

IDENTIFICATION OF TRANSLOCATION t(11;18) API2/MLT

AMPLI-SET/API2-MLT

Cat. n.1.400.1

The translocation t(11;18)(q21;q21) represents the most common abnormality in the marginal zone lymphoma extranodal that rise from the lymphoid tissue mucosa- associated (MALT). This translocation results in the expression of a chimeric transcript obtained from the fusion of AP12 gene (Apoptosis inhibitor gene 2) on chromosome 11 to the gene MLT (MALT lymphoma translocation gene) on chromosome 18q21.

Il kit Ampli-SET-API2/MLT permette di identificare, mediante l'uso della Reverse Transcription-Polymerase Chain Reaction (RT-PCR), la traslocazione t(11;18). The analysis of the API2/MLT fusion transcript is based on primers design on opposite sites of fusion regions so that the PCR product will include the specific fusion sequence.

Principle of assay:

- extraction of RNA
- retro-transcription
- amplification
- detection on agarose gel

Applicability: extracted and purified RNA

Number of tests: 45.

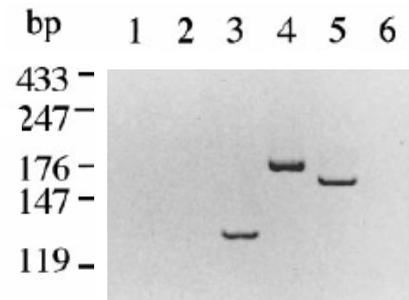
REAGENTS AND STORAGE

RETROTRANSCRIPTION	
Mix RT	-20°C
Reverse Transcriptase (40U/μl)	-20°C
Rnase inhibitor (40U/μl)	-20°C
Random primers	-20°C
RNsa/DNase -free water	-20°C
AMPLIFICATION	
Mix PCR API2/MLT	-20°C
Taq Polymerase (5U/μl)	-20°C
RNsa/DNase -free water	-20°C
Positive control	-20°C

Stability: over 18 months if correctly stored.

ANALYSIS OF RESULTS

The positive samples for the AP12/MLT rearrangement will produce discrete bands.



References:

- Akagi T et al. Oncogene 18:5785-5794, 1999
Auer IA et al. Ann Oncol 8:979, 1997
Dierlamm J et al. Blood 93:3601-3609, 1999
Izumiyama K et al. Oncogene 22:8085-8092, 2003
Lucas PC et al. J Biol Chem;276:19012-19019, 2001
Ott G. et al. Cancer Res 57:3944, 1997
Sanchez-Izquierdo D et al. Blood 101:4539-4546, 2003