



RNA Extraction from Breast Cancer Xenografts and Lymph Node Metastases at The Institute of Cancer Research

CONTEXT

Within the context of The Institute of Cancer Research, *ex vivo* tumor tissues are being analyzed to study the gene expression of breast tumors and their metastases.

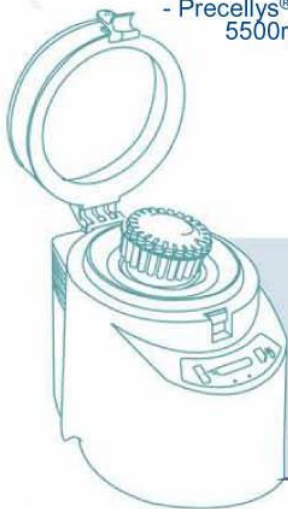
Breast cancer xenografts from different cell lines (MDA-MB-435 and GI-101), as well as their lymph node metastases, were frozen in liquid nitrogen after collection.

MATERIAL

- Precellys®24
- Precellys®24 kit CK14 (small ceramic beads)
- Sample : tumors from breast cancer xenografts and lymph node metastases (frozen)
- Buffer : 600µl of lysis buffer (RTL and β-mercaptoethanol)

PROTOCOL

- Precellys®24 parameters:
5500rpm, 1x20 sec.



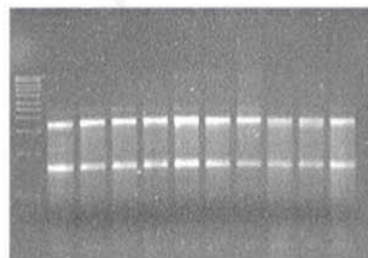
RESULTS

In collaboration with The Institute of Cancer Research, McElain Laboratories, Sutton UK.

RNA extraction are performed following 10 different protocols :

Number	Sample	Time	Number	Sample	Time
1	Primary Tumors	10 Sec.	6	Primary Tumors	20 Sec.
2		3 x 10 Sec.	7		20 Sec.
3		20 Sec.	8	Lymph node metastases	20 Sec.
4		20 Sec.	9		20 Sec.
5		20 Sec.	10		20 Sec.

RNA gel procedure :
0.5 µg of total RNA diluted in 15µl of denaturing loading buffer (containing urea) denatured and run on a 1% agarose gel in 1x TBE at 140v for 1hour.



1 2 3 4 5 6 7 8 9 10
Gel electrophoresis analysis after the 10 protocols. Efficiency is validated on the primary tumors and lymph node metastases.



CONCLUSION

The Precellys® kits allow a quick and effective homogenization of the xenograft tissues, and the total RNA extracted with an appropriate kit following tissue lysis with the Precellys® kits is of good quality.

For more details, please contact precellys@bertin.fr or visit our website

