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Long non coding RNAs era in liver cancer

Francesca Guerrieri

4 Abstract

Hepatocellular carcinoma (HCC) is one of the most common malignancies leading to high mortality rates in the general population and the sixth most common cancer worldwide. HCC is characterized by deregulation of multiple genes and signaling pathways. These genetic effects can involve both protein coding genes as well as non coding RNA genes. Long non coding RNAs (lncRNAs) are transcripts longer than 200 nt, constituting a subpopulation of ncRNAs. Their biological effects are not well understood compared to small non coding RNA (microRNAs), but they have been recently recognized to exert a

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