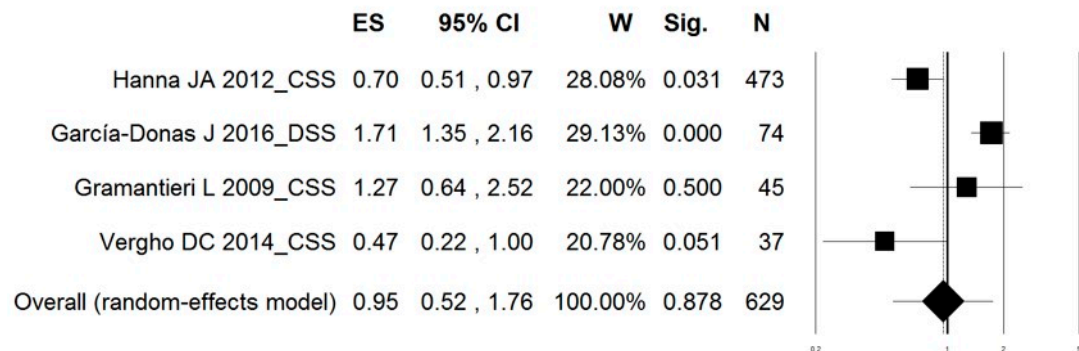
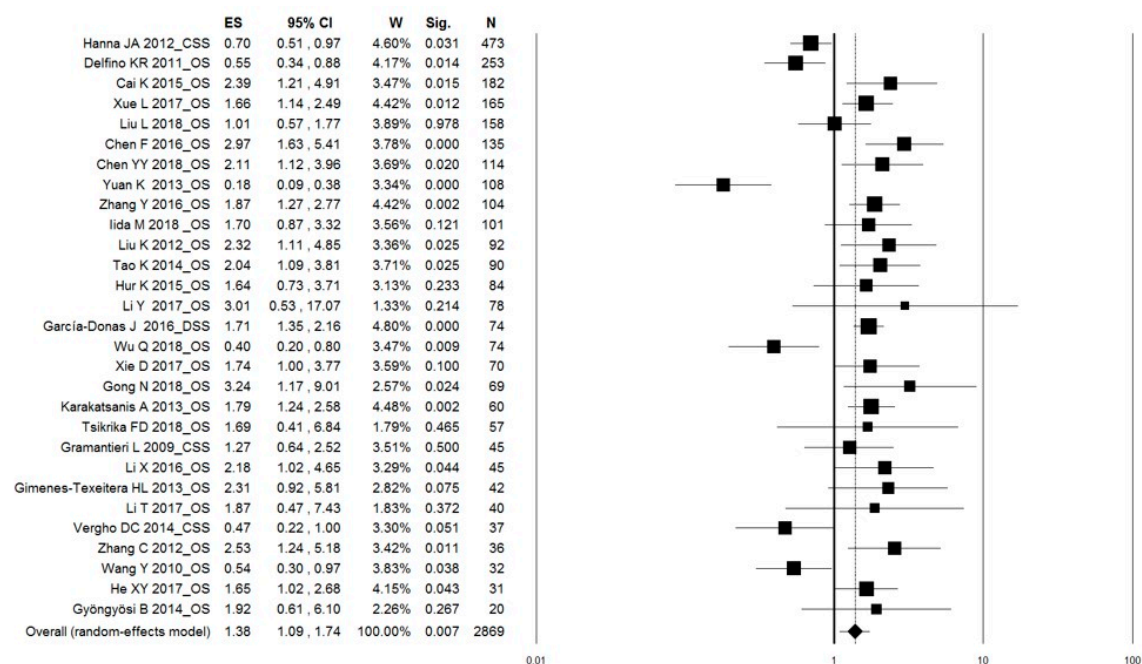


# Supplementary Materials: Prognostic role of miR-221 and miR-222 expression in cancer patients: A systematic review and meta-analysis

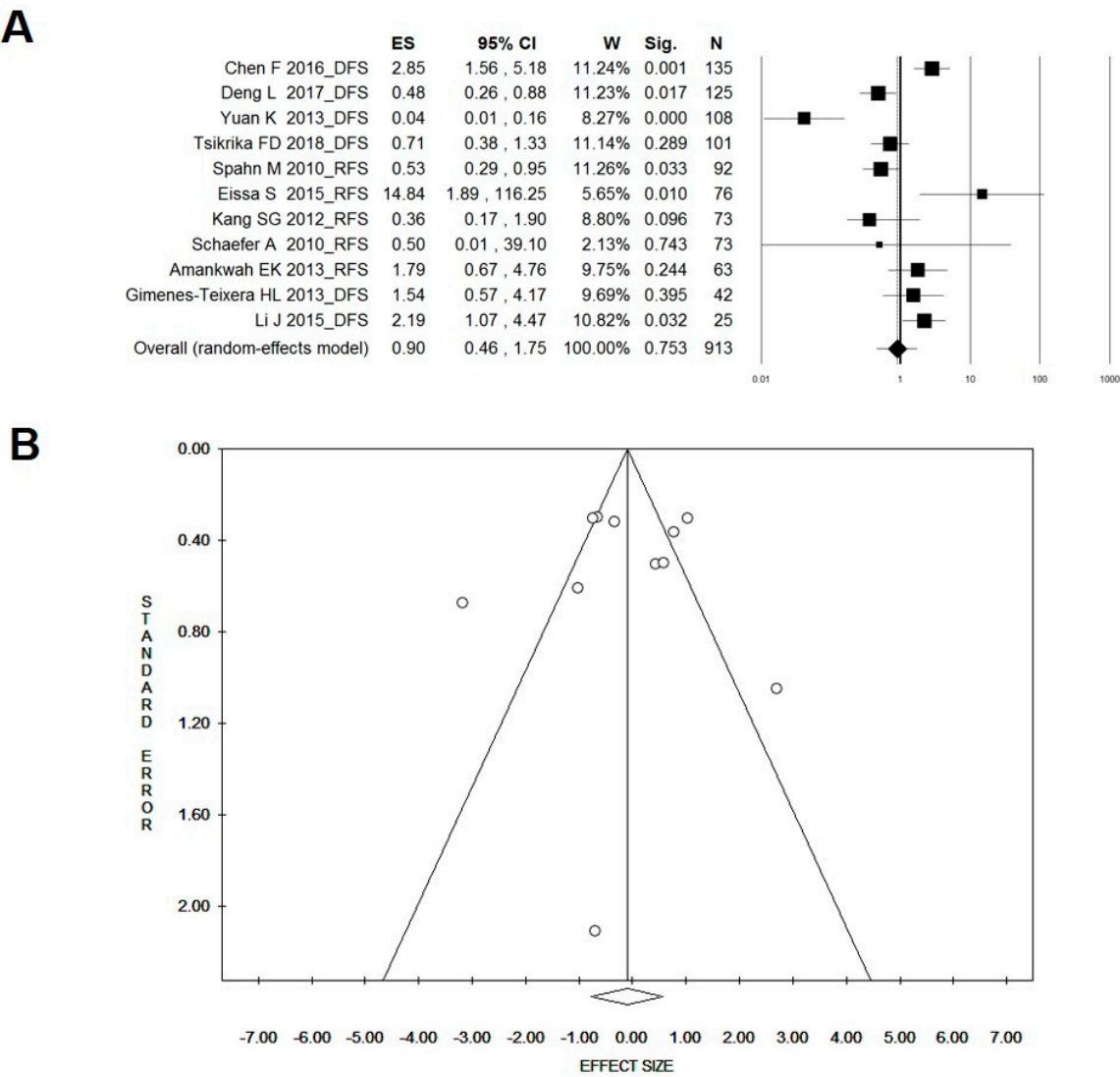
Gloria Ravegnini, Sarah Cargnin, Giulia Sammarini, Federica Zanotti, Justo Lorenzo Bermejo, Patrizia Hrelia, Salvatore Terrazzino and Sabrina Angelini



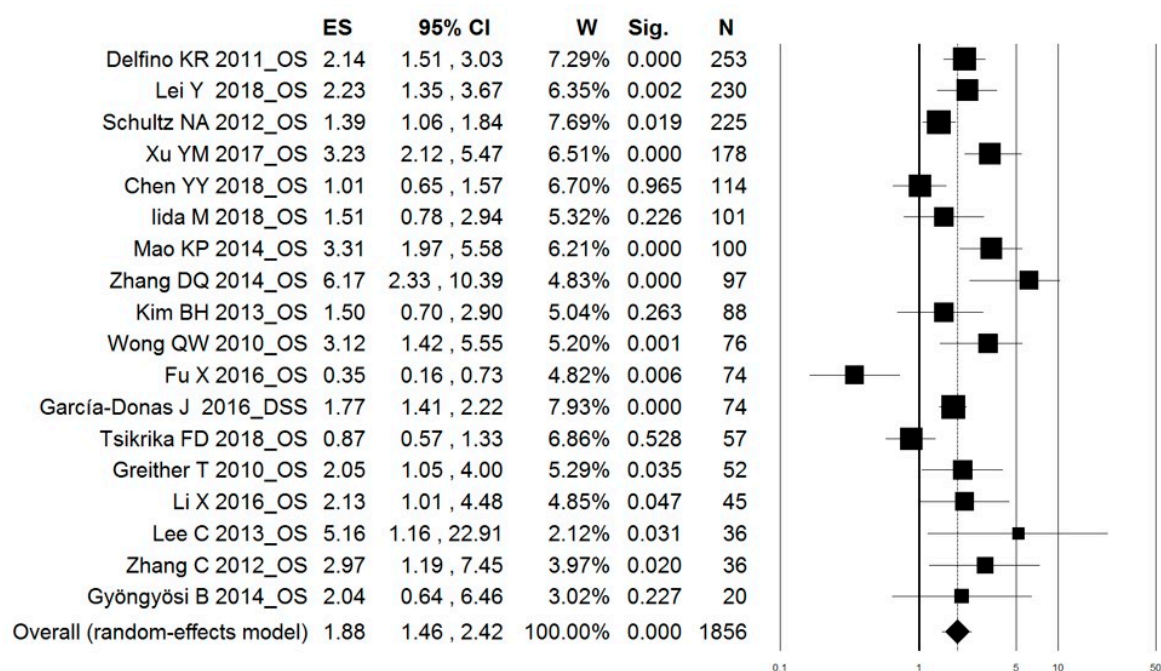
**Figure S1.** Forest plots for the effect of miR-221 overexpression and CSS and DSS



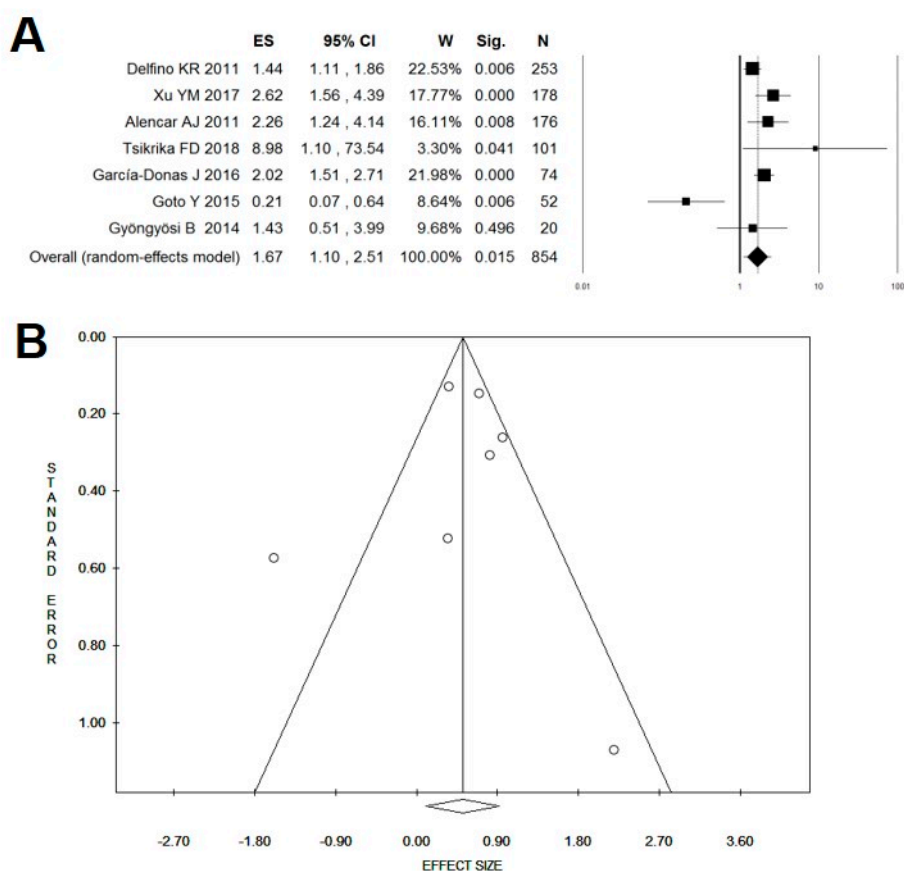
**Figure S2.** Forest plots for the effect of miR-221 overexpression and OS/CSS/DSS



**Figure S3.** Forest (A) and Funnel (B) plots for the effect of miR-221 overexpression on DFS/RFS. ES, effect size (i.e., hazard ratio); W, weight; Sig, statistical significance; N, total number of cancer patients included in the survival analysis.



**Figure S4.** Forest plots for the effect of miR-222 overexpression and OS/DSS



**Figure S5.** Forest (A) and Funnel (B) plots for the effect of miR-222 overexpression on progression-free survival. ES, effect size (i.e., hazard ratio); W, weight; Sig, statistical significance; N, total number of cancer patients included in the survival analysis.

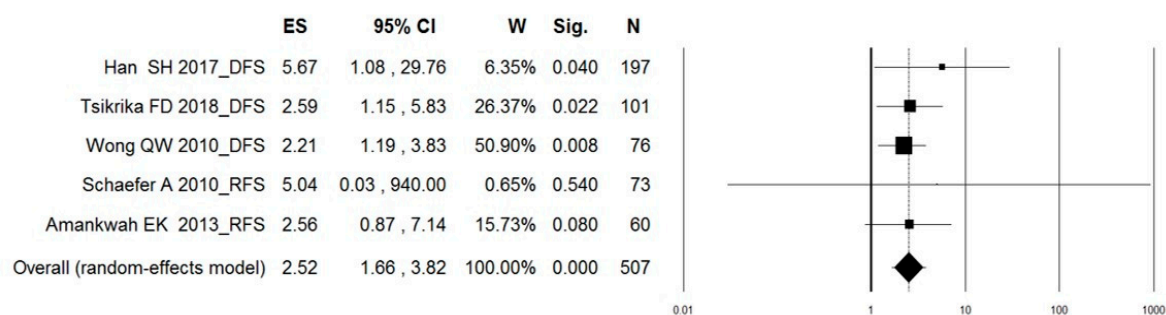


Figure S6. Forest plots for the effect of miR-222 overexpression and RFS/DFS

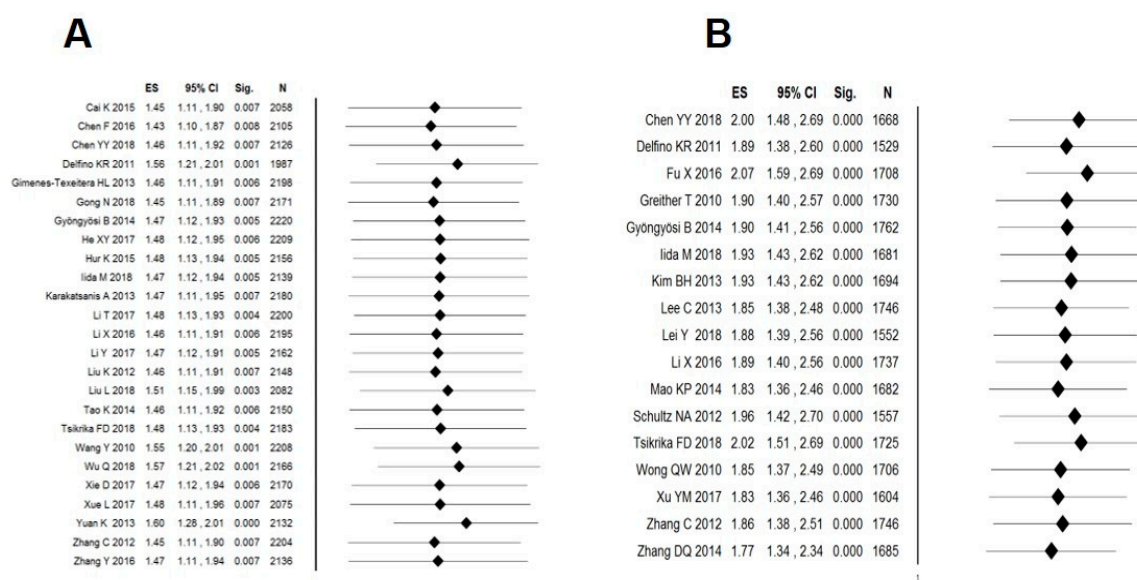


Figure S7. Sensitivity analysis of the meta-analysis for miR-221 (A) and miR-222 (B)



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