

## Supplementary Figures

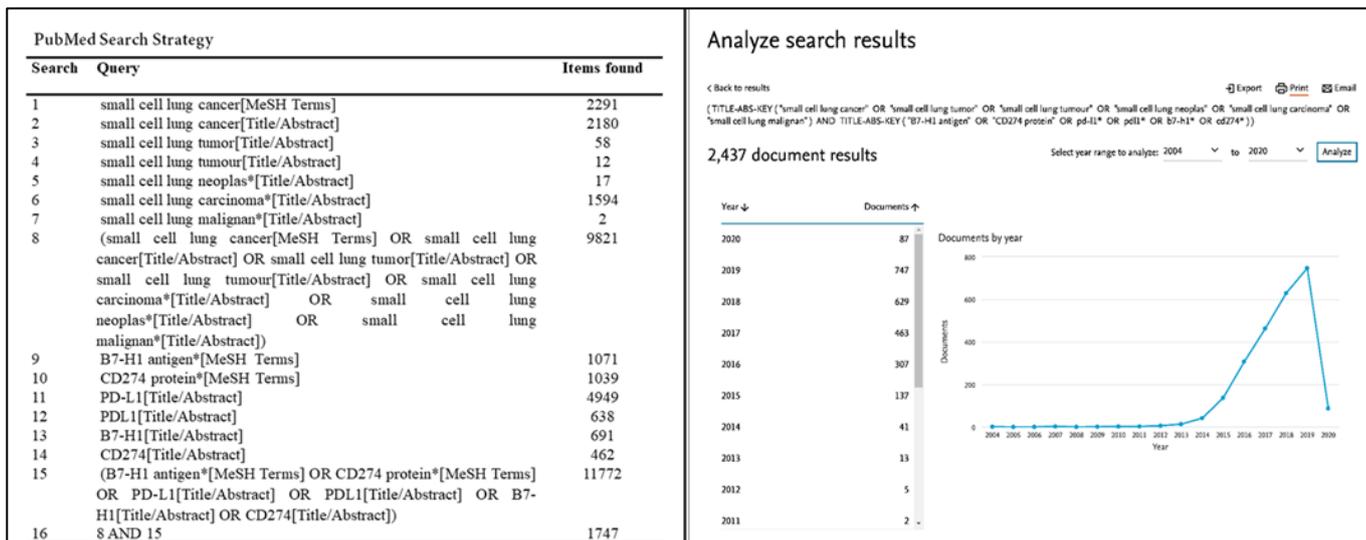


Figure S1: Search strategy -PubMed/Scopus

Author (year)	Representative sample of relevant population	Selection of the non-exposed cohort	Ascertainment of exposure	Demonstration that outcome of study does not present at the start of the study	Design of analysis controlled for cofounders	Assessment of outcome	Follow-upadequate for outcomes to occur	Quality Score
Ishii et al., 2015	*	*	*	**				5
Komiya and Madan, 2015		*	*	**				4
Schultheis et al., 2015		*	*	**		*		6
Yu et al., 2016	*	*	*	**	*	*	*	8
Takada et al., 2016		*	*	**		*		5
Antonia et al 2016	**	*	**	**	**	**	*	12
Berghoff et al., 2016	*		*	**		**	*	7
Miao et al., 2017		*	*	**		*	*	6
Chang et al., 2017	*		*	**		**	*	7
Inamura et al., 2017		*	*	**		*	*	6
Tsuruoka et al., 2017		*	*	**	*	**	*	8
Ott et al, 2017	*	*	**	**	**	**	**	12
Yasuda et al, 2018		*		**		*		4
Bonanno et al., 2018	*	*	*	**	*	*		7
Wang et al., 2018		*	*	**	*	*	*	6
Ichiki et al., 2018		*	*	**		**		7
Liu et al., 2018		*	*	**	*	**	*	8
Gadgeel et al, 2018		*	**	**	**	**	**	11
Chung et al., 2018	*	*	*	**	**	*	*	8
Kim et al., 2018	*	*	*	**	**	*	*	8
Xu et al., 2019		*	*	**		**	*	7
Zhao et al., 2019	*	*	*	**		**	*	9
Carvajal-Hausdorf et al., 2019		*	*	**	*	*		6
Pujol et al., 2019	**	*	**	**	**	**	**	13
Sun et al., 2019		*	*	**	*	**	*	8
Pas-Ares et al., 2019	**	*	**	**	**	**	**	13
Recket al., 2020	*	*	**	**	**	**	**	12

**Figure S2: Assessment of the quality of methodological characteristics on included studies utilising the Newcastle-Ottawa Scale (NOS) scale**  
**\*\*=criterion is satisfied; \*=criterion is partly satisfied; no \* =criterion is not satisfied or cannot be determined. Maximum quality score=16; 0–5 points were considered low quality, 5–9 points were considered medium quality and 10–16 points were considered as high quality.**

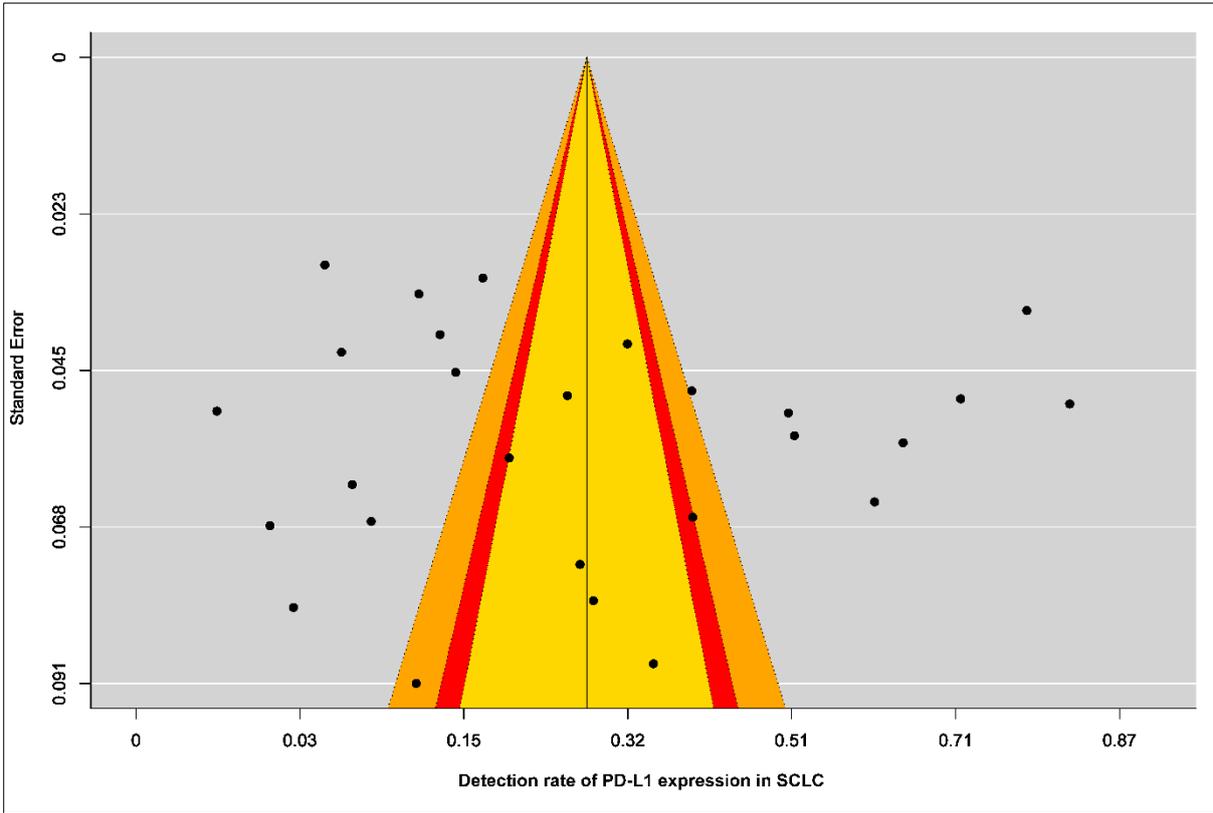
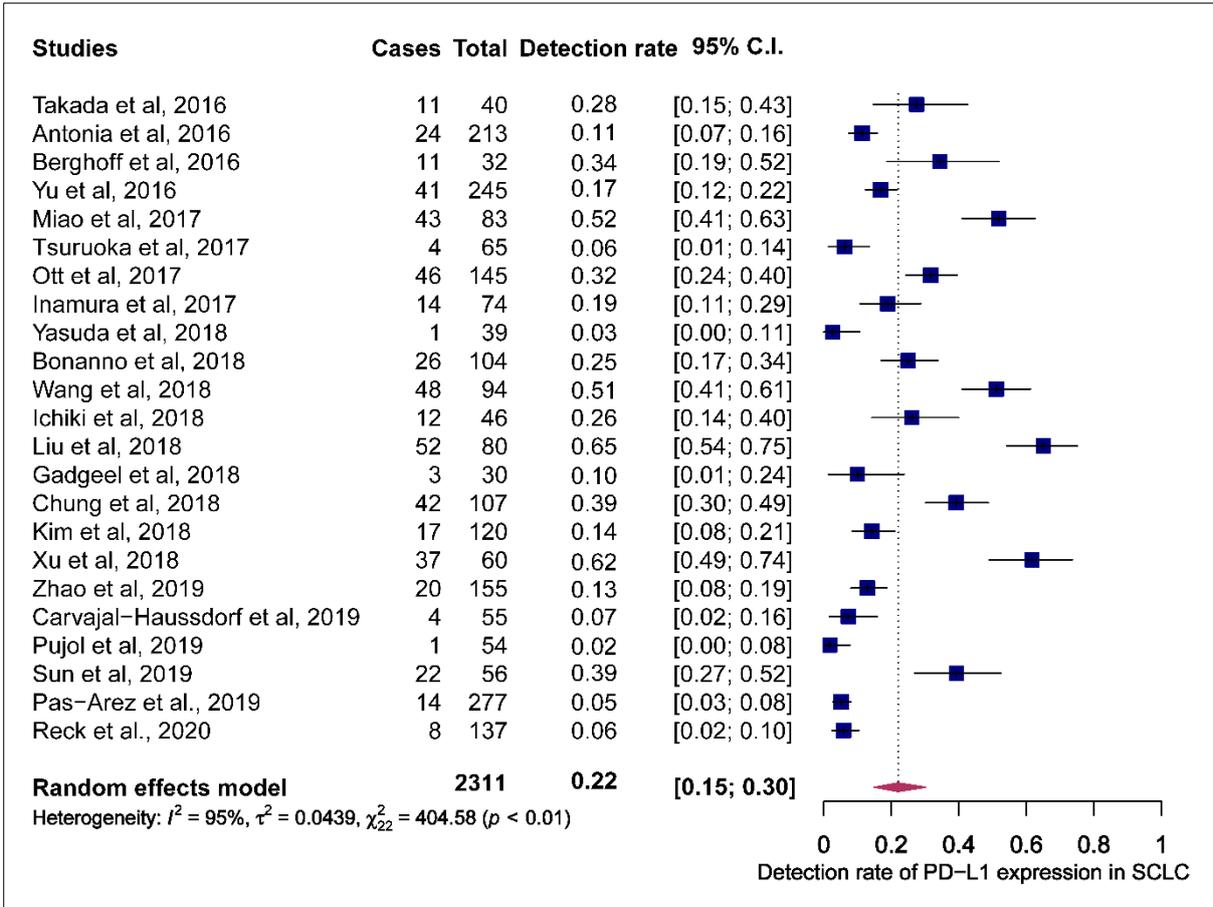


Figure S3: Begg's funnel plots with 90% CI (gold), 95% CI (red) and 99% CI (orange) for publication bias testing for the prevalence of PD-L1 expression in SCLC.



**Figure S4: Forest plot of studies reporting the detection rate of PD-L1 expression in SCLC after the leave-one-out sensitivity analysis to remove outliers. The PD-L1 detection rates and 95%CI of each study is represented with a horizontal line and the square area mirrors the weight of the study. A random-effect model was utilised.**

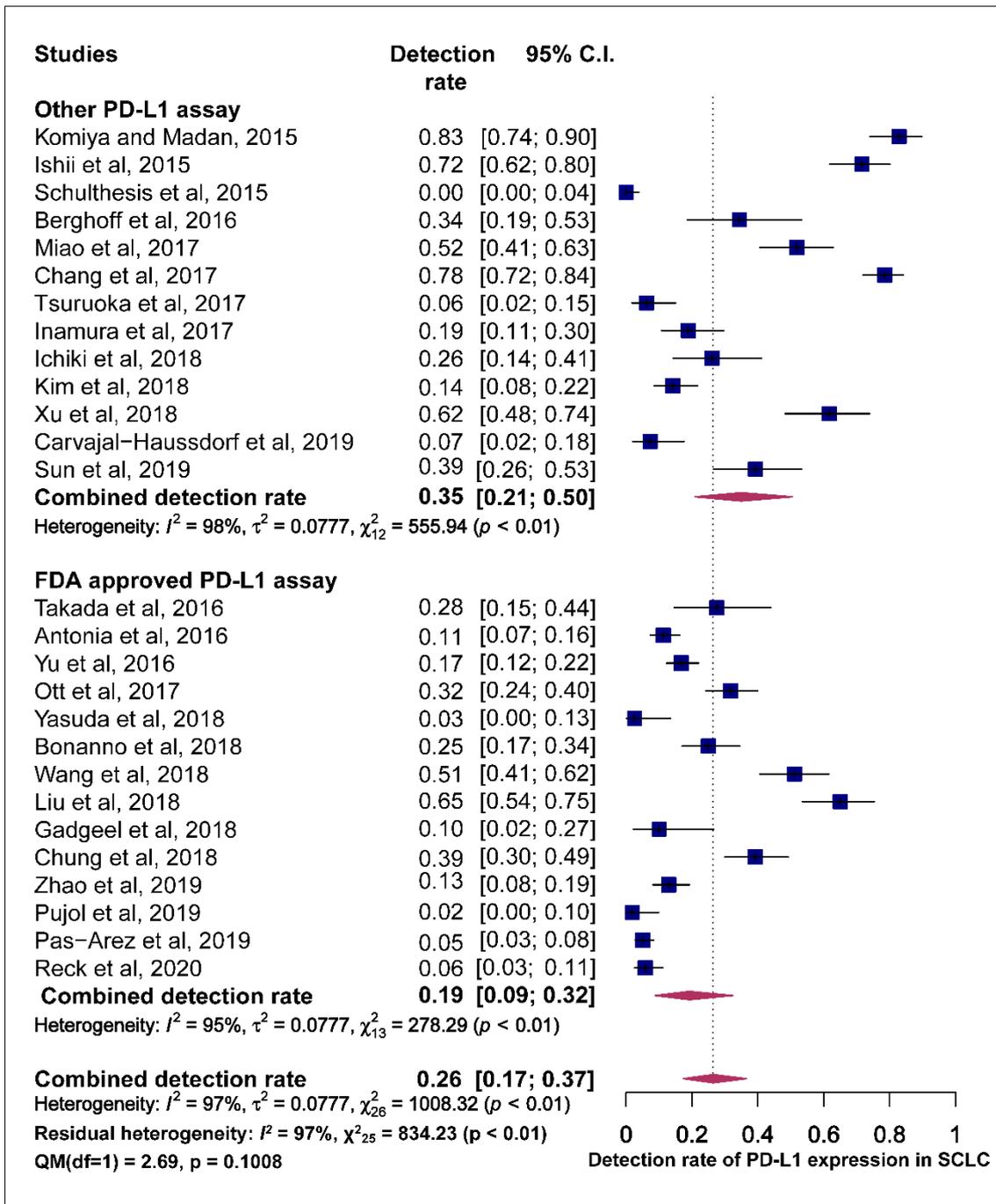


Figure S5: Forest plot of subgroup analysis of association between the use of FDA approved PD-L1 assays or not and prevalence of PD-L1 expression. The PD-L1 detection rates and 95%CI of each study is represented with a horizontal line and the square area mirrors the size effect of each study. A random-effect model was utilised.

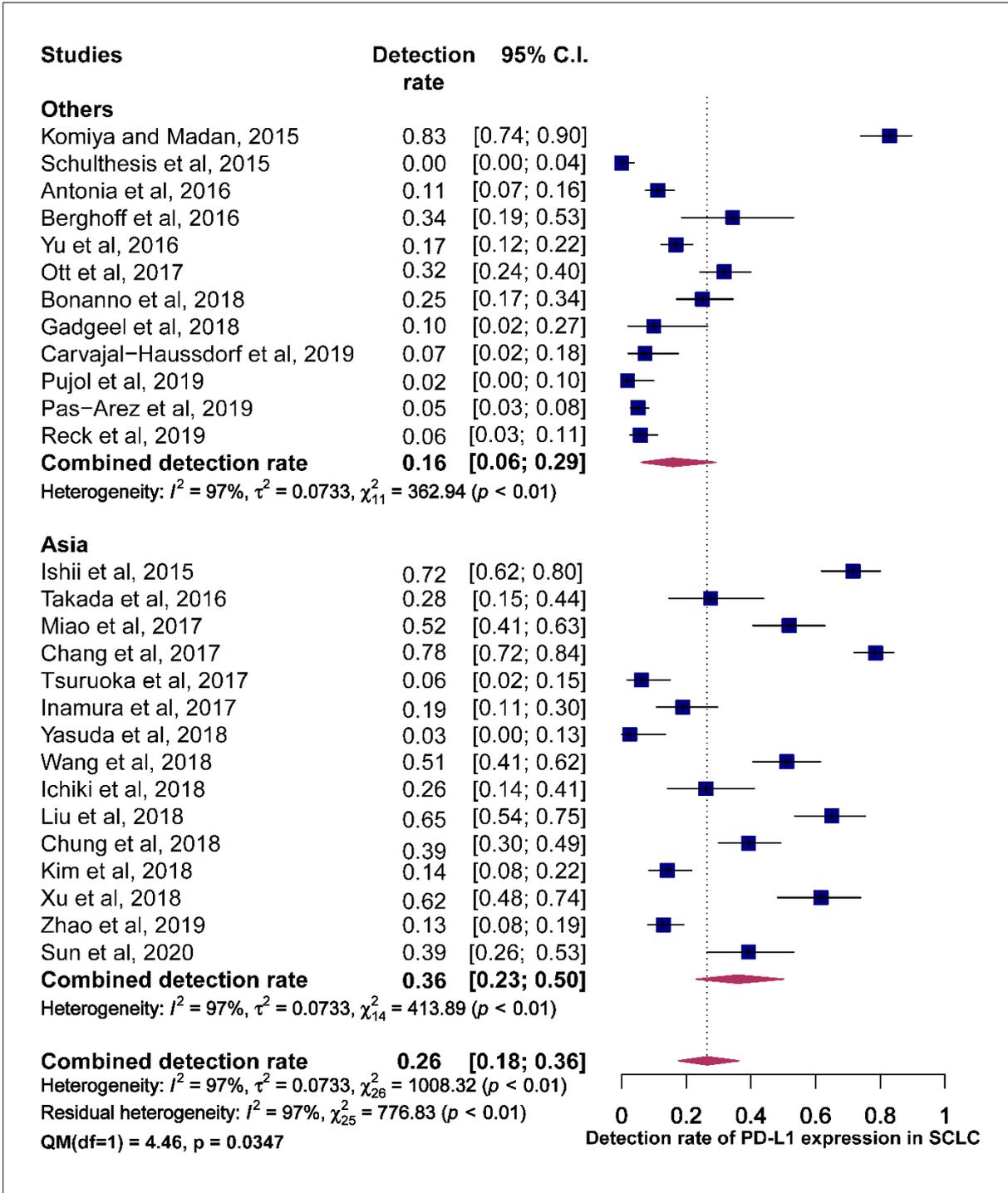
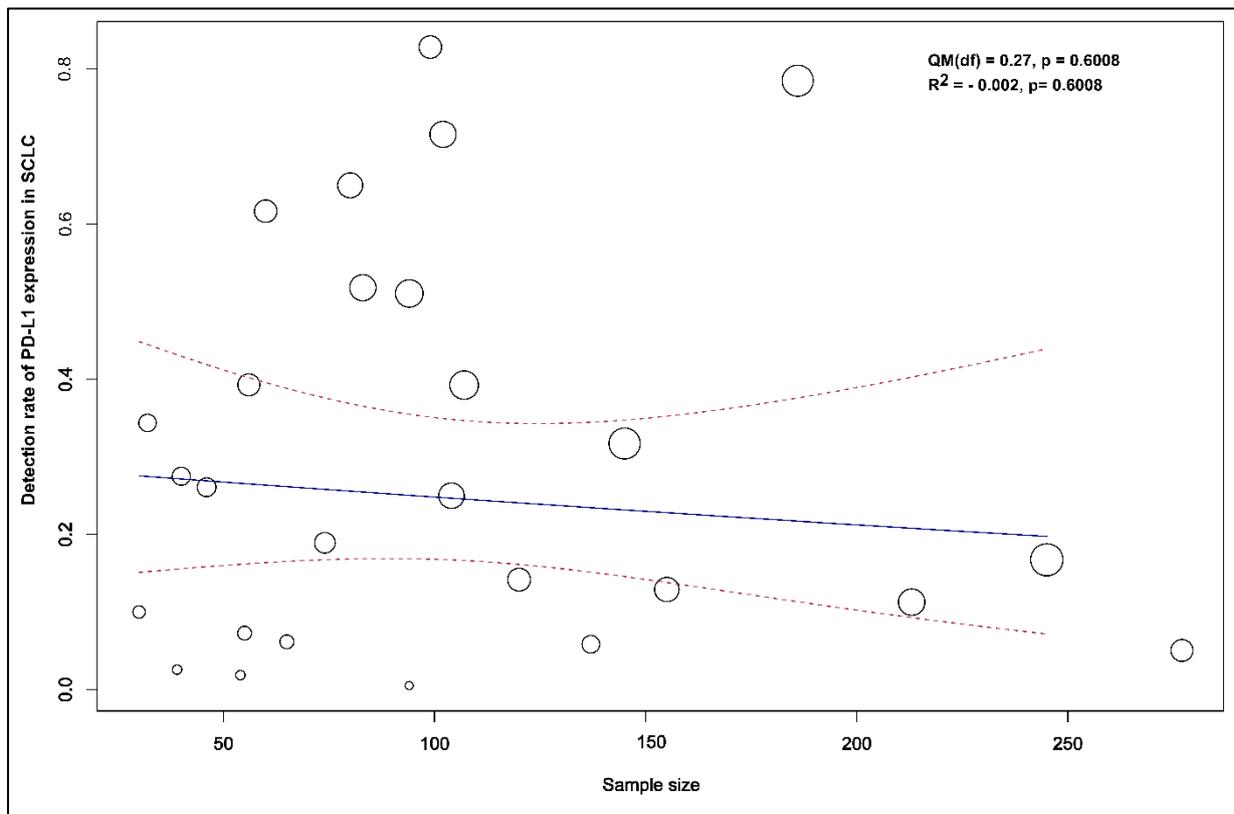


Figure S6: Forest plot of subgroup analysis of association between geographical distribution of published articles and prevalence of PD-L1 expression. The PD-L1 detection rates and 95%CI of each study is represented with a horizontal line and the square area mirrors the weight of the study. A random-effect model was utilised.



**Figure S7:** Meta-regression plot showing the association of sample size with pooled estimate of the prevalence of PD-L1 expression. Blue line represents the regression line of best fit, the red lines represent the lower and upper 95% confidence intervals (Cis), Each open circle represents a study. The size of the circle indicates the precision of the effect estimate and the weight given to that study

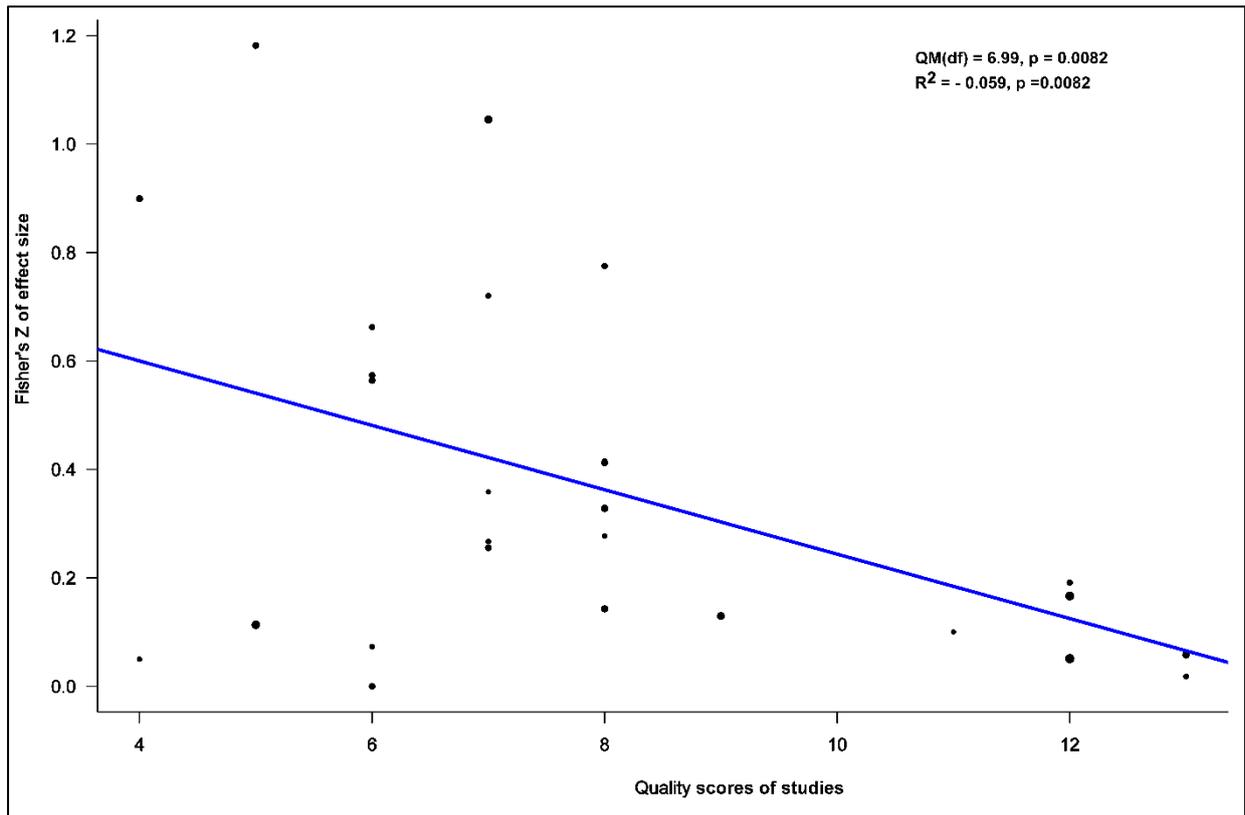


Figure S8: Meta-regression plot showing the association of quality scores of the study's methodological characteristics with pooled estimate of the prevalence of PD-L1 expression.

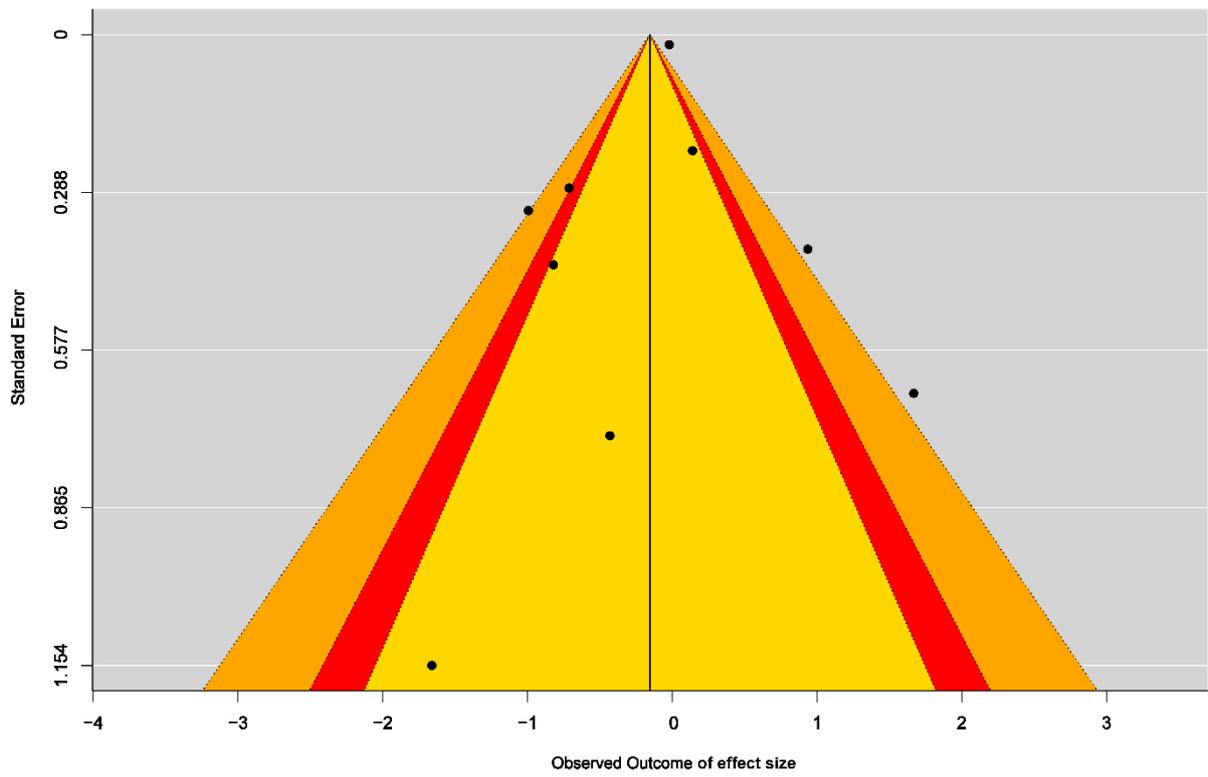


Figure S9: Begg's funnel plots with 90% CI (gold), 95% CI (red) and 99% CI (orange) for publication bias testing for the prognosis of PD-L1 expression in SCLC