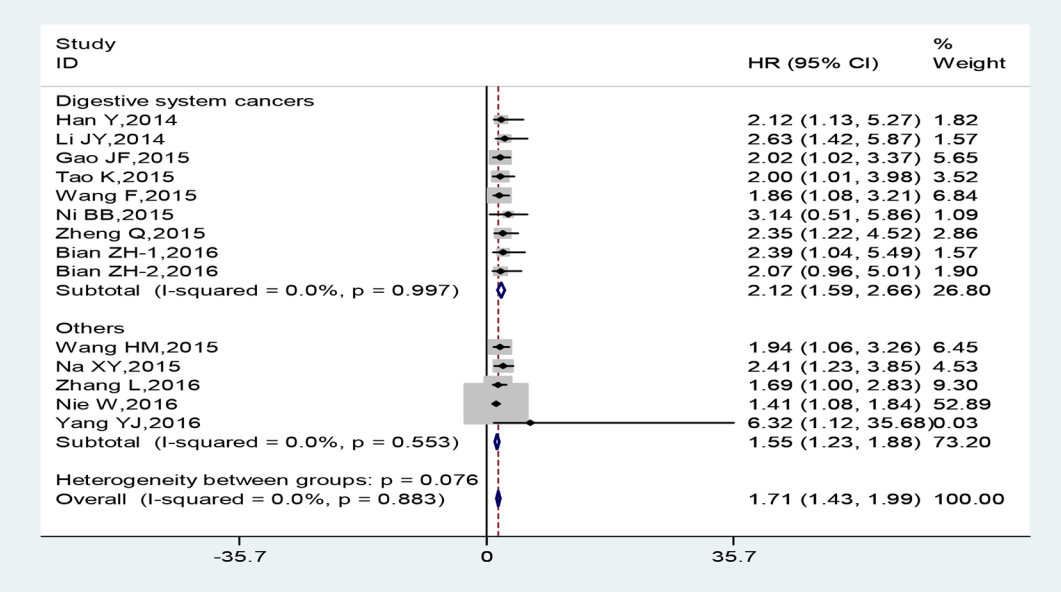
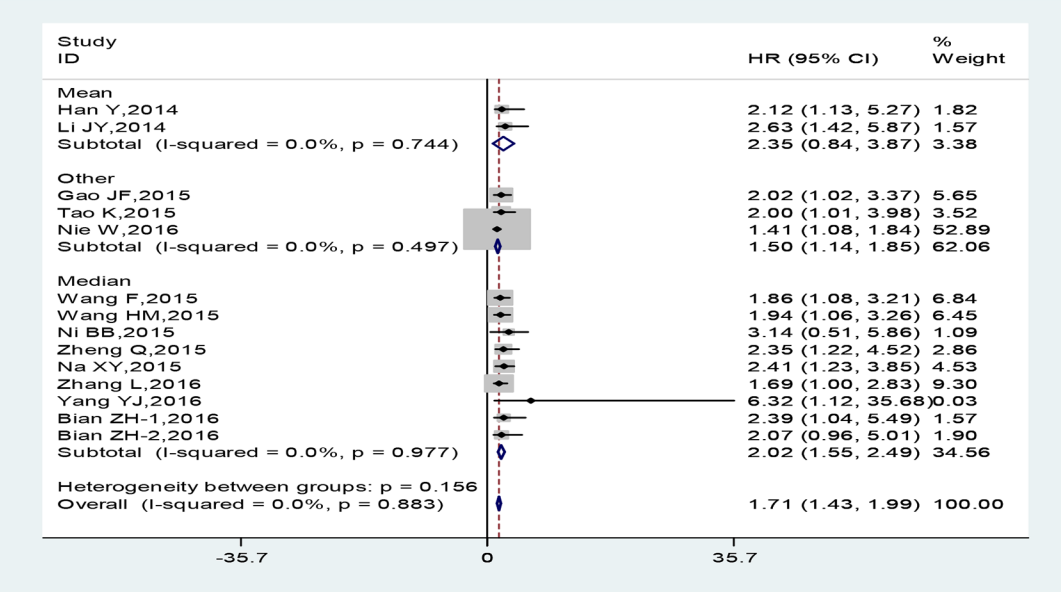
Prognostic value of long non-coding RNA UCA1 in human solid tumors

Supplementary Materials

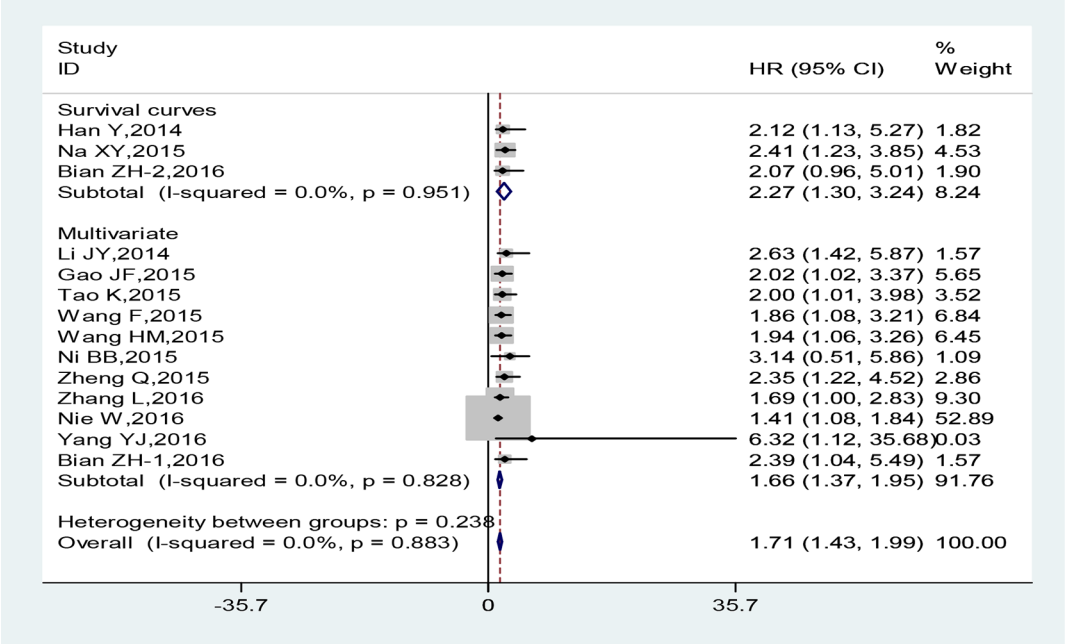
(1) Association between lncRNA-UCA1 expression and OS:



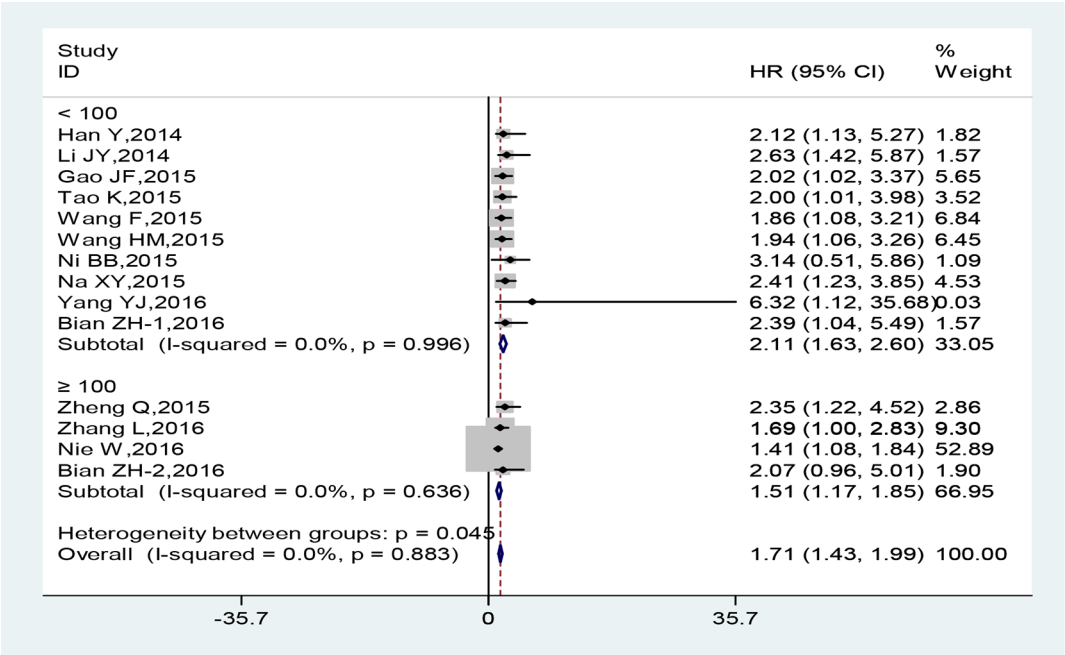
Supplementary Figure S1: Meta-analysis for the pooled HRs of OS stratified by the cancer type.



Supplementary Figure S2: Meta-analysis for the pooled HRs of OS stratified by the cutoff value.

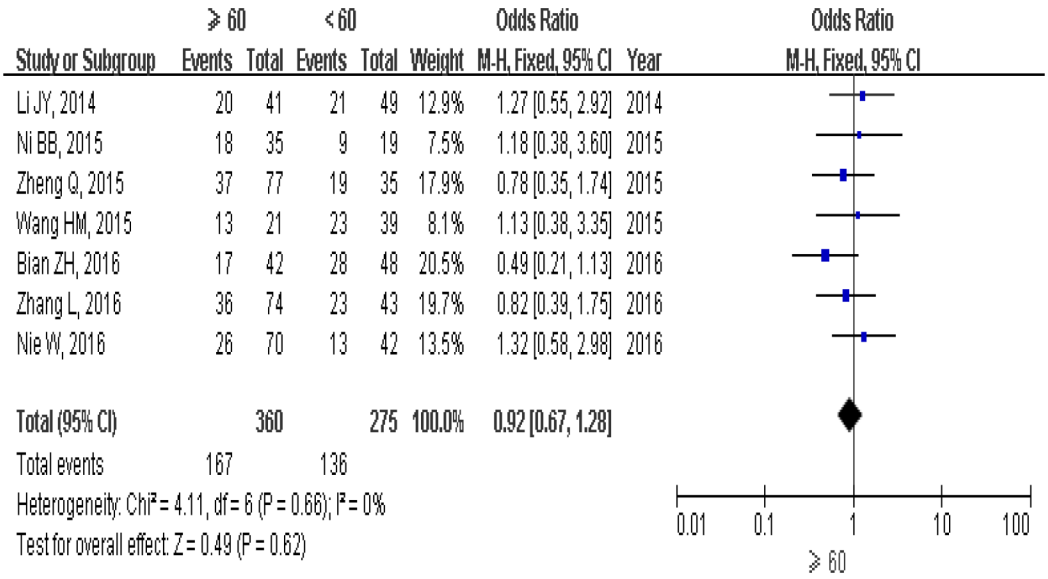


Supplementary Figure S3: Meta-analysis for the pooled HRs of OS stratified by the analysis type.

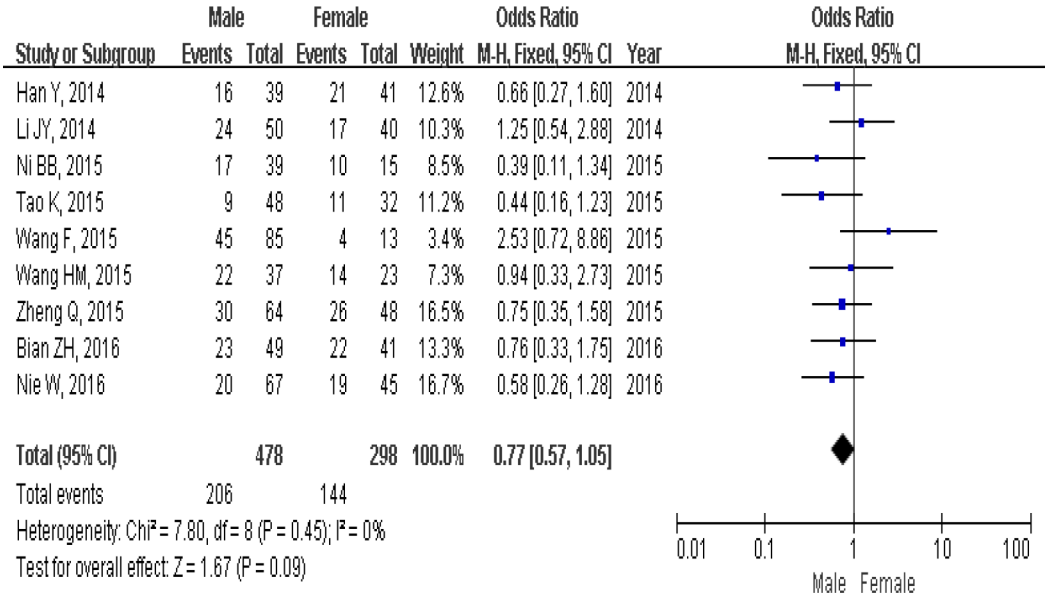


Supplementary Figure S4: Meta-analysis for the pooled HRs of OS stratified by the sample size.

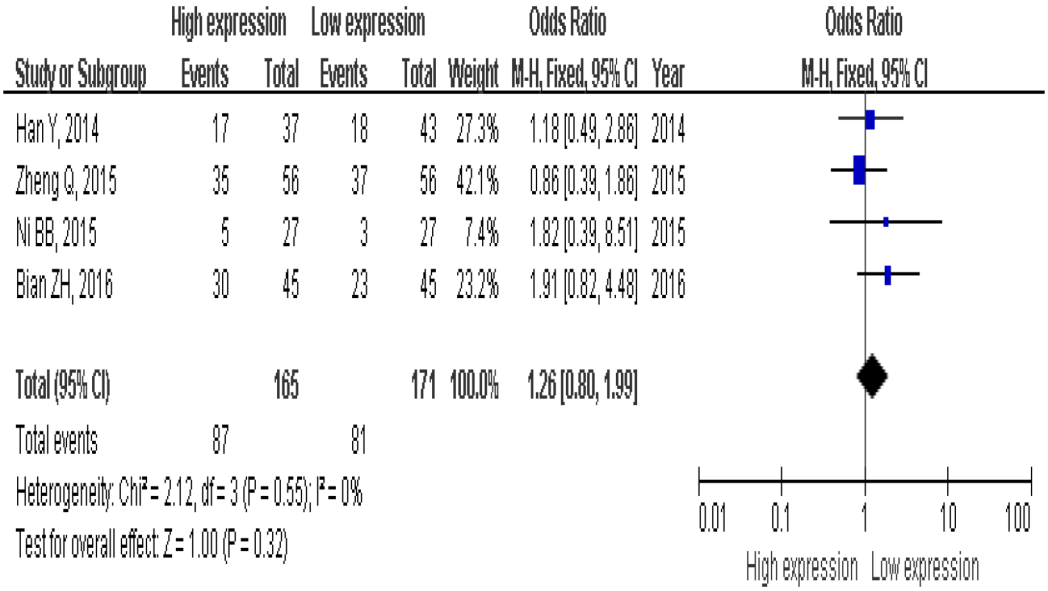
(2) Association between lncRNA-UCA1 expression and clinicopathological parameters:



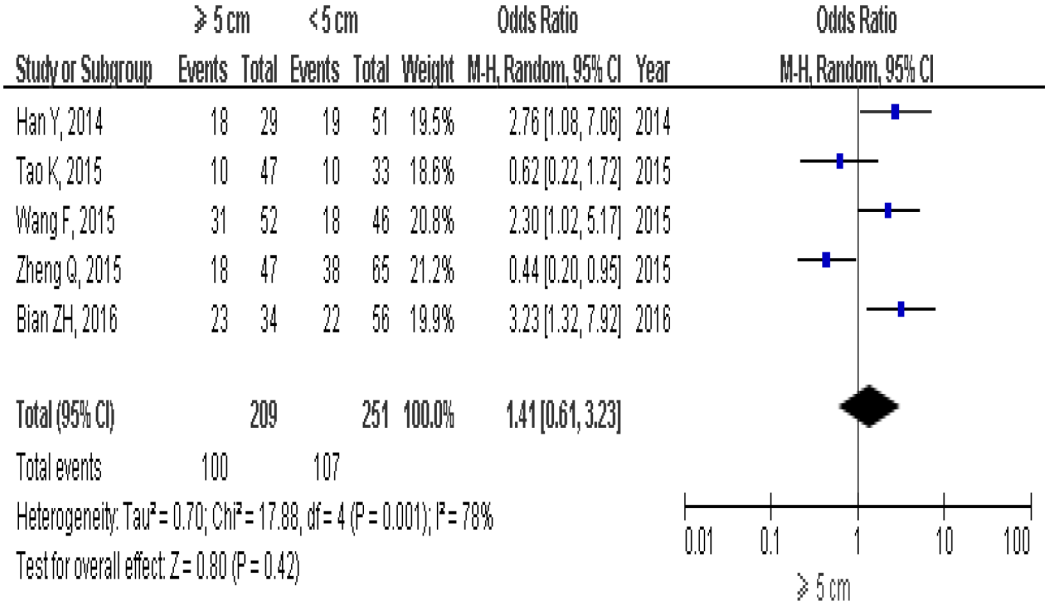
Supplementary Figure S5: The correlation between lncRNA-UCA1 expression and age.



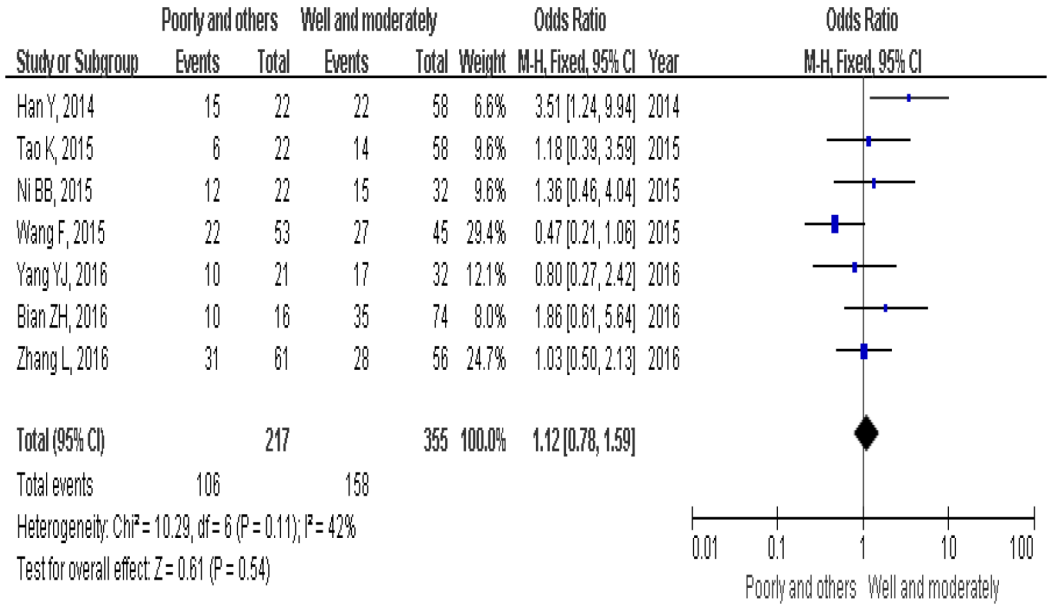
Supplementary Figure S6: The correlation between lncRNA-UCA1 expression and gender (male vs. female).



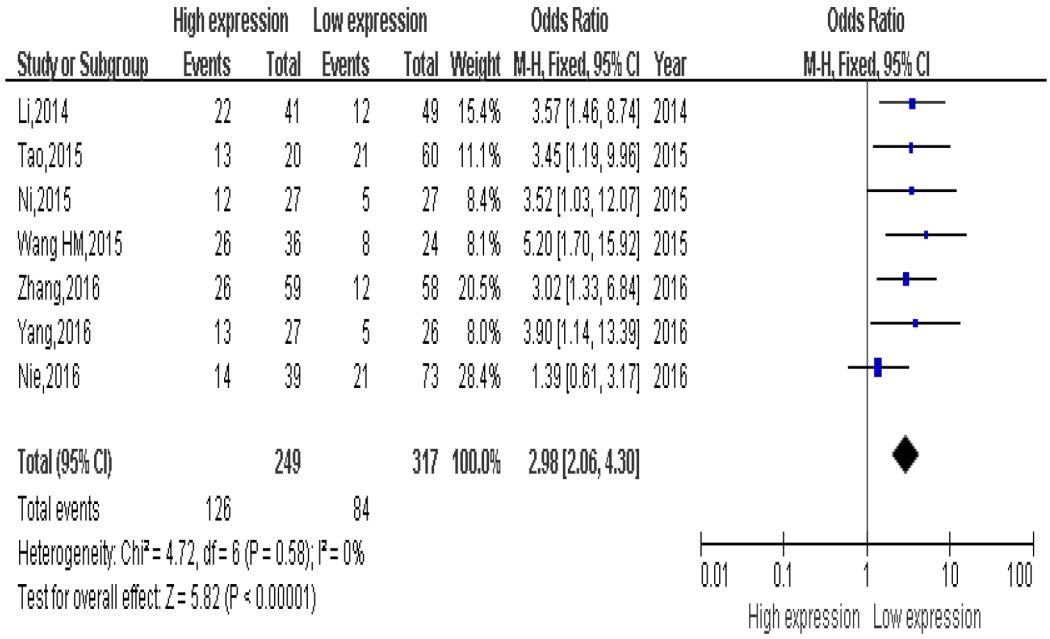
Supplementary Figure S7: The correlation between lncRNA-UCA1 expression and lymphatic invasion (present vs. absent).



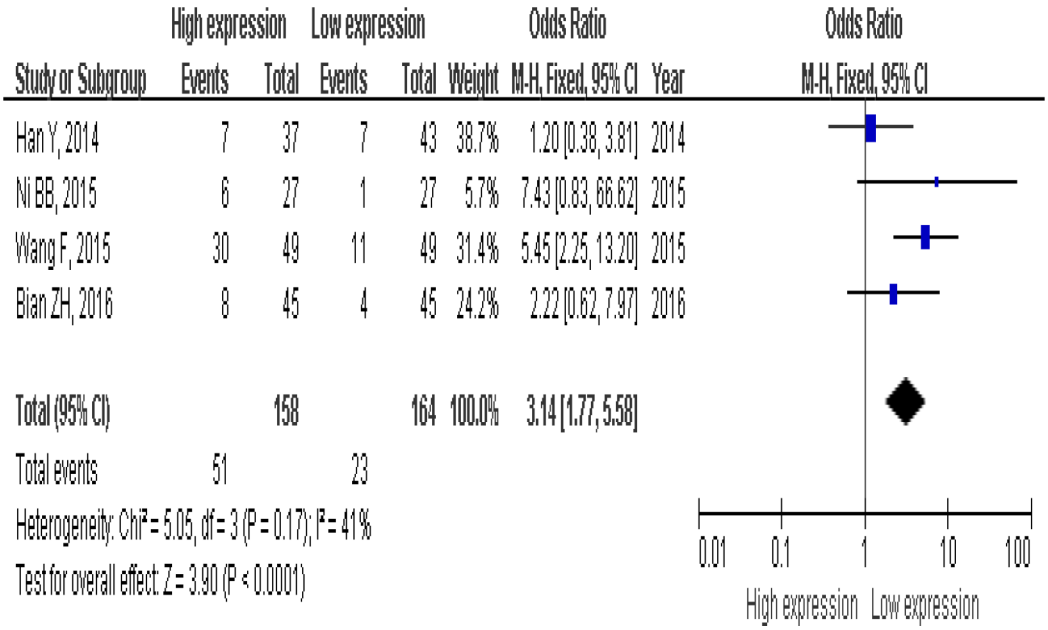
Supplementary Figure S8: The correlation between lncRNA-UCA1 expression and tumour size.



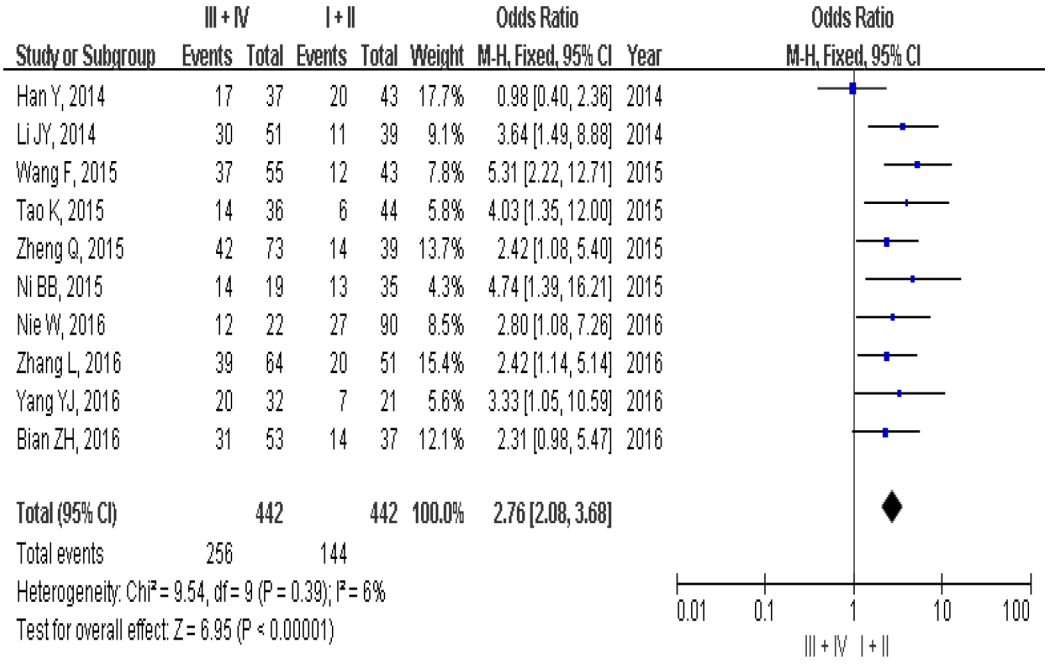
Supplementary Figure S9: The correlation between lncRNA-UCA1 expression and tumor differentiation (poorly/ others vs. well/moderately).



Supplementary Figure S10: The correlation between lncRNA-UCA1 expression and lymph node metastasis (yes vs. no).



Supplementary Figure S11: The correlation between lncRNA-UCA1 expression and distant metastasis (yes vs. no).



Supplementary Figure S12: The correlation between lncRNA-UCA1 expression and TNM stage (III–IV vs. I–II).