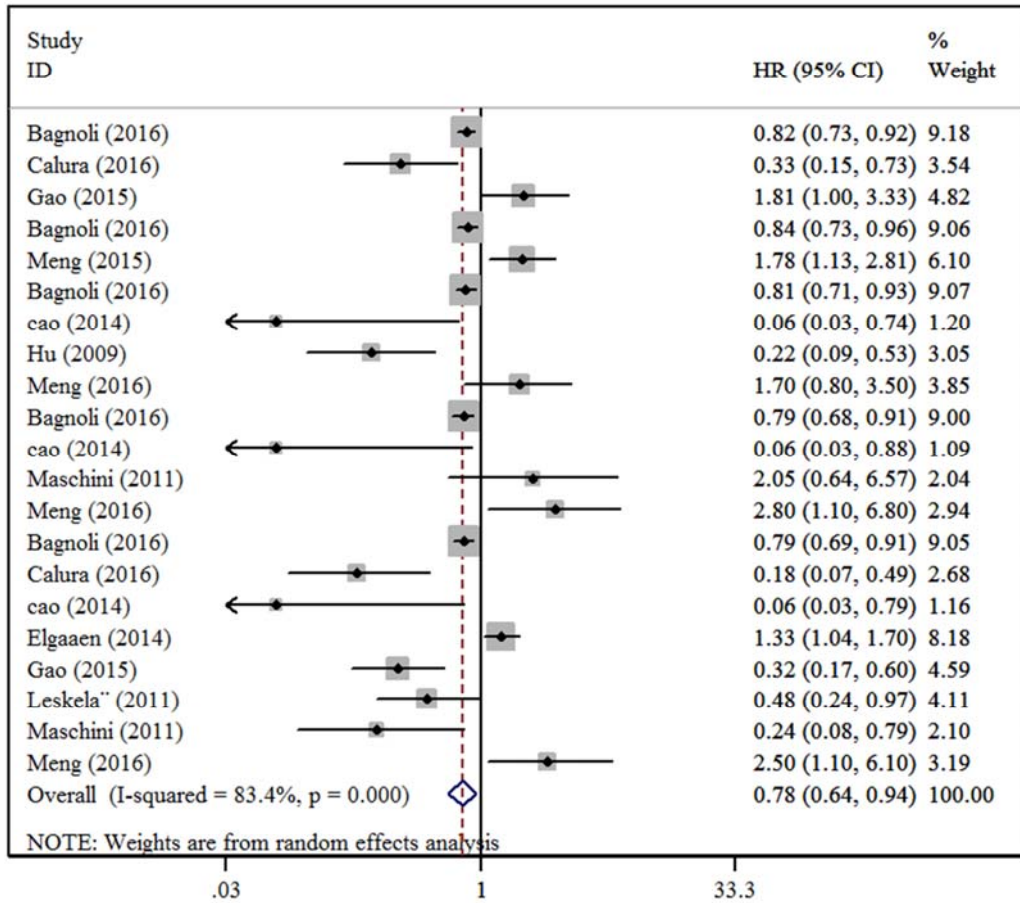


Supplementary Table 1. Reporting recommendations for tumour marker prognostic studies (REMARK)

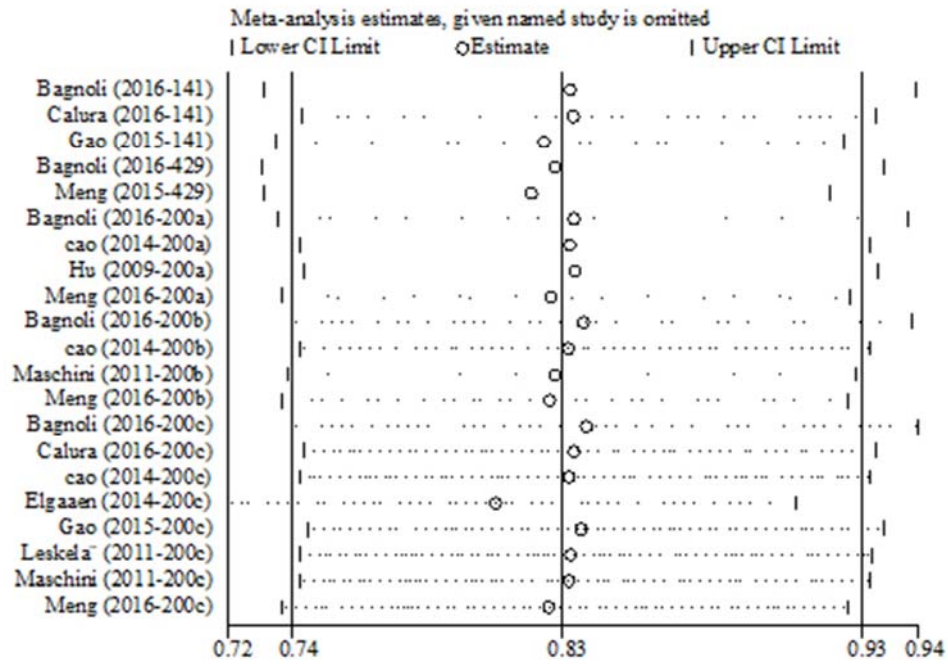
Items	Authors										
	Bagno li	Calur a	Ca o	Elgae n	Ga o	Hu	Kapetan ak is	Le e	Leskelä	Marchi ni	Me
Introduction											
1.objectives and hypotheses	√		√	√	√	√		√	√	√	√
Materials and Methods											
Patients											
2.characteristics	√	√	√	√	√	√	√	√	√	√	√
3.treatment methods	√										
Specimen characteristics											
4.biological material	√	√	√	√	√	√	√	√	√	√	√
Assay methods											
5. method protocol	√		√	√	√		√	√	√	√	√
Study design											
6.method of case selection	√	√	√	√	√	√	√	√	√	√	√
7.define clinical and points examined	√	√	√	√	√	√	√	√	√	√	√
8.candidate variable	√										
9.sample size	√	√	√	√	√	√	√	√	√	√	√
Statistical analysis methods											
10.statistical methods	√	√	√	√	√	√	√	√	√	√	√
11.cutpoint	√	√	√	√	√	√	√	√	√	√	√
Results											
Data											
12.number of patients	√	√	√	√	√	√	√	√	√	√	√
13.demographic characteristics	√	√	√	√	√	√	√	√	√	√	√
Analysis and presentation											
14. the relation of the marker to standard prognostic variables	√	√	√	√	√	√	√	√	√	√	√
15.univariate analyses	√	√	√	√	√	√	√	√	√	√	√
16.multivariable	√	√	√				√	√		√	√

analyses											
17. effects and confidence intervals	√	√	√		√		√	√	√	√	√
18. further investigations						√					√
Discussion											
19. interpret the results	√	√	√	√	√	√	√	√	√	√	√
20. implications for future research	√	√		√	√	√		√	√	√	√

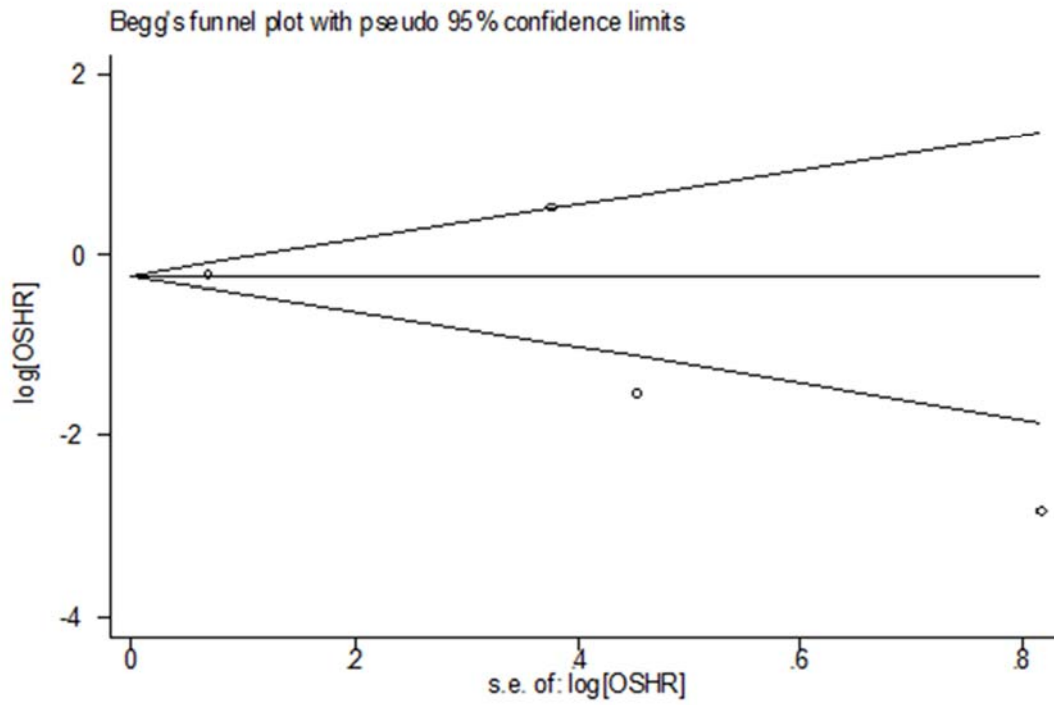
Supplementary Figure 1. Forest plot showing the combined HR from included studies for the association between the expression levels of miR-200 family and OS. OS = overall survival.



Supplementary Figure 2. Sensitivity analyses of studies regarding miR-200 family expression and OS. OS = overall survival.

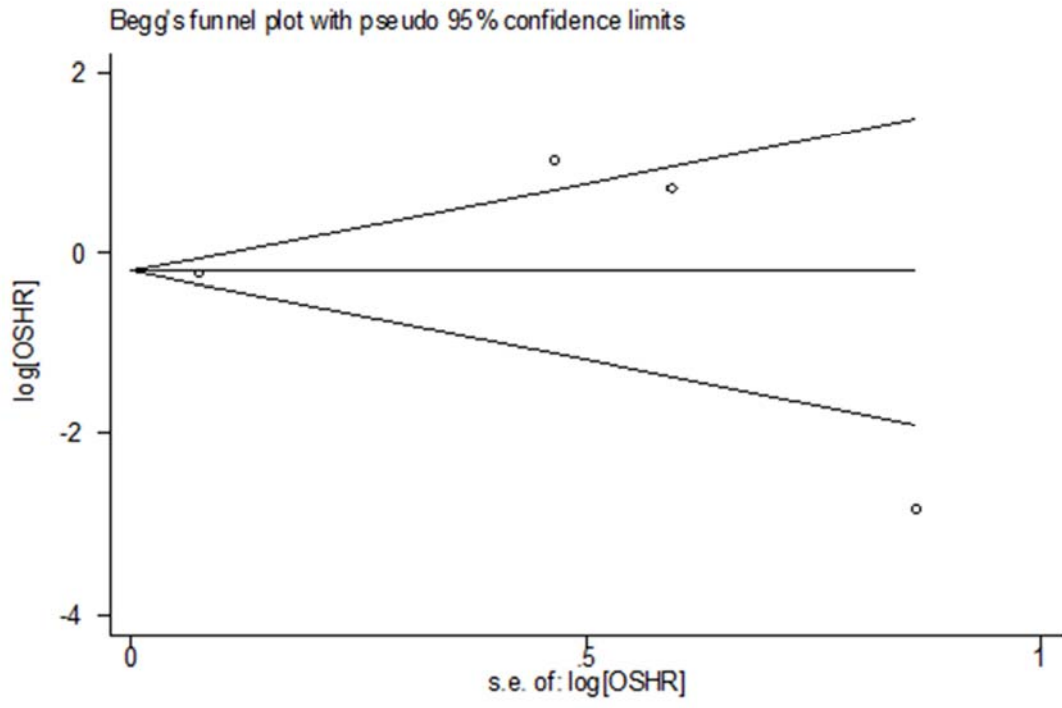


(A)

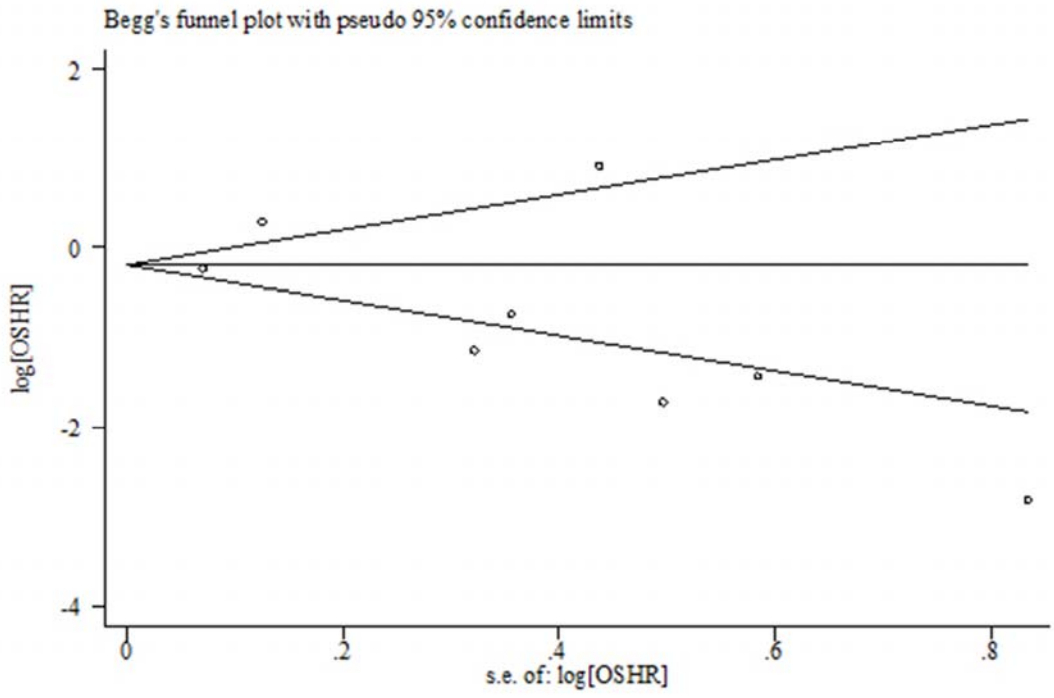


Supplementary Figure 3. Begg's funnel plot of standard error for assessing publication bias. A, miR-200a. B, miR-200b. C, miR-200c. D, miR-141.

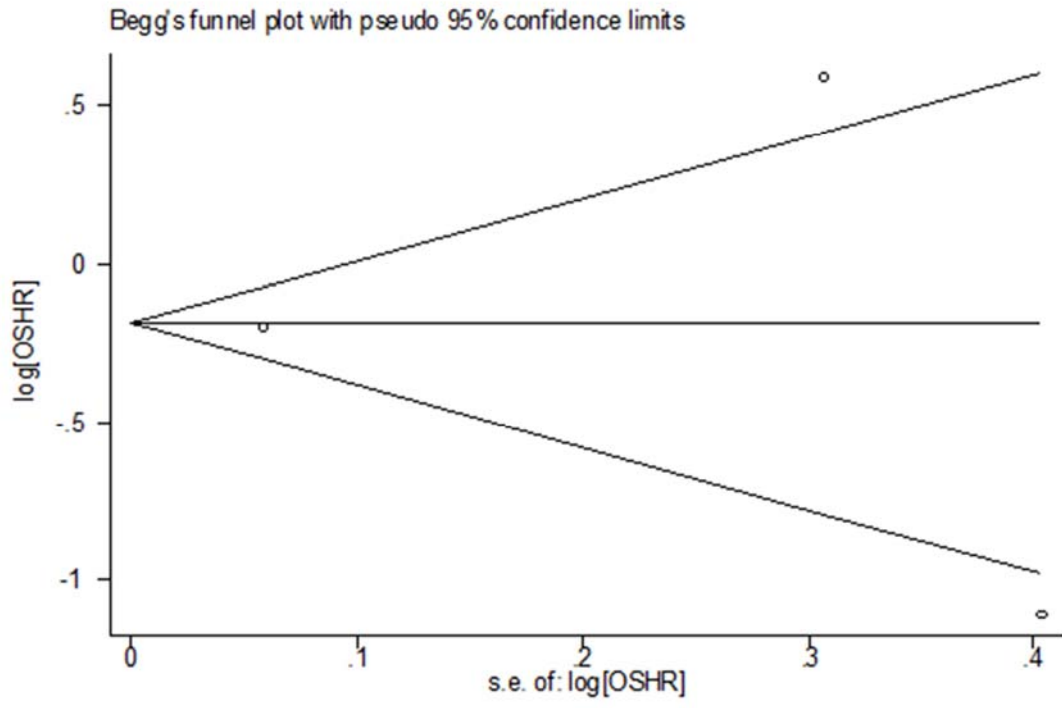
(B)



(C)



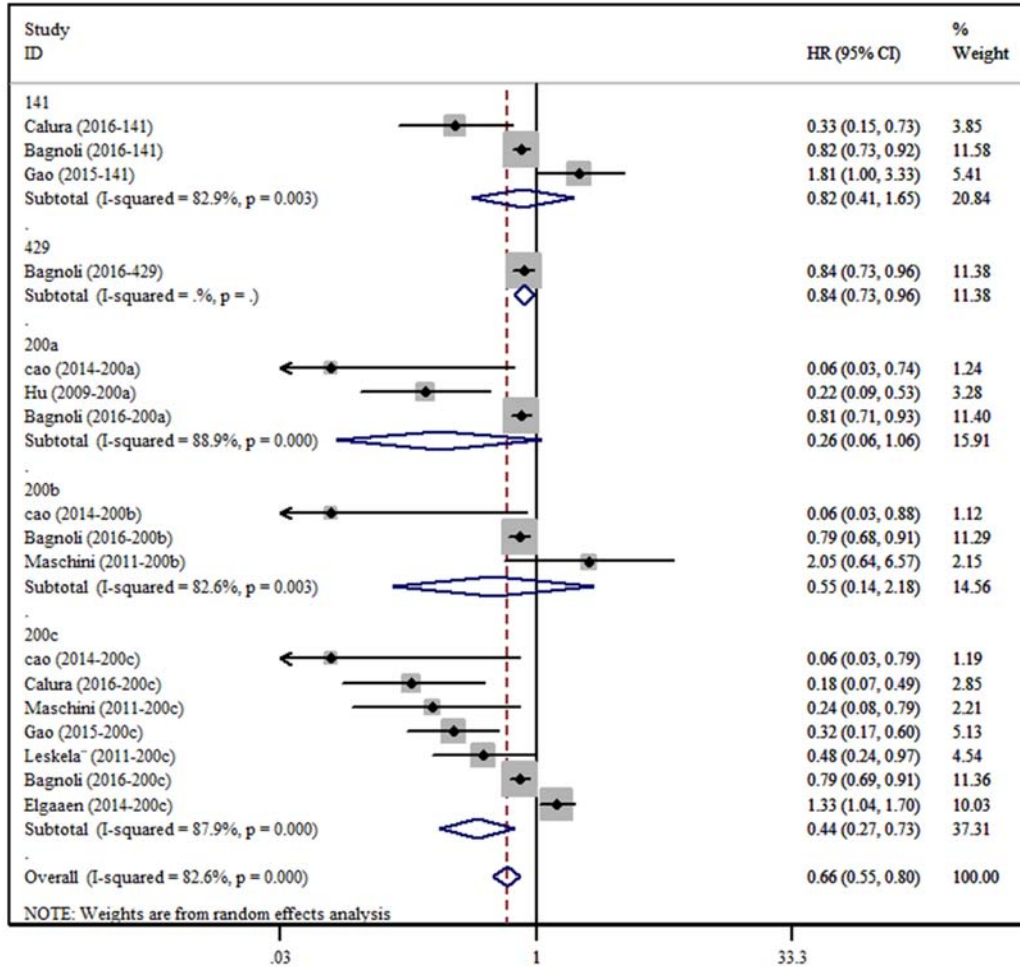
(D)



Supplementary Figure 3. Begg's funnel plot of standard error for assessing publication bias. A, miR-200a. B, miR-200b. C, miR-200c. D, miR-141.

Supplementary Figure 4. Forest plots of subgroup analysis by study samples regarding the miR-200 family expression and OS of patients with ovarian cancer. A, the study samples were tissues. B, the study samples were blood.

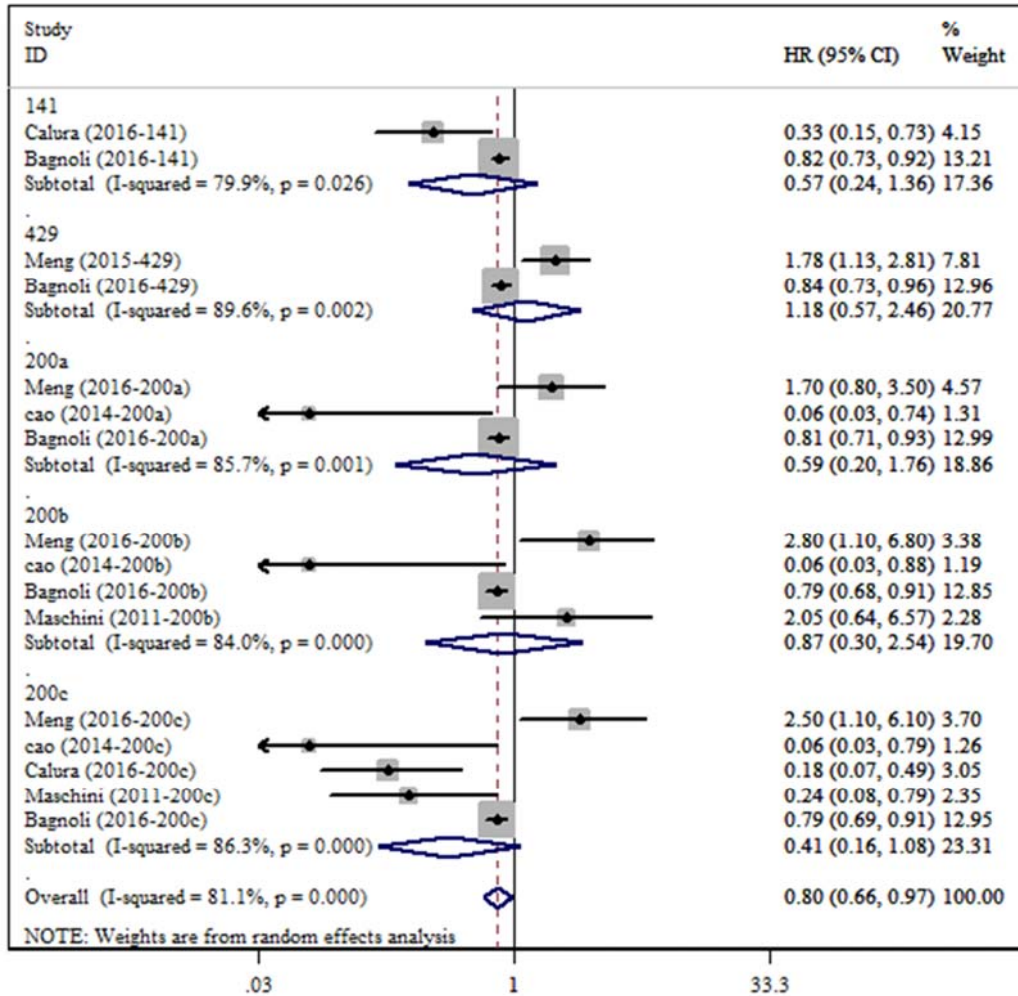
(A)



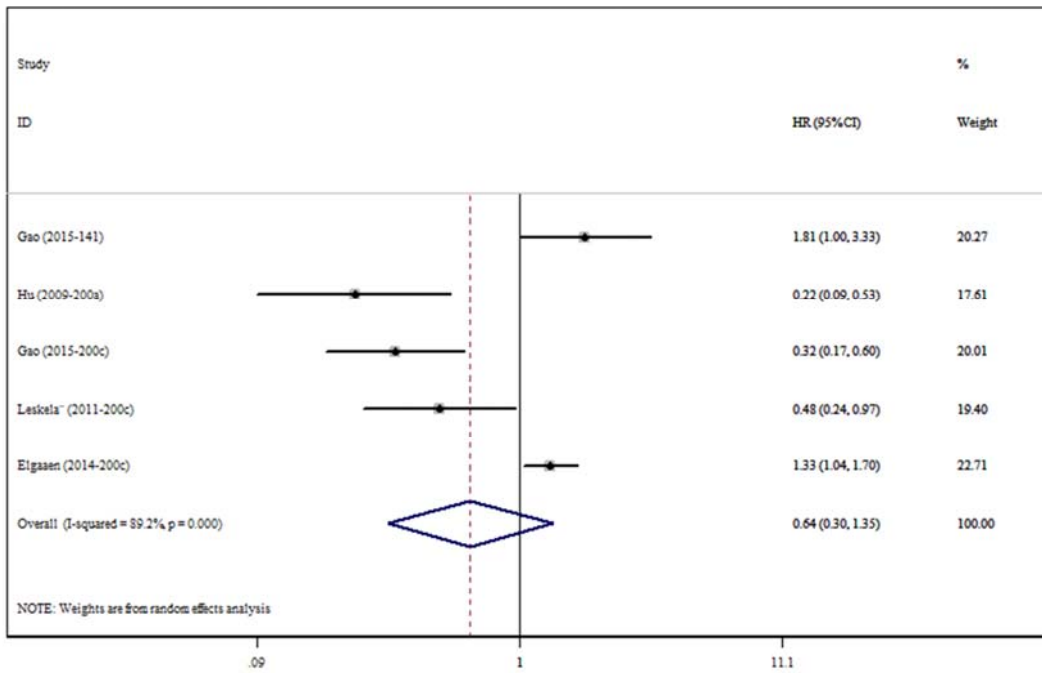
(B)

Supplementary Figure 5. Forest plots of subgroup analysis by study kinds regarding the miR-200 family expression and OS of patients with ovarian cancer. A, the study kinds were multivariate study. B, the study kinds were univariate study. C, divided miR-200 family into two clusters based on chromosomal location. OS = overall survival.

(A)

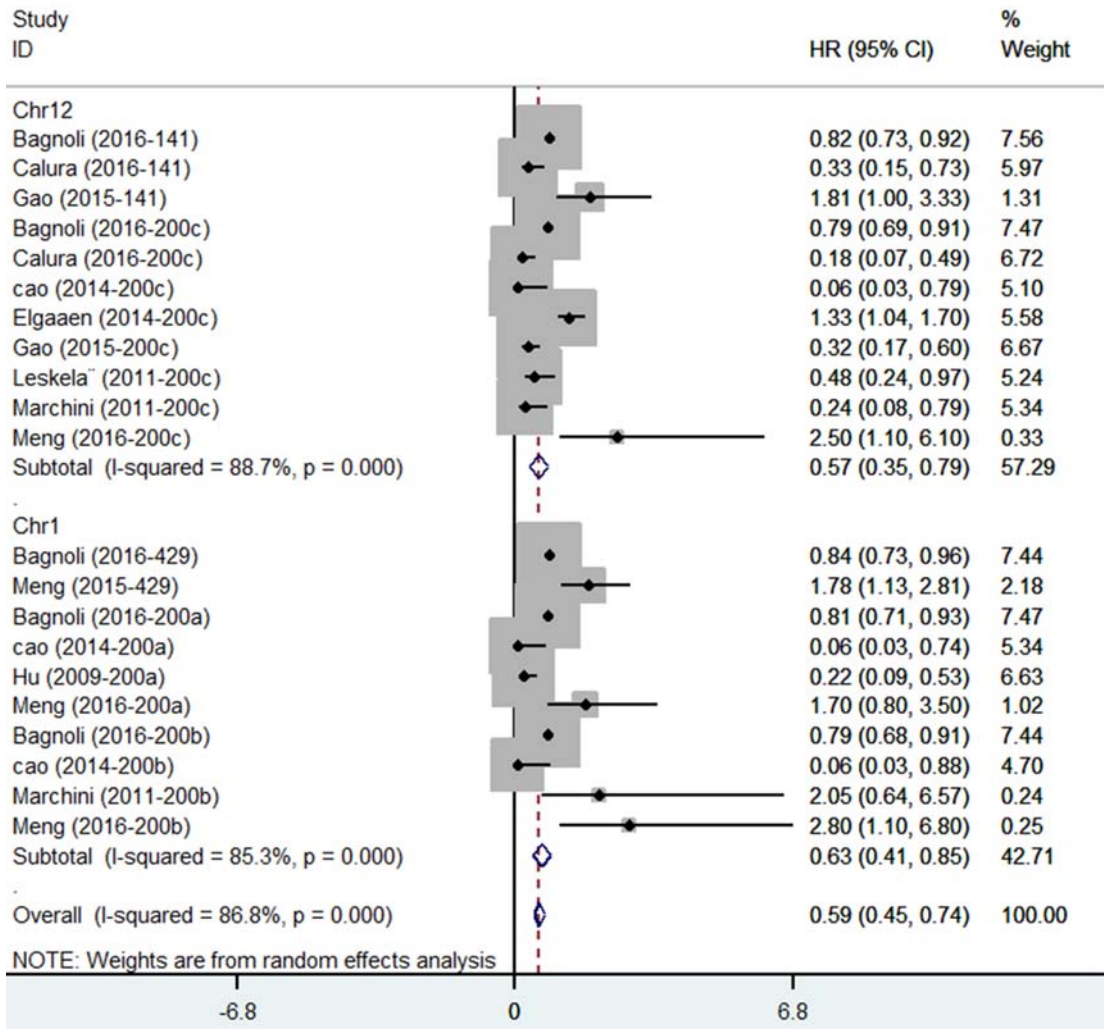


(B)



Supplementary Figure 5. Forest plots of subgroup analysis by study kinds regarding the miR-200 family expression and OS of patients with ovarian cancer. A, the study kinds were multivariate study. B, the study kinds were univariate study. C, divided miR-200 family into two clusters based on chromosomal location. OS = overall survival.

(C)



Supplementary Figure 6. Forest plot showing the combined HR from included studies for the association between the expression levels of miR-200 family and PFS. PFS = progression free survival.

