

1. Record Nr.	TD17006628
Autore	Ferrari, Anna
Titolo	Depicting the role of CDKN2A/ARF alterations in adult BCR-ABL1-positive acute lymphoblastic leukemia patients: from genomic deletions to prognostic impact [Tesi di dottorato]
Lingua di pubblicazione	Non definito
Formato	Tesi di dottorato
Livello bibliografico	Monografia
Note	In relazione con <a href="http://amsdottorato.unibo.it/3617/">http://amsdottorato.unibo.it/3617/</a>
Sommario	<p>This 9p21 locus, encode for important proteins involved in cell cycle regulation and apoptosis containing the p16/CDKN2A (cyclin-dependent kinase inhibitor 2a) tumor suppressor gene and two other related genes, p14/ARF and p15/CDKN2B. This locus, is a major target of inactivation in the pathogenesis of a number of human tumors, both solid and haematologic, and is a frequent site of loss or deletion also in acute lymphoblastic leukemia (ALL) ranging from 18% to 45% 1. In order to explore, at high resolution, the frequency and size of alterations affecting this locus in adult BCR-ABL1-positive ALL and to investigate their prognostic value, 112 patients (101 de novo and 11 relapse cases) were analyzed by genome-wide single nucleotide polymorphisms arrays and gene candidate deep exon sequencing. Paired diagnosis-relapse samples were further available and analyzed for 19 (19%) cases. CDKN2A/ARF and CDKN2B genomic alterations were identified in 29% and 25% of newly diagnosed patients, respectively. Deletions were monoallelic in 72% of cases and in 43% the minimal overlapping region of the lost area spanned only the CDKN2A/2B gene locus. The analysis at the time of relapse showed an almost significant increase in the detection rate of CDKN2A/ARF loss (47%) compared to diagnosis (<math>p = 0.06</math>). Point mutations within the 9p21 locus were found at very low level with only a non-synonymous substitution in the exon 2 of CDKN2A. Finally,</p>

correlation with clinical outcome showed that deletions of CDKN2A/B are significantly associated with poor outcome in terms of overall survival ( $p = 0.0206$ ), disease free-survival ( $p = 0.0010$ ) and cumulative incidence of relapse ( $p = 0.0014$ ). The inactivation of 9p21 locus by genomic deletions is a frequent event in BCR-ABL1-positive ALL. Deletions are frequently acquired at the leukemia progression and work as a poor prognostic marker.

---

Localizzazioni e accesso

[http://memoria.depositolegale.it/\\*/http://amsdottorato.unibo.it/3617/1/Ferrari\\_Anna\\_Tesi.pdf](http://memoria.depositolegale.it/*/http://amsdottorato.unibo.it/3617/1/Ferrari_Anna_Tesi.pdf)

---