- PP = Histopathology
- PC = Non Gynae Cytology
- PG = Gynae Cytology
- TT = Blood Transfusion

6. **Selecting a new patient** – Return to the Patient select screen by clicking on the Open icon (or Ctrl-O).
7. **Selection by specimen number** – If the specimen number for the enquiry is known, on the Patient Select screen press tab to move to the Specimen number field and enter the specimen number. Press tab then Enter.
8. **Tool Bar Icons**

New: - Displays the Patient Select window so a new *enquiry* can be performed.

Display Specimen Details: - The current *specimen* details are displayed in a format as in *Order Entry*.

Cumulative enquiry: - displays the *tests* of the current *specimen* in a cumulative format. Historic *specimens* are also displayed. *Specimen* data is presented by the most recent on the left to the oldest on the right.

If there are more *tests* or *specimens* than can be displayed in the window, then scroll bars will enable the *user* to access the remaining *tests* or *specimens*.

Note tests from previous *specimens* are only displayed if they match the *tests* displayed on the current *specimen*. For example if you are looking at an FBC result ONLY previous FBCs will be displayed.

9. **To exit Enquiries** – click on the **x** at the top right of the window, or press Alt-F4 or select exit from drop down File menu.

Return to Patient Select screen using the New icon.

The screenshot shows a software window titled "APX362115 ADAM.PETER - Enquiry". The window has a menu bar with "File", "Edit", "Options", "Patient Flags", and "Help". Below the menu bar is a toolbar with various icons. The main area is divided into several sections:

- Patient Demographics:** Case No.: APX362115, Pat Name: ADAM, PETER, Date of Birth: 01/01/1953, U, <PREVIOUS>. Ward/Dr.: Pathology Location / Dr B A Green. Col Date: 30/10/1998, Cut-up Path: HARROP.
- Specimen List (Tree View):** A tree view on the left shows a folder "APX362115" containing several specimen numbers (e.g., PP123123/98, PP001025/98) and their sequences (Seq 1, Seq 2).
- Report Content:** The main area displays details for specimen PP001025/98, dated 30/10/1998 at 12:51 PM. It includes "Report sequence 2", "Technically Validated" by Dave Campbell, and a "Z Logical Name" of 10/11/1998 10:31. The report text includes:
 - Gross Description:** Site of tumour ascending colon, Maximum tumour diameter 1.35 cm, Presence of tumour perforation Yes.
 - For Rectal Tumours:** Tumour is above the peritoneal reflection, Distance from pectinate line 1.05 cm.
 - Histology Type:** Adenocarcinoma Yes (to include mucinous and signet ring adenocarcinomas), If No, state other, Differentiation by predominating area poor.
 - Local Invasion:** Submucosa (pT1) N, Muscularis propria (pT2) N, Beyond Muscularis propria (pT3) Y.

Annotations with arrows point to various features: "Patient Flags" points to the menu bar; "4 lines of patient demographic details as defined in MDD" points to the patient information section; "Scroll bar to move up & down the report text." points to the vertical scroll bar on the right; "Tree view" showing multiple report sequence for one specimen number. points to the specimen list; and "Separator bar may be dragged to right & left." points to the vertical line between the specimen list and the report content.

"Tree view" showing multiple report sequence for one specimen number.

Separator bar may be dragged to right & left.