

# INPLASY PROTOCOL

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**Review Stage at time of this submission:** Piloting of the study selection process.

**Conflicts of interest:**  
None declared.

## INTRODUCTION

**Review question / Objective:** Despite improvement in expounding the molecular mechanism of long noncoding RNAs (lncRNAs) in malignant tumors, efforts to find clinically relevant prognosis-associated lncRNAs are necessary.

However, the clinical prognostic significance of LINC01133 in tumors is not fully understood.

**Condition being studied:** Inclusion criteria were as follows: (1) the study included a rigorous pathological diagnosis of the associated tumor; (2) Prognostic

## The clinical prognostic value of lncRNA LINC01133 in Chinese patients with cancer: a meta-analysis

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**Condition being studied:** Inclusion criteria were as follows: (1) the study included a rigorous pathological diagnosis of the associated tumor; (2) Prognostic characteristics and survival curve analysis of different expression levels of LINC01133 in patients with malignant tumor; (3) Complete information is provided concerning the association between different expression levels of LINC01133 and clinicopathological data of malignant tumors. (3) Studies that can visualize hazard risk (HR) and 95% confidence intervals (CI) or be able to evaluate relevant data through computation.

**INPLASY registration number:** This protocol was registered with the International Platform of Registered Systematic Review and Meta-Analysis Protocols (INPLASY) on 23 June 2021 and was last updated on 23 June 2021 (registration number INPLASY202160087).

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## METHODS

**Search strategy:** The search strategies are as follows, including: (((clinicalpathological feature) OR ((prognosis) OR (overall survival)) OR (clinicalpathological characteristics))) OR (( "Neoplasms/mortality"[Mesh] OR "Neoplasms/pathogenicity"[Mesh] OR "Neoplasms/pathology"[Mesh] OR "Neoplasms/statistics and numerical data"[Mesh] OR "Neoplasms/surgery"[Mesh] OR "Neoplasms/therapy"[Mesh] ))) AND ((LINC01133) OR (((("long non-coding RNA LINC01133, human" [Supplementary Concept]))) AND "RNA, Long Noncoding"[Mesh])).

**Participant or population:** The study included a rigorous pathological diagnosis of the associated tumor (1229).

**Intervention:** Inapplicability.

**Comparator:** Inapplicability.

**Study designs to be included:** RCT.

**Eligibility criteria:** (1) the study included a rigorous pathological diagnosis of the associated tumor; (2) Prognostic characteristics and survival curve analysis of different expression levels of LINC01133 in patients with malignant tumor; (3) Complete information is provided concerning the association between different expression levels of LINC01133 and clinicopathological data of malignant tumors. (3) Studies that can visualize hazard risk (HR) and 95% confidence intervals (CI) or be able to evaluate relevant data through computation.

**Information sources:** Electronic databases.

**Main outcome(s):** The merged results of this meta-analysis suggest that LINC01133 overexpression may have a predictive effect on prognosis of digestive cancers, especially in colorectal cancer, indicating that LINC01133 is expected to be a target for digestive tumor prognostic monitoring and provide a new therapeutic strategy for biologically targeted therapy.

**Quality assessment / Risk of bias analysis:** The publication bias of the meta-analysis was evaluated using Begg's test. The results demonstrated that this meta-analysis did not exist significant publication bias for OS ( $P=0.858$ ). In addition, sensitivity analysis was applied to test the stability of the pooled results.

**Strategy of data synthesis:** Hazard Ratio or odds ratio was combined using Review Manager (RevMan) 5.3 software. Publication bias and sensitivity analysis was estimated using Stata 14 software. Q test and I<sup>2</sup> were performed to estimate the heterogeneity of results. When I<sup>2</sup>50%), the pooled results selected the random-effects model for data analysis. LINC01133 was significantly correlated with poor prognosis when the results revealing  $HR>1$ . Conversely, high LINC01133 expression indicated a good prognosis when  $HR<1$ .

**Subgroup analysis:** We conducted a subgroup-analysis to analyze the merged effect on digestive tumors. The results indicated that LINC01133 have a certain effect of predicting good prognosis for digestive tumors ( $HR=0.5$ , 95%CI: 0.28–0.92,  $P=0.03$ ), especially in colorectal cancer ( $HR=0.53$ , 95%CI: 0.35–0.80,  $P=0.003$ ). In addition, elevated LINC01133 expression was obviously and negatively associated with TNM stage of tumors (III/IV vs. I/II, Odd Ratio (OR)=0.44, 95%CI=0.23–0.84,  $P=0.01$ ), lymphatic metastasis (Positive vs. Negative, OR=0.30, 95%CI=0.19–0.48,  $P<0.001$ ) and smoking status (Ever vs. Never, OR=0.44, 95%CI=0.25–0.79,  $P=0.006$ ) in Chinese cancer patients.

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**Sensitivity analysis:** The sensitivity analysis illustrated that eliminating any single study did not significantly change the combined HR, suggesting the results were stable and reliable.

**Country(ies) involved:** China.

**Keywords:** Overall survival; Hazard Ratio; Prognosis; LINC01133; Cancer.

**Contributions of each author:**

Author 1 - Fei Xie.

Author 2 - Jie Wang.

Author 3 - Hao Hua.

Author 4 - Qin Yang.