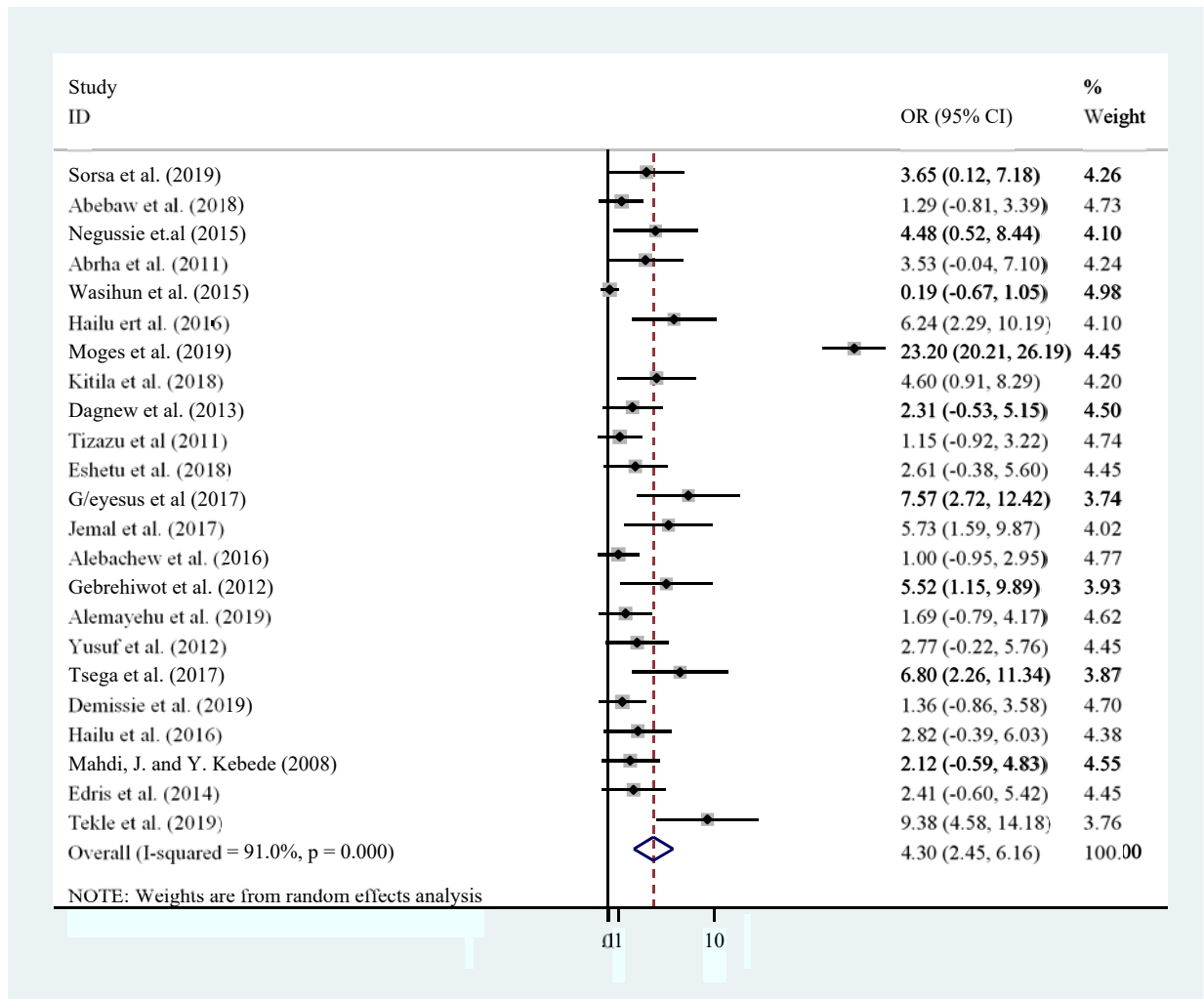
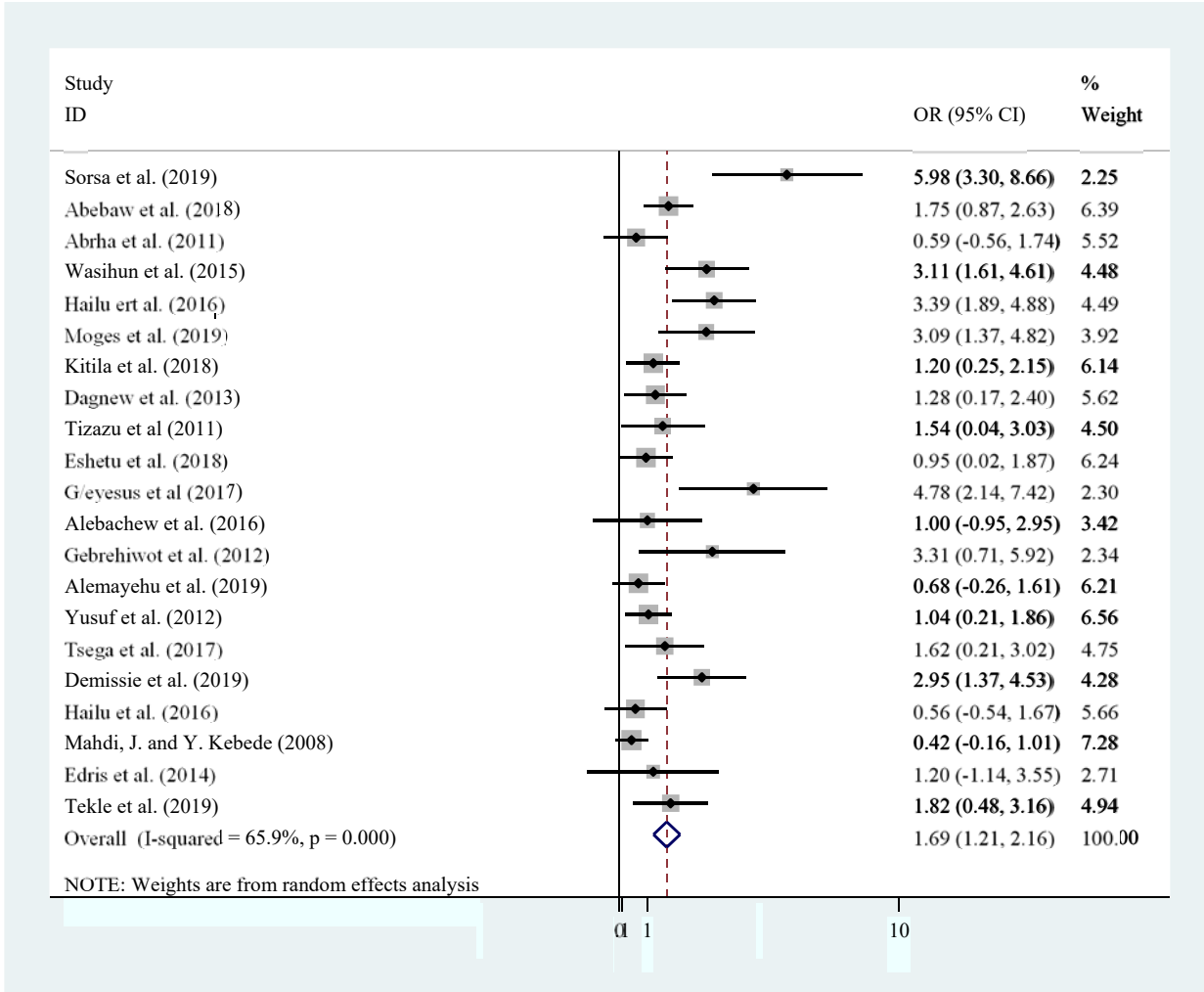


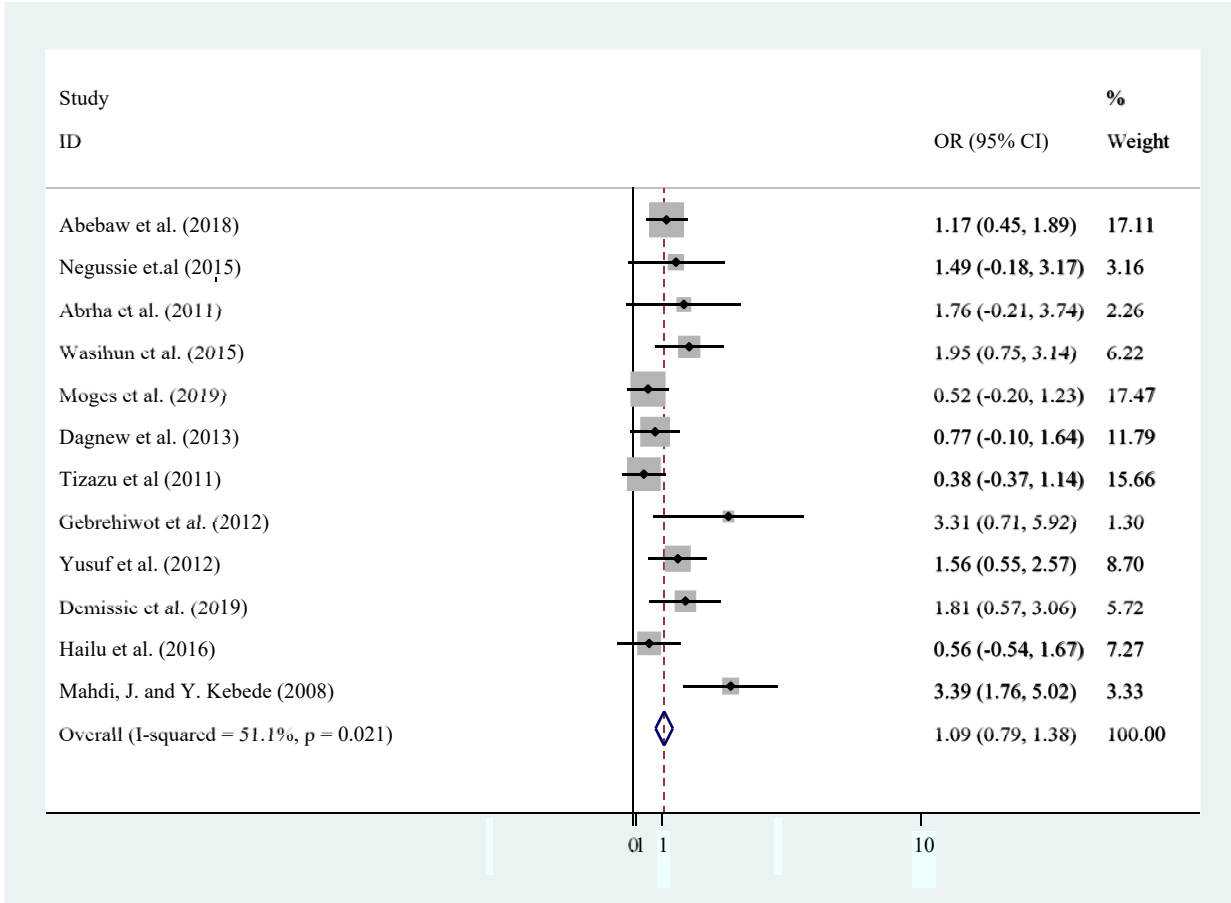
## Supplementary figures



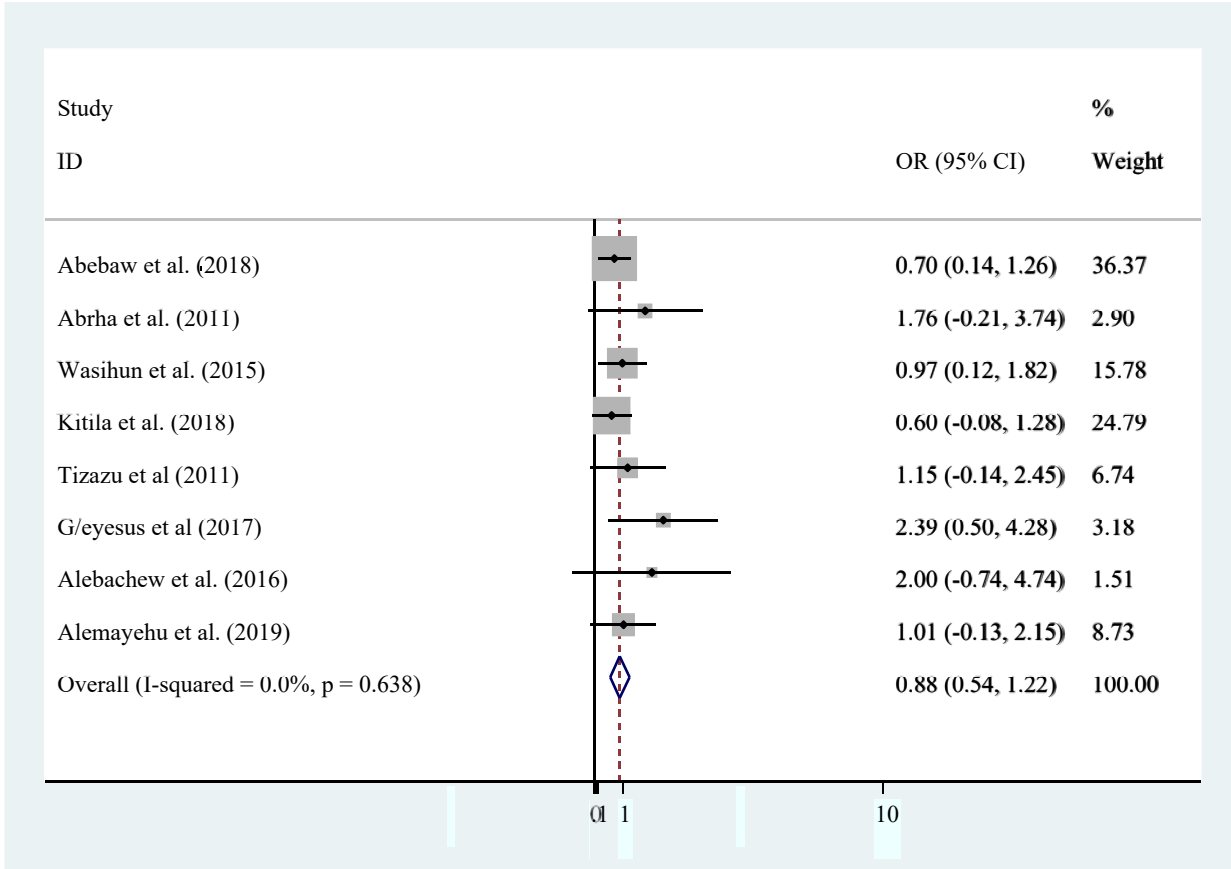
**Fig.S1:** Forest plot showed the pooled prevalence of Klebsiella species isolated among patients with suspected bloodstream infections in Ethiopia



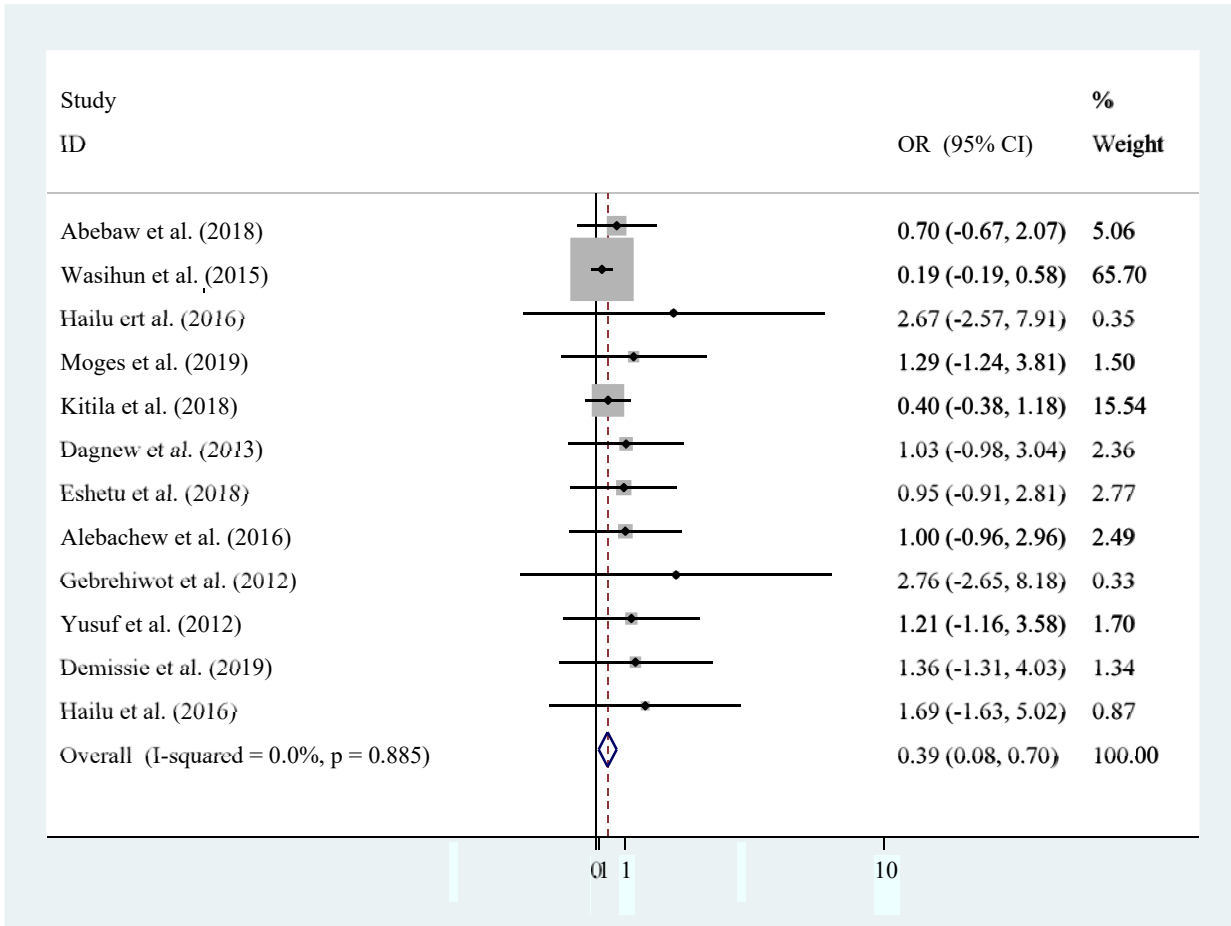
**Fig.S2:** Forest plot showed the pooled prevalence of *E.Coli* isolated among patients with suspected bloodstream infections in Ethiopia



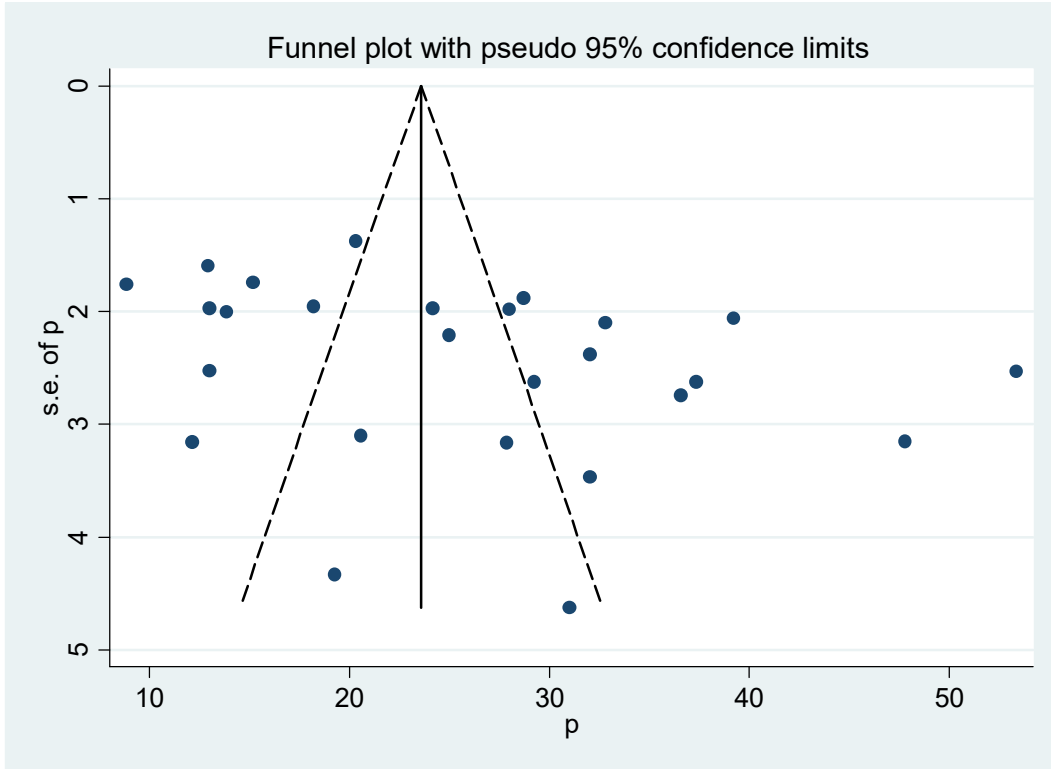
**Fig.S3:** Forest plot showed the pooled prevalence of *Salmonella Species*-isolated among patients with suspected bloodstream infections in Ethiopia



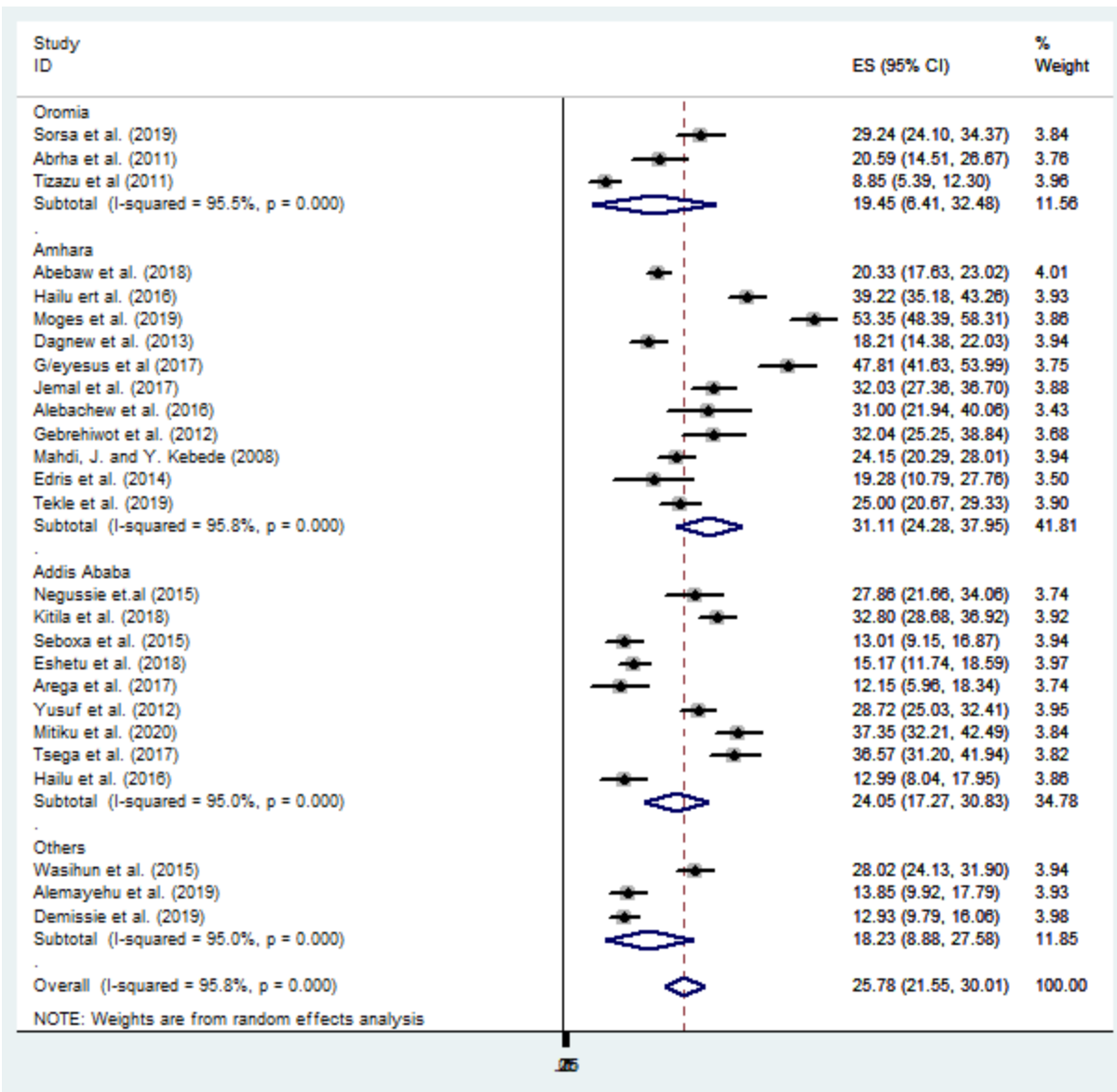
**Fig. S4:** Forest plot showed the pooled prevalence of *S.pyogenes* isolated among patients with suspected bloodstream infections in Ethiopia



**Fig. S5:** Forest plot showed the pooled prevalence of *Pseudomonas* Species isolated among patients with suspected bloodstream infections in Ethiopia

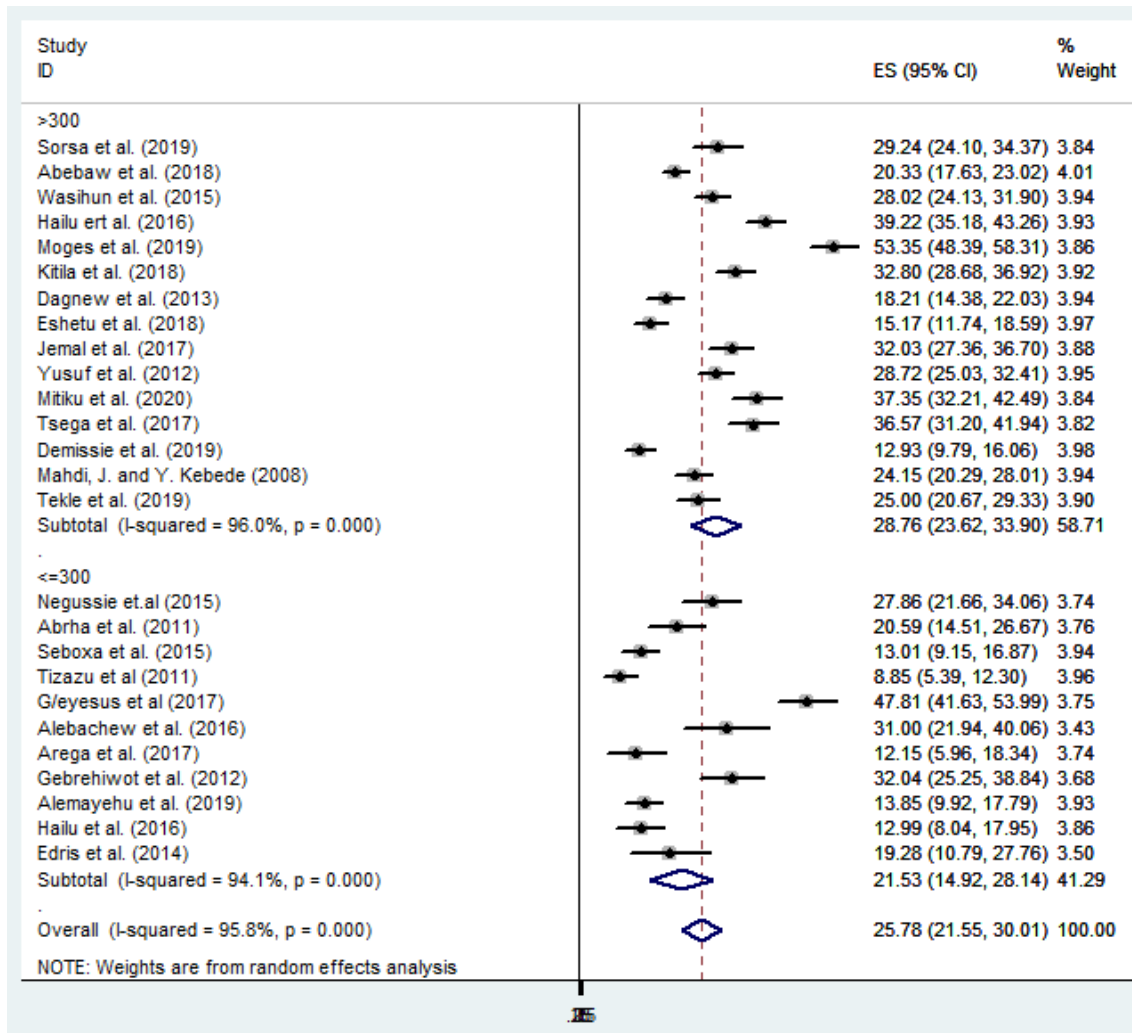


**Fig. S6:** funnel plots showing publication bias of included studies, p (prevalence) of bacterial isolates represented in the x-axis and SE (standard error) in the y-axis



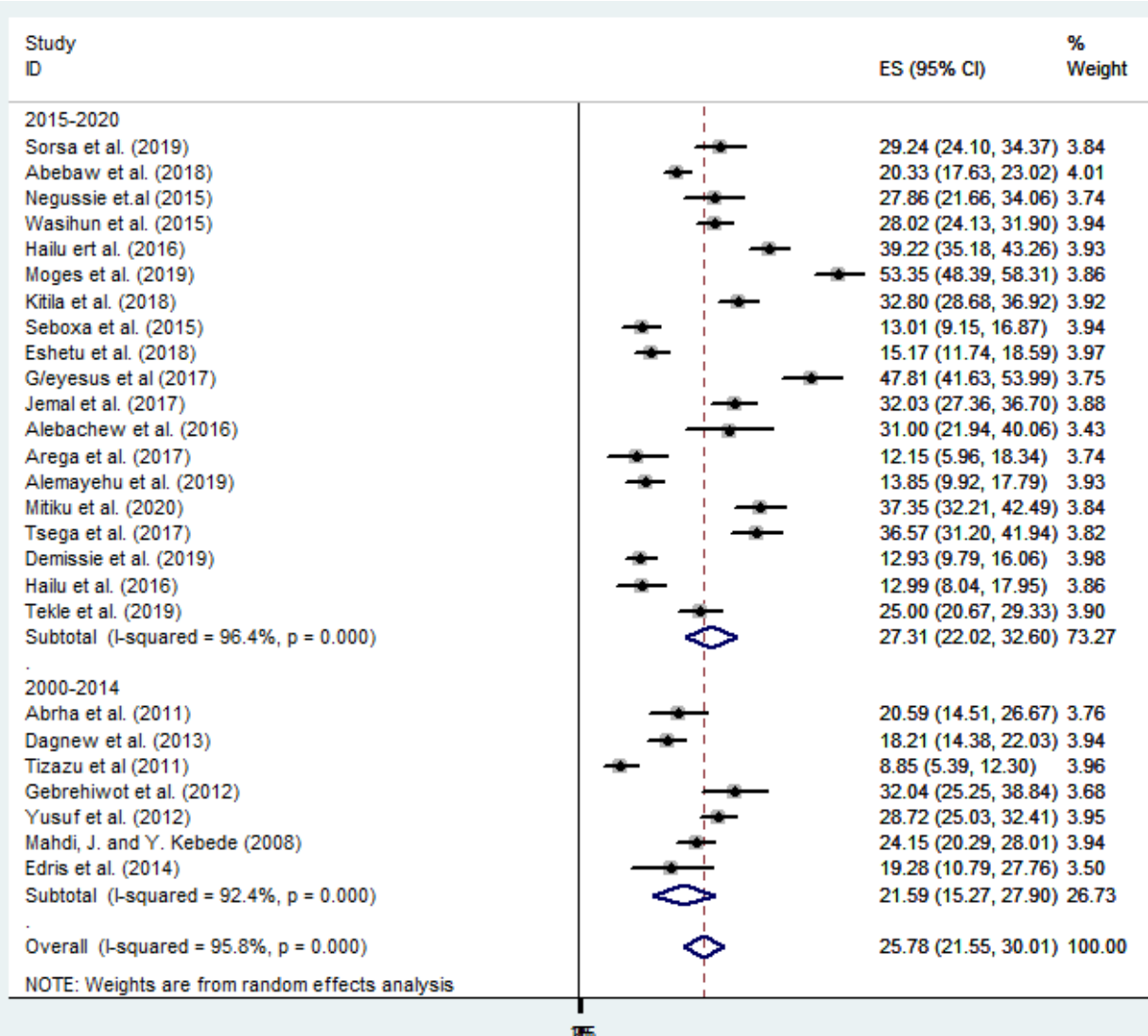
NB: Others: South nation nationality people, Tigray, Dire Dawa

**Fig. S7.** Subgroup analysis based on study area/region for the pooled prevalence of bacterial isolates causing blood stream infection in Ethiopia, 2020

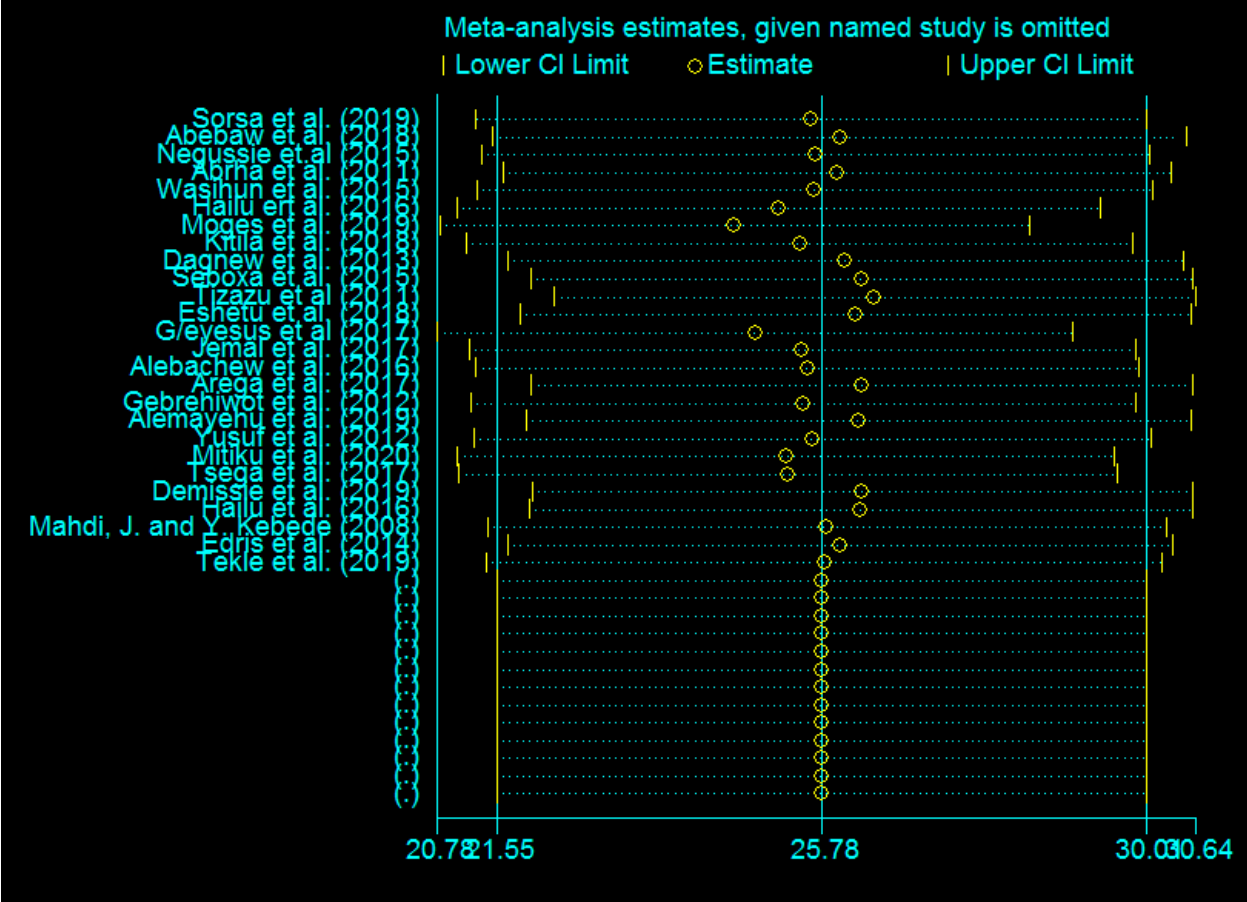


**Fig S8.** Subgroup analysis based on sample size for the pooled prevalence of bacterial isolates causing blood stream infection in Ethiopia, 2020.





**Fig. S9.** Subgroup analysis based on year publication for the pooled prevalence of bacterial isolates causing blood stream infection in Ethiopia, 2020



**Fig. S10:** Sensitivity analysis pooled prevalence pooled prevalence of bacterial isolates causing blood stream infection in Ethiopia