# Prognostic value of immune checkpoint molecules in head and neck cancer: a meta-analysis

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#### ABSTRACT

Immune checkpoint molecules are important targets in cancer immunotherapy, but their association with prognosis in patients with head and neck cancer is controversial. In this meta-analysis, we searched for 12 immune checkpoint molecules in the PubMed, Embase and Cochrane Library databases and retrieved 52 studies with 7127 participants. Among the molecules included in the search, indoleamine 2, 3-dioxygenase (IDO), programmed death ligand 1 (PD-L1), and programmed death 1 (PD-1) met the inclusion criteria for further analysis. Higher expression of IDO was associated with poorer overall survival in head and neck cancer patients (P = 0.011), but higher expression of PD-L1 correlated with better overall survival specifically in nasopharyngeal carcinoma patients (P = 0.043), and was associated with better overall survival in Caucasian subjects (P = 0.02), nasopharyngeal carcinoma patients (P = 0.015), and studies with small sample sizes (P = 0.001). PD-1 had no prognostic significance. There was no publication bias affecting the results. Thus, among the immune checkpoint molecules, IDO and PD-L1 are potential prognostic predictors in head and neck cancer.

#### **INTRODUCTION**

Head and neck cancer (HNC) is the sixth most common malignancy worldwide [1]. Most patients exhibit advanced-stage disease, including regional lymph node involvement, and 10% of patients have distant metastases [2]. The traditional treatment options for HNC are surgery, radiotherapy and chemotherapy [3], which have severe adverse effects. Furthermore, some patients do not benefit much from these treatments, and are likely to relapse. Anatomic complexities often lead to malfunctions in speaking, swallowing and breathing after treatments, hampering patients' long-term quality of life [4]. Although there have been certain advances in treatment, the overall survival of HNC patients is still unsatisfactory, and the five-year survival rate is less than 50% [5-7].

Immunosuppressive patients are prone to suffer from HNC [8], although the predominant causes of HNC are tobacco and alcohol consumption [4] and viral infections [9, 10]. Among the functions of the immune cells, immune checkpoint activity has been reported to be involved in the surveillance of tumor development and progression [11]. Immune checkpoint molecules including programmed death 1 (PD-1) [12, 13], indoleamine 2, 3-dioxygenase (IDO) [14, 15], B7-H3 [16, 17], lymphocyte activation gene 3 (LAG-3) [18], cytotoxic T-lymphocyte-associated protein 4 (CTLA-4) [19], programmed death ligand 2 (PD-L2) [20], V- domain Ig suppressor of T cell activation (VISTA) [21], B7-H4 [22] and programmed death ligand 1 (PD-L1) [23-25] have been used as markers to evaluate the prognosis of HNC. However, the survival rates of patients with high expression of immune checkpoint molecules have differed according to the overexpressed molecule.

In the present study, we performed a systematic review of the available literature on this topic in PubMed, Embase and the Cochrane Library. Then, we conducted a meta-analysis of the survival rates (including overall survival [OS], disease-free survival [DFS], progressionfree survival [PFS], disease-specific survival [DSS] and distant metastases-free survival [DMFS]) of patients expressing different levels of immune checkpoint molecules.

#### **RESULTS**

#### **Study characteristics**

The characteristics of the included studies are shown in Table 1. There were 52 prospective studies comparing contemporary series of patients (level of evidence: 3b) in 51 articles. These studies included 7127 patients and

met the criteria for meta-analysis. The literature selection procedure is presented in Figure 1. The included articles were evaluated by the Newcastle-Ottawa Scale (NOS; Supplementary Table 1), and all the articles were published between 2010 and 2018. Roughly half of the studies were conducted in Asia (n=23), while the remainder were conducted in Europe (n=18), North and South America (n=6), Oceania (n=4) and Africa (n=1). Thus, the samples included in this meta-analysis covered most of the continents of the world. In terms of the immune checkpoint molecules, the majority of the studies evaluated PD-L1 (n=40), while the rest assessed PD-1 (n=8) and IDO (n=4). The sample sizes of the included studies ranged from 38 to 517. With reference to the mean value of all the samples, 17 studies were considered to have a large sample size (n > 139), while 35 had a small sample size  $(n \le 139)$ . Forty-three studies explored the prognostic value of their chosen immune checkpoint molecule for OS, 19 for DFS, 6 for PFS, 5 for DSS and 3 for DMFS.

#### Methodological quality of the included studies

The quality of the included studies was generally high. Most of the studies mentioned the length of the followup period, and the majority provided adequate follow-



Figure 1. Flow diagram of studies identified, included and excluded.

Author and year	Target	Country / Region	Ethnicity	Tumor location	Sampl e size	Gender M/F	Cut-off value	Detection method	TNM stage	Outcom e	HR estimatio n	Study design	NOS score
Ahn et al. 2017[23]	PD-L1	Korea	Asian	OSCC	68	45/23	Grade > 1	IHC	I-IV	OS DFS	reported	Р	7
Badoual et al. 2013[12]	PD-1	France	Caucasia n	HNSCC	64	NA	> median	IF	I-IV	OS	reported	Р	6
Balempas et al. 2017[36]	PD-L1	Germany	Caucasia n	HNSCC	161	131/30	> 5%	IHC	I-IV	OS DMFS	reported	Р	7
Ben-Haj- Ayed et al. 2016[14]	IDO	Tunisia	Caucasia n	NPC	71	48/23	> median	IHC	I-IV	OS DFS	reported	Р	7
Birtalan et al. 2017[24]	PD-L1	Hungary	Caucasia n	HNSCC	106	90/16	Score > 0%	IHC	I-IV	DSS	reported	Р	6
Budczies et al. 2016[25]	PD-L1	Germany	Caucasia n	HNSCC	517	NA	> median	qRT-PCR	NA	OS DFS	reported	Р	5
Chan et al. 2017[46]	PD-L1	USA	Caucasia n	NPC	161	117/44	≥1%	IHC	I-IV	OS PFS	reported	Р	6
Chang et al. 2017[47]	PD-L1	Philippine s	Asian	NPC	56	43/13	> 1%	IHC	I-IV	OS	reported	Р	5
Chen et al. 2015[48]	PD-L1	Taiwan	Asian	OSCC	218	145/73	> 5%	IHC	I-IV	OS	reported	Р	7
Chen et al. 2017[49]	PD-L1	China	Asian	HNSCC	496	NA	> 5%	qRT-PCR	I-IV	OS	reported	Р	7
Cho et al. 2011[50]	PD-L1	Korea	Asian	OSCC	45	32/13	Grade > 1	IHC	I-IV	OS	estimated	Р	6
De Meulenaere et al. 2017[51]	PD-L1	Belgium	Caucasia n	OSCC	99	82/17	> 1%	IHC	I-IV	OS DFS	reported	Р	6
Fang et al. 2014[52]	PD-L1	China	Asian	NPC	139	113/26	> 35%	IHC	I-IV	DFS	estimated	Р	6
Feng et al. 2017[53]	PD-L1	USA	Caucasia n	OSCC	119	74/45	< 30 µm	IHC	I-IV	OS	estimated	Р	6
Fiedler et al. 2018[54]	PD-L1	Germany	Caucasia n	HNSCC	82	73/9	> 5%	IHC	I-IV	OS	reported	Р	7
Hanna et al. 2018[37]	PD-L1	USA	Caucasia n	OSCC	81	49/32	> 10%	IHC	I-IV	OS	reported	Р	7
Hong et al. 2016[55]	PD-L1	Australia	Caucasia n	OSCC	99	79/20	> 25%	IHC	I-IV	OS	reported	Р	6
Hsu et al. 2010[13]	PD-1	Taiwan	Asian	NPC	46	39/7	> median	IHC	NA	OS DFS	reported	Р	4

### Table 1. Characteristics of included studies.

Kansy et al. 2017[56]	PD-1	Germany	Caucasia n	HNSCC	56	NA	NA	FACS	I-IV	DFS	reported	Р	6
Kim et al. 2016[57]	PD-1	Korea	Asian	HNSCC	402	302/100	> 5%	IHC	I-IV	OS DFS	reported	Р	6
Kim et al. 2016[58]	PD-1	Korea	Asian	OSCC	133	120/13	> 5%	IHC	I-IV	OS	reported	Р	7
Kogashiwa et al. 2017[35]	PD-L1	Japan	Asian	OSCC	84	57/27	> 5%	IHC	I-IV	OS PFS	reported	Р	7
Laimer et al. 2011[15]	IDO	Austria	Caucasia n	OSCC	88	67/21	> 4	IHC	I-IV	OS	reported	Р	7
Larbcharoens ub et al. 2018[59]	PD-L1	Thailand	Asian	NPC	114	77/67	≥ 5%	IHC	I-IV	OS	estimated	Р	7
Lee et al. 2016[60]	PD-L1	Hong Kong	Asian	NPC	104	85/19	> 1	IHC	I-IV	PFS DMFS OS	reported	Р	5
Li et al. 2017[61]	PD-L1	China	Asian	NPC	62	40/14	> 20%	IHC	I-IV	DFS	reported	Р	5
Lin et al. 2015[30]	PD-L1	Taiwan	Asian	OSCC	305	236/69	> 1	IHC	I-IV	OS	reported	Р	6
Muller et al. 2017[62]	PD-L1	Germany	Caucasia n	HNSCC	293	82/16 142/53 (224/69)	Score $\geq 1$	IHC	I-IV	OS	reported	Р	6
Ock et al. 2016[63]	PD-L1	South Korea	Asian	HNSCC	141	40/10 61/30 (101/40)	≥ 5%	IHC	I-IV	OS	reported	Р	6
Oguejiofor et al. 2017[64]	PD-L1	UK	Caucasia n	OPSCC	124	NA	> 5%	IHC	I-IV	OS	reported	Р	7
Oliveira- Costa et al. 2015[65]	PD-L1	Brazil	Caucasia n	OSCC	142	125/17	≥ 5%	IHC	I-III	DSS	reported	Р	6
Ono et al. 2017[66]	PD-L1	Japan	Asian	HPSCC	83	79/4	≥1%	IHC	III-IV	OS PFS	reported	Р	6
Ono et al. 2018[67]	PD-L1	Japan	Asian	NPC	66	54/12	$\geq$ 5%	IHC	I-IV	OS PFS	reported	Р	7
Ou et al. 2017[68]	PD-L1	France	Caucasia n	HNSCC	38	NA	≥1%	IHC	III-IV	OS PFS	estimated	Р	7
Qu et al. 2018[69]	PD-L1	China	Asian	NPC	96	72/24	> 10%	IHC	I-IV	DMFS	estimated	Р	6
Riobello et al. 2018[70]	PD-L1	Spain	Caucasia n	SSCC	53	37/16	≥ 5%	IHC	I-IV	OS DFS DSS	reported	Р	5
Roper et al. 2017[71]	PD-L1	Australia	Caucasia n	HNSCC	74	64/10	> 5%	IHC	NA	DFS	reported	Р	6
Satgunaseela n et al.	PD-L1	Australia	Caucasia n	OSCC	217	130/87	Score $\geq 1$	IHC	NA	DSS	estimated	Р	6

2016[72]													
Schneider et al. 2018[73]	PD-1 PD-L1	Austria	Caucasia n	HNSCC	129	97/28	> 5%	IHC	I-IV	OS DFS	reported	Р	7
Seppälä et al. 2016[74]	IDO	Finland	Caucasia n	OSCC	58	29/29	> 0	IHC	I-III	OS	reported	Р	6
Solomon et al. 2018[75]	PD-L1	Australia	Caucasia n	OSCC	190	157/33	≥ 5%	IHC	I-IV	OS	reported	Р	7
Steuer et al. 2018[76]	PD-1	USA	Caucasia n	OPSCC	97	81/16	Score > 1	IHC	I-IV	OS	reported	Р	7
Strati et al. 2017[77]	PD-L1	Greece	Caucasia n	HNSCC	113	75/19	NA	qRT-PCR	I-IV	OS PFS	reported	Р	5
Straub et al. 2016[78]	PD-L1	Germany	Caucasia n	OSCC	80	54/26	> 5%	IHC FISH	I-IV	OS DFS	estimated	Р	7
Tang et al. 2017[79]	PD-1	China	Asian	NPC	96	NA	NA	IHC	NA	OS	estimated	Р	6
Ukpo et al. 2013[80]	PD-L1	USA	Caucasia n	OPSCC	181	162/19	> 5%	IHC	I-IV	OS DFS DSS	reported	Р	7
Vassilakopou lou et al. 2016[81]	PD-L1	Greece	Caucasia n	LSCC	260	249/11	> 59th percentile of AQUA score	IHC	I-IV	OS DFS	reported	Р	7
Ye et al. 2013[82]	IDO	China	Asian	LSCC	187	179/8	NA	IHC	I-IV	OS DFS	reported	Р	6
Zhang et al. 2015[83]	PD-1 PD-L1	China	Asian	NPC	139	113/26	H-score PD-1 > 0 PD-L1 > 35	IHC	I-IV	DFS	estimated	Р	7
Zheng et al. 2017[84]	PD-L1	China	Asian	NPC	85	63/22	Score > 2	IHC	I-IV	OS	estimated	Р	6
Zhu et al. 2017[38]	PD-L1	China	Asian	NPC	209	150/59	$\geq 5\%$	IHC	I-IV	OS DFS	reported	Р	7

PD-L1: programmed death ligand 1; PD-1: programmed death 1; IDO: indoleamine 2, 3-dioxygenase; M/F: male/female; NA: not available; OSCC: oral squamous cell carcinoma; HNSCC: head and neck squamous cell carcinoma; NPC: nasopharyngeal carcinoma; OPSCC: oropharyngeal squamous cell carcinoma; HPSCC: hypopharyngeal squamous cell carcinoma; SSCC: sinonasal squamous cell carcinoma; LSCC: laryngeal squamous cell carcinoma; cut-off value: the value that can be diagnosed as positive/high expression of an immune checkpoint molecule; AQUA: automated quantitative analysis; IHC: immunohistochemistry; IF: immunofluorescence; qRT-PCR: quantitative reverse transcription polymerase chain reaction; FACS: fluorescence-activated cell sorting; FISH: fluorescence in situ hybridization; OS: overall survival; DFS: disease-free survival; DMFS: distant metastases-free survival; DSS: disease-specific survival; PFS: progression-free survival; P: prospective; NOS: Newcastle–Ottawa Quality Assessment Scale.

Ahn et al 2017	Study D	HR (95% CI)	% Weight
Ahn et al 2017	PD-L1		
Baengas tul 2017 Baengas tul 2017 Den et al 2018 Den et al 2016 Den et al 2017 Den et al 2017 Den et al 2016 Den et al 2016 Den et al 2016 Den et al 2016 Den et al 2017 Den et al 2016 Den et al	Ahn et al 2017	0.32 (0.11, 0.94)	1.80
bucches et al 2016 Dener et al 2017 Dener et al 2017 Tener et al 2017 Tener et al 2018 Dener et al 2016 Dener et al 2017 Dener et al 2016 Dener et al 2017 Dener et al 2016 Dener et a			
Dhan et al 2017       041 (0.47, 1.40)       265         Dhan et al 2017       207 (133, 322)       321         Dhan et al 2017       157 (101, 251)       318         De et al 2017       051 (02, 051)       277         De et al 2017       158 (0.47, 3.39)       158         De et al 2017       158 (0.47, 3.39)       158         Tename et al 2018       058 (0.45, 0.47)       380         Atoma et al 2018       058 (0.46, 0.43)       360         Atoma et al 2018       058 (0.46, 0.43)       360         Atoma et al 2016       058 (0.41, 0.22)       128         Don et al 2017       258 (0.41, 0.22)       278         Don et al 2017       058 (0.41, 0.22)       278         Don et al 2016       059 (0.42, 0.12)       276 <td< td=""><td></td><td></td><td></td></td<>			
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Den et al 2015 Den et al 2017 Che et al 2018 Che che et al 2018 Che che che et al 2018 Che che che che ch			
Dame at 2017       150 (101, 2.51)       3 18         Che et al 2017       0.51 (022, 0.91)       2.87         Fang et al 2017       1.50 (023, 0.91)       2.87         Fang et al 2018       0.51 (023, 0.91)       2.87         Harma et al 2018       0.56 (0.45, 0.74)       3.00         Landrage tal 2016       0.54 (0.42, 1.21)       2.97         Kopga bia 2017       0.56 (0.45, 0.74)       3.00         Landradom et al 2018       0.76 (0.42, 1.24)       2.80         Landradom et al 2016       0.54 (0.42, 1.12)       2.93         Kopga bia vet al 2017       0.56 (0.52, 1.12)       2.93         Landradom et al 2016       0.76 (0.42, 1.24)       2.80         Dan et al 2017       0.86 (0.36, 1.84)       2.25         Dan et al 2018       0.87 (0.70, 2.42)       2.76         Dan et al 2016       0.87 (0.71, 1.27)       2.83         Standrad et al 2016       0.87 (0.71, 1.27)       2.83         Standradom et al 2016       0.87 (0.71, 1.27)       2.75         Standrad al 2016			
Dhe wit al 2011       07 (0 02.2.7.0)       1.40         Fend et al 2017       1.15 (0 66, 20)       2.57         Fend et al 2016       1.15 (0 66, 2.01)       2.52         interance et al 2016       0.58 (0 24, 0.74)       3.50         Gogashiwe et al 2017       0.28 (0 16, 0.43)       3.60         Latorbanesub et al 2016       0.63 (0 22, 1.10)       1.55         Latorbanesub et al 2016       0.63 (0 22, 1.10)       1.85         Latorbanesub et al 2016       0.63 (0 32, 1.13)       2.84         Latorbanesub et al 2016       0.63 (0 32, 1.13)       2.84         Unite et al 2016       0.63 (0 33, 1.04)       2.90         Date et al 2017       2.93 (1 16, 7.24)       2.12         Date et al 2016       0.67 (0 2.3, 1.55)       2.27         Date et al 2016       0.70 (0 40, 1.21)       2.55         Date et al 2016       0.70 (0 40, 1.21)       2.55         Schneder et al 2016       0.70 (0 40, 1.21)       2.55         Schneder et al 2016       0.70 (0 40, 1.21)       2.55         Schneder et al 2016       0.70 (0 40, 1.21)       2.55         Schneder et al 2016       0.70 (0 40, 1.21)       3.50         Vasalidational (Lequared = 72.8%, p = 0.000)       0.56 (0 31, 10.50)       2.42			
De al 2017 Farg el 12017 Farg el 2017 Farg el 2017 Farg el 2016 Farg el 2017 Farg el 2016 Farg el 2017 Farg el 2016 Farg el 2017 Farg e			
Faceg et al 2017       115 (0 06, 2.0)       2.92         Fieder et al 2018       0.86 (0.45, 0.73, 3.0)       1.96         Hanna et al 2018       0.86 (0.46, 0.73, 3.0)       2.92         Action et al 2016       0.86 (0.46, 0.73, 3.0)       2.96         Cogastiva, et al 2017       0.26 (0.16, 0.43, 3.0)       2.90         Latchdracesub, et al 2018       0.76 (0.43, 1.34)       2.90         Latchdracesub, et al 2016       0.76 (0.43, 1.34)       2.90         Doc et al 2016       0.86 (0.16, 1.32)       2.11         Opact et al 2016       0.86 (0.16, 1.32)       2.12         Occ et al 2016       0.86 (0.16, 1.32)       2.12         Occ et al 2016       0.86 (0.16, 1.32)       2.12         Opac et al 2017       0.83 (0.38, 1.04)       2.25         Onc et al 2017       0.83 (0.38, 1.02)       2.12         Opac et al 2016       0.76 (0.42, 1.21)       2.99         Schender et al 2018       0.76 (0.42, 1.22)       2.99         Schender et al 2016       1.96 (0.28, 5.57)       1.04         Ukro et al 2016       1.96 (0.28, 5.57)       1.04         Ukro et al 2016       0.76 (0.41, 1.22)       3.99         Schender et al 2016       0.76 (0.41, 0.28, 5.26)       1.75			
Fieder et al 2018 tama et al 2019 (Sogastiwa et al 2017 (Sogastiwa et al 2017 (Sogastiwa et al 2017 (Sogastiwa et al 2017 (Des (La			
Hanne et al 2016       0.58 (0.46, 0.74)       3.60         Hong et al 2017       0.56 (0.40, 0.28, 1.12)       2.51         Cognativa et al 2016       0.76 (0.43, 1.34)       2.90         Latocharoensub et al 2016       0.76 (0.43, 1.34)       2.90         Latocharoensub et al 2016       0.76 (0.43, 1.34)       2.90         Latocharoensub et al 2016       0.76 (0.43, 1.34)       2.90         Dock et al 2016       0.83 (0.86, 0.94, 0.92)       1.81         Opt et al 2017       0.83 (0.36, 0.94, 0.92)       2.81         Once et al 2016       0.87 (0.27, 1.95)       2.27         Dot et al 2017       0.27 (0.24, 1.22)       0.99         Schender et al 2016       1.35 (0.74, 4.94)       2.81         Schender et al 2016       1.35 (0.74, 4.94)       2.81         Schender et al 2016       1.50 (0.86, 0.28)       1.89         Straub et al 2016       1.45 (0.28, 1.57)       1.04         Ukpo et al 2017       0.66 (0.34, 0.96)       3.02         Vasalakopoulus et al 2016       0.87 (0.71, 1.05)       2.56         Schender et al 2016       0.56 (0.74, 1.22)       3.59         Vasalakopoulus et al 2016       0.56 (0.34, 0.96)       3.02         Schender et al 2016       0.56 (0.34, 0.96)       3.0			
Hong at 2016 Kogashwa et al 2017 Lice at al 2018 Lice at al 2018 Lice at al 2017 Lice at al 2017 Lice at al 2017 Lice at al 2017 Dock et al 2018 Dock			
Cogenative et al 2017       0.26 (0.16, 0.43)       3.06         Lathochrancensub et al 2018       0.76 (0.43, 1.34)       2.90         Lathochrancensub et al 2016       0.68 (0.22, 1.60)       1.85         Lin et al 2017       2.99 (1.16, 7.24)       2.12         Dock et al 2016       0.46 (0.16, 1.32)       1.44         Opact et al 2017       0.46 (0.16, 1.32)       1.44         Opact et al 2017       0.46 (0.16, 1.32)       1.44         Onc et al 2017       0.67 (0.28, 1.55)       2.27         Onc et al 2018       0.67 (0.28, 1.55)       2.27         Schmodt et al 2018       0.770 (0.40, 1.21)       2.95         Schmod et al 2016       0.770 (0.40, 1.21)       2.95         Stratu et al 2016       0.770 (0.40, 1.21)       2.95         Stratu et al 2016       0.770 (0.40, 1.21)       2.95         Value et al 2016       0.83 (0.33, 1.02)       3.12         Value et al 2016       0.56 (0.34, 0.55)       3.02         Schmod et al 2016       0.70 (0.42, 2.05)       1.77         Schue et			
Labcharossub et al 2018 Lee et al 2016 In et al 2017 Ock et al 2018 Ock et			
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Multer et al 2017 Cock et al 2016 Que et al 2017 Cock et al 2018 Schneider et al 2016 Cock et al 2017 Cock et al 2017 Cock et al 2018 Schneider et al 2018 S			
Dek et al 2018       0.46 (0.16, 1.32)       1.84         Oper et al 2017       0.83 (0.36, 1.94)       2.25         Once et al 2018       0.67 (0.28, 1.55)       2.27         Out et al 2017       0.22 (0.04, 1.22)       0.99         Schneider et al 2018       0.70 (0.40, 1.21)       2.95         Schneider et al 2018       0.70 (0.40, 1.21)       2.95         Stonm et al 2016       1.90 (0.68, 5.28)       1.75         Straub et al 2016       1.90 (0.68, 5.23, 50)       1.75         Straub et al 2016       0.83 (0.39, 1.02)       3.12         Straub et al 2017       0.96 (0.34, 0.55)       3.02         Vassilakopoulou et al 2016       0.83 (0.39, 1.02)       3.12         Straub et al 2017       0.56 (0.34, 0.55)       3.02         Vassilakopoulou et al 2016       0.87 (0.71, 1.07)       7.53         Stubtotall (I-squared = 72.8%, p = 0.000)       0.87 (0.71, 1.07)       7.53         PD-1       136 (0.02, 0.75)       0.95         Stubtotall (I-squared = 76.7%, p = 0.000)       0.93 (0.22, 0.25)       1.79         Stubtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.71, 1.07)       1.73         Stubtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.71, 1.07)       1.53         OD       0.93 (0.79			
Ogu et al 2017       0.83 (0.36, 1.94)       2.25         One et al 2017       1.30 (0.70, 2.42)       2.78         Dor et al 2018       0.67 (0.29, 1.55)       2.27         Dor et al 2018       0.70 (0.40, 1.21)       2.95         Schneider et al 2018       0.70 (0.40, 1.21)       2.95         Schneider et al 2018       0.70 (0.40, 1.21)       2.95         Schneider et al 2016       1.90 (0.86, 5.28)       1.89         Ukpo et al 2016       1.90 (0.86, 5.28)       1.89         Vassilakopoulou et al 2016       0.95 (0.74, 1.22)       3.59         Vassilakopoulou et al 2016       0.63 (0.39, 1.02)       3.12         Zhen et al 2017       0.56 (0.40, 0.95)       3.02         Subtotal (I-squared = 72.8%, p = 0.000)       0.57 (0.31, 1.07)       77.53         PD-1       0.13 (0.02, 0.75)       0.95         Badoual et al 2016       0.70 (0.42, 2.05)       1.79         Schneider et al 2018       0.50 (0.34, 0.95)       3.02         Subtotal (I-squared = 72.8%, p = 0.000)       0.51 (0.15, 56.26)       0.91         Schneider et al 2016       0.70 (0.42, 2.05)       1.79         Schneider et al 2016       0.70 (0.42, 2.05)       1.79         Subtotal (I-squared = 78.7%, p = 0.000)       0.93 (0.42, 2.02			
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Dno et al 2018       0.67 (0.29, 1.55)       2.27         Out et al 2017       0.22 (0.04, 1.22)       0.99         Schneider et al 2018       0.70 (0.40, 1.21)       2.96         Schneider et al 2018       0.70 (0.40, 1.21)       2.95         Schneider et al 2018       0.70 (0.40, 1.21)       2.95         Schneider et al 2018       0.70 (0.40, 1.21)       2.95         Strait et al 2017       1.90 (0.68, 5.28)       1.89         Ukpo et al 2016       0.95 (0.74, 1.22)       3.59         Zhang et al 2017       0.56 (0.74, 1.22)       3.59         Schneider et al 2016       0.56 (0.31, 0.02)       3.12         Zheng et al 2017       0.56 (0.34, 0.95)       3.02         Subtotal (I-squared = 72.8%, p = 0.000)       0.87 (0.71, 1.07)       77.53         PD-1       360-1120, 562)       2.42         Schneider et al 2013       0.57 (0.31, 1.05)       2.66         Hsu et al 2016       0.57 (0.31, 1.05)       2.60         Schneider et al 2017       0.32 (0.11, 0.95)       1.77         Schneider et al 2018       0.32 (0.11, 0.95)       1.77         Schneider et al 2016       0.33 (0.42, 2.02)       1.29         DO       0.57 (0.31, 1.05)       2.58       2.20 (1.20, 4.22)			
Du et al 2017 Robelio et al 2018 Solomon et al 2018 Solomon et al 2018 Solomon et al 2018 Solomon et al 2018 Strati et al 2017 Straub et al 2016 Uspo et al 2016 Uspo et al 2017 Zheng et al 2017 De 1 Badoual et al 2017 De 1 Badoual et al 2017 Subtotal (I-squared = 72.8%, p = 0.000) De 57 (0.31, 1.05) Solomon et al 2016 Liss (0.24, 2.05) Solomon et al 2016 Solomon et al 2017 De 1 Badoual et al 2017 Solomon et al 2018 Subtotal (I-squared = 76.7%, p = 0.000) De 30 (0.42, 2.02) Solomon et al 2016 Subtotal (I-squared = 59.8%, p = 0.059) Correl (I-squared = 74.8%, p = 0.000) De 30 (0.42, 2.02) De			
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Schneider et al 2018       0.70 (0.40, 1.21)       2.95         Solmon et al 2018       1.90 (0.68, 5.28)       1.89         Strait et al 2016       7.96 (2.65, 23.90)       1.75         Straub et al 2016       0.95 (0.74, 1.22)       3.59         Vassilakopoulou et al 2016       0.63 (0.39, 1.02)       3.12         Zheng et al 2017       0.56 (0.34, 0.95)       3.02         Stubt et al 2017       0.56 (0.34, 0.95)       3.02         Subtotal (I-squared = 72.8%, p = 0.000)       0.87 (0.71, 1.07)       77.53         PD-1       0.56 (0.34, 0.95)       3.02         Schneider et al 2013       0.55 (0.31, 1.05)       2.60         You et al 2016       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       0.70 (0.24, 2.05)       1.79         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       122         DO       1.07 (0.49, 2.35)       2.38       1.70 (0.49, 2.35)       2.38         Limmer et al 2016       1.07 (0.49, 2.35)       2.38       3.97 (1.55, 10.15)       2.05         DO       1.07 (0.49, 2.17)       1.94       3.97 (1.55, 10.15)       2.05         Subtotal (I-squared = 59.8%, p = 0.059)	Ou et al 2017	0.22 (0.04, 1.22)	0.99
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Strati et al 2017       7.96 (2 65, 23.90)       1.75         Stratu et al 2016       1.45 (0 28, 7.57)       1.04         Ukpo et al 2016       0.95 (0 74, 1.22)       3.59         Vessilakopoulou et al 2016       0.63 (0 39, 1.02)       3.12         Zheng et al 2017       5.61 (0 18, 172.93)       0.31         Des de 12 2017       0.66 (0 34, 0.95)       3.02         Subtotal (I-squared = 72.8%, p = 0.000)       0.87 (0 71, 1.07)       77.53         PD-1       0.13 (0 02, 0.75)       0.95         Badoual et al 2016       0.13 (0 02, 0.75)       0.95         Kim et al 2016       0.57 (0 31, 1.05)       2.80         Kim et al 2016       0.70 (0 24, 2.05)       1.79         Schneider et al 2018       2.60 (1 20, 5.62)       2.42         Statuer et al 2017       0.39 (0 42, 2.02)       12.95         DO       0.93 (0 42, 2.02)       12.95       1.07 (0 49, 2.35)       2.38         Laimer et al 2016       1.07 (0 49, 2.35)       2.38       3.97 (1.55, 10.15)       2.06         DO       0.93 (0 42, 2.02)       12.95       3.97 (1.55, 10.15)       2.05       2.02 (1 20, 4.02)       9.52         Do       0.98 (0,79, 1.17)       1.94       3.97 (1.55, 10.15)       2.05       2.06 (1.	Schneider et al 2018	0.70 (0.40, 1.21)	2.95
Straub et al 2016       1.45 (0.28, 7.57)       1.04         Ukpo et al 2013       0.95 (0.74, 1.22)       3.59         Vassilakopoulou et al 2016       0.63 (0.39, 1.02)       3.12         Zheng et al 2017       0.56 (0.34, 0.95)       3.02         Stubtotal (I-squared = 72.8%, p = 0.000)       0.87 (0.71, 1.07)       77.53         PD-1       0.56 (0.34, 0.95)       3.02         Badoual et al 2016       0.57 (0.31, 1.05)       2.80         Kim et al 2016       0.57 (0.31, 1.05)       2.80         Kim et al 2016       0.57 (0.31, 1.05)       2.80         Subtotal (I-squared = 76, 7%, p = 0.000)       0.97 (0.24, 2.05)       1.79         Subtotal (I-squared = 76, 7%, p = 0.000)       0.93 (0.42, 2.02)       1.25         DO       0       0.93 (0.42, 2.02)       1.25         DO       0       0.93 (0.42, 2.02)       1.25         Subtotal (I-squared = 59.8%, p = 0.059)       0.96 (0.79, 1.17)       1.00 0.00	Solomon et al 2018	1.90 (0.68, 5.28)	1.89
Ukpo et al 2013       0.95 (0.74, 1.22)       3.59         Vassilakopoulou et al 2016       0.63 (0.39, 1.02)       3.12         Zheng et al 2017       5.61 (0.18, 172.93)       0.31         Zhu et al 2017       0.56 (0.34, 0.95)       3.02         Subtotal (I-squared = 72.8%, p = 0.000)       0.87 (0.11, 1.07)       77.53         PD-1       0.13 (0.02, 0.75)       0.95         Badoalet al 2013       0.13 (0.02, 0.75)       0.95         Hsu et al 2016       0.13 (0.02, 0.75)       0.95         Kim et al 2016       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       2.60 (1.02, 5.62)       2.42         Stubtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       1.295         DO       0.51       0.93 (0.42, 2.02)       1.295         DO       0.51       1.07 (0.49, 2.35)       2.38         Laimer et al 2016       2.20 (1.20, 4.02)       5.25         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)	Strati et al 2017	7.96 (2.65, 23.90)	1.75
vassilakopoulou et al 2016       0.63 (0.39, 1.02)       3.12         Zheng et al 2017       5.61 (0.18, 172, 93)       0.31         Subtotal (I-squared = 72.8%, p = 0.000)       0.56 (0.34, 0.95)       3.02         PD-1       0.13 (0.02, 0.75)       0.95         Badoual et al 2013       9.51 (1.55, 58.26)       0.91         Schneider et al 2016       0.13 (0.02, 0.75)       0.95         Kim et al 2016       0.13 (0.02, 0.75)       0.95         Schneider et al 2018       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       0.32 (0.11, 0.95)       1.77         Tang et al 2017       1.50 (0.86, 3.41)       2.30         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         DO       0.93 (0.42, 2.02)       12.95       2.35         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         DO       0.93 (0.42, 2.02)       12.95       2.38         Laimer et al 2016       1.07 (0.49, 2.35)       2.38         Laimer et al 2016       1.70 (1.06, 2.71)       3.15         Sepolat et al 2016       2.02 (1.20, 4.02)       9.52         Oceanal (I-squared = 59.8%, p = 0.059)       2.02 (1.20, 4.02)       9.52         Operall (I-squared	Straub et al 2016	1.45 (0.28, 7.57)	1.04
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Zheng et al 2017       5.61 (0.18, 172.93)       0.31         Zhu et al 2017       0.56 (0.34, 0.95)       3.02         Subtotal (I-squared = 72.8%, p = 0.000)       0.87 (0.71, 1.07)       77.53         PD-1       0.13 (0.02, 0.75)       0.95         Badoual et al 2016       0.13 (0.02, 0.75)       0.95         Kim et al 2016       0.57 (0.31, 1.05)       2.80         Kim et al 2016       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       0.32 (0.11, 0.95)       1.77         Tang et al 2017       1.50 (0.66, 3.41)       2.30         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95           1.50 (0.66, 3.41)       2.30         Subtotal (I-squared = 76.7%, p = 0.000)        0.93 (0.42, 2.02)       12.95                                   Scheider et al 2018	Vassilakopoulou et al 2016	0.63 (0.39, 1.02)	3.12
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Subtotal (I-squared = 72.8%, p = 0.000)       0.87 (0.71, 1.07)       77.53         PD-1       38doual et al 2013       0.13 (0.02, 0.75)       0.95         Hsu et al 2010       9.51 (1.55, 58.26)       0.91         Kim et al 2016       0.57 (0.31, 1.05)       2.80         Scheider et al 2018       0.70 (0.24, 2.05)       1.79         Steuer et al 2018       0.32 (0.11, 0.55)       1.77         Tang et al 2017       0.32 (0.11, 0.55)       1.77         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         DO       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.07 (0.49, 2.35)       2.38         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52         Oxerall (I-squared = 74.8%, p = 0.000)       0.96 (0.79, 1.17)       100.00	Zhu et al 2017		3.02
Badoual et al 2013       0.13 (0.02, 0.75)       0.95         Hsu et al 2010       9.51 (1.55, 58.26)       0.91         Kim et al 2016       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       2.60 (1.20, 5.62)       2.42         Steure et al 2017       0.32 (0.11, 0.95)       1.77         Tang et al 2017       0.90 (0.66, 3.41)       2.30         Subtrat (I-squared = 76.7%, p = 0.000)       0.91 (1.65, 12.19)       1.94         Pon-Haj-Ayed et al 2016       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.07 (0.49, 2.35)       2.38         Sepatal et al 2016       1.07 (0.49, 2.35)       2.38         Laimer et al 2013       3.97 (1.55, 10.15)       2.05         Ver et al 2016       2.20 (1.20, 4.02)       9.52         Ver et al 2016       2.20 (1.20, 4.02)       9.52         Ver et al 2013       3.97 (1.55, 10.15)       2.05         Ver et al 2013       2.20 (1.20, 4.02)       9.52         Ver et al 2013       9.96 (0.79, 1.17)       100.00	Subtotal (I-squared = 72.8%, p = 0.000)		77.53
Hsu et al 2010 Kim et al 2016 Scheeider et al 2018 Scheeider et al 2018 Scheeider et al 2018 Subtotal (I-squared = 76.7%, p = 0.000) DO Ben-Haj-Ayed et al 2016 Laimer et al 2011 Sepefaile et al 2016 Laimer et al 2016 Correct al 2016 Laimer et al 2016 Laimer et al 2016 Correct al 2016 Laimer et al 2016 Laimer et al 2016 Correct al 2016 Co	PD-1		
Kim et al 2016       0.57 (0.31, 1.05)       2.80         Kim et al 2016       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       2.60 (1.20, 5.62)       2.42         Steuer et al 2017       0.32 (0.11, 0.95)       1.77         Tang et al 2017       1.50 (0.66, 3.41)       2.30         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         IDO       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.70 (1.06, 2.71)       3.15         Sepedia et al 2016       4.49 (1.65, 12.19)       1.94         Ye et al 2013       3.97 (1.55, 10.15)       2.05         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52         Overall (I-squared = 74.8%, p = 0.000)       0.96 (0.79, 1.17)       100.00	Badoual et al 2013	0.13 (0.02, 0.75)	0.95
Kim et al 2016       0.57 (0.31, 1.05)       2.80         Kim et al 2016       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       2.60 (1.20, 5.62)       2.42         Subotal (I-squared = 76.7%, p = 0.000)       0.32 (0.11, 0.95)       1.77         Subotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         Joint       1.50 (0.66, 3.41)       2.30         Subotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         Joint       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.70 (1.06, 2.71)       3.15         Sepublia et al 2016       4.49 (1.65, 12.19)       1.94         Ye et al 2013       3.97 (1.55, 10.15)       2.05         Subotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       5.52         Joint       0.96 (0.79, 1.17)       100.00	Hsu et al 2010	9.51 (1.55, 58.26)	0.91
Kim et al 2016       0.70 (0.24, 2.05)       1.79         Schneider et al 2018       2.60 (1.20, 5.62)       2.42         Steuer et al 2017       0.32 (0.11, 0.95)       1.77         Tang et al 2017       1.50 (0.66, 3.41)       2.30         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95               DO            Sep-Haj-Ayed et al 2016       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.70 (1.06, 2.71)       3.15         Sepolat et al 2016       1.70 (1.06, 2.71)       3.15         Subtotal (I-squared = 59.8%, p = 0.059)       2.90 (1.55, 10.15)       2.05	Kim et al 2016		2.80
Schneider et al 2018       2.60 (1.20, 5.62)       2.42         Steuer et al 2018       0.32 (0.11, 0.95)       1.77         Tarag et al 2017       1.50 (0.66, 3.41)       2.30         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         DO	Kim et al 2016		1.79
Steuer et al 2018       0.32 (0.11, 0.95)       1.77         Tang et al 2017       1.50 (0.66, 3.41)       2.30         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         DO	Schneider et al 2018	2.60 (1.20, 5.62)	2.42
Tang et al 2017       1.50 (0.66, 3.41)       2.30         Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         DO       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.70 (1.06, 2.71)       3.15         Sepeñal et al 2016       4.49 (1.65, 12.19)       1.94         Ye et al 2013       3.97 (1.55, 10.15)       2.05         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52           0.96 (0.79, 1.17)       100.00	Steuer et al 2018		1.77
Subtotal (I-squared = 76.7%, p = 0.000)       0.93 (0.42, 2.02)       12.95         IDO       1.07 (0.49, 2.35)       2.38         Laimer et al 2016       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.70 (1.06, 2.71)       3.15         Septala et al 2016       4.49 (1.65, 12.19)       1.94         Ye et al 2013       3.97 (1.55, 10.15)       2.05         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52         Overall (I-squared = 74.8%, p = 0.000)       0.96 (0.79, 1.17)       100.00	_ 1_		
Ben-Haj-Ayed et al 2016       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.70 (1.06, 2.71)       3.15         Seppäla et al 2016       4.49 (1.65, 12.19)       1.94         Ye et al 2013       3.97 (1.55, 10.15)       2.05         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52           0.96 (0.79, 1.17)       100.00			12.95
Ben-Haj-Ayed et al 2016       1.07 (0.49, 2.35)       2.38         Laimer et al 2011       1.70 (1.06, 2.71)       3.15         Seppäla et al 2016       4.49 (1.65, 12.19)       1.94         Ye et al 2013       3.97 (1.55, 10.15)       2.05         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52           0.96 (0.79, 1.17)       100.00	DO		
Laimer et al 2011 Seppala et al 2016 Ye et al 2013 Subtotal (I-squared = 59.8%, p = 0.059) Overall (I-squared = 74.8%, p = 0.000) Overall (I-squared = 74.8%, p = 0.000)	Ben-Haj-Ayed et al 2016	1.07 (0.49, 2.35)	2.38
Seppälä et al 2016       4.49 (1.65, 12.19)       1.94         Ye et al 2013       3.97 (1.55, 10.15)       2.05         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52         Overall (I-squared = 74.8%, p = 0.000)       0.96 (0.79, 1.17)       100.00			
Ye et al 2013       3.97 (1.55, 10.15)       2.05         Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52         Overall (I-squared = 74.8%, p = 0.000)       0.96 (0.79, 1.17)       100.00			
Subtotal (I-squared = 59.8%, p = 0.059)       2.20 (1.20, 4.02)       9.52         Overall (I-squared = 74.8%, p = 0.000)       0.96 (0.79, 1.17)       100.00	Ye et al 2013		
	Subtotal (I-squared = 59.8%, p = 0.059)	-	
VOTE: Weights are from random effects analysis	Overall (I-squared = 74.8%, p = 0.000)	0.96 (0.79, 1.17)	100.00
	NOTE: Weights are from random effects analysis		

# Figure 2: Overall forest plot of stratified analysis based on the type of molecule for the association of immune checkpoint molecules with OS.

up data for more than five years. Nevertheless, almost none of the prospective studies had an exposed cohort that sufficiently represented the general characteristics of the population in the community, as this factor was not considered in the study design. None of the studies were designed with adequate comparability of cohorts, due to their failure to match exposed and non-exposed individuals and/or adjust for confounders. Methods for handling missing data and intention-to-treat analysis were not adequately described in the majority of the studies.

# Immune checkpoint molecule expression and prognosis of HNC patients

Forty-three studies with 6225 patients reported the relationship between OS and at least one of the three immune checkpoint molecules in HNC. The expression

	Variable	Study no	Sample size	HR (95% CI)	P value -	Hetero	geneity
	variable	Study no.	Sample size	HK (95% CI)	P value -	$\mathbf{I}^2$	P value
OS	Overall	43	6225	0.964 (0.791-1.175)	0.714	74.8%	< 0.001
	Immune che	ckpoint molect	iles	· · · · · ·			
	PD-L1	32	4854	0.874 (0.711-1.073)	0.197	72.8%	< 0.001
	PD-1	7	967	0.926 (0.424-2.025)	0.848	76.7%	< 0.001
	IDO	4	404	2.197 (1.199-4.023)	0.011	59.8%	0.059
	Ethnicity	•	101	2.137 (11133 11023)	0.011	591070	0.029
	Asian	19	2938	0.923 (0.651-1.307)	0.650	77.1%	< 0.001
	Caucasian	24	3287	0.995 (0.779-1.270)	0.965	73.8%	< 0.001
	Tumor locati		5207	0.555 (0.775 1.270)	0.905	/5.0/0	-0.001
	OSCC	13	1477	0.879 (0.586-1.317)	0.532	85.0%	< 0.001
	NPC	10	1008	0.862 (0.618-1.203)	0.383	33.7%	0.139
	OPSCC	4	592	0.878 (0.532-1.450)	0.611	47.1%	0.129
	HPSCC	1	83	1.300 (0.700-2.415)	0.407		-
	SSCC	1	53	1.355 (0.739-2.485)	0.326	_	_
	LSCC	2	447	1.517 (0.252-9.126)	0.649	91.4%	0.001
	Sample size	2		1.517(0.252-9.120)	0.079	J1.7/0	0.001
	Large	14	3721	1.044 (0.803-1.356)	0.748	74.0%	< 0.001
	Small	29	2504	0.915 (0.687-1.220)	0.748	74.0%	< 0.001
DEC							
DFS	Overall	19	2901	1.097 (0.733-1.642)	0.652	92.5%	< 0.001
	Inhibitory in	nmune checkp	oint molecules				
	PD-L1	13	2010	0.874 (0.523-1.459)	0.606	94.1%	< 0.001
	IDO	2	258	1.725 (0.611-4.869)	0.303	59.5%	0.116
	PD-1	4	633	1.931 (0.716-5.211)	0.194	87.5%	< 0.001
	Ethnicity			,			
	Asian	8	1252	1.131 (0.506-2.533)	0.764	93.6%	< 0.001
	Caucasian	11	1649	1.060 (0.760-1.479)	0.731	73.9%	< 0.001
	Tumor locati		1019	1.000 (0.700 1.177)	0.7.51	101970	0.001
	OSCC	3	247	0.609 (0.208-1.788)	0.367	70.8%	0.033
	NPC	6	666	1.339 (0.581-3.085)	0.494	92.5%	< 0.001
	SSCC		53	1.834 (0.955-3.522)	0.494	92.370	
		1				-	-
	OPSCC	1	181	1.090 (0.783-1.518)	0.610	-	-
	LSCC	2	447	1.282 (0.242-6.783)	0.770	85.9%	0.008
	Sample size						
	Large	6	1756	0.844 (0.595-1.198)	0.343	75.5%	< 0.001
	Small	13	1145	1.225 (0.764-1.963)	0.399	88.9%	< 0.001
PFS	Overall	6	545	0.996 (0.585-1.685)	0.989	68.5%	0.007
	Inhibitory in	nmune checkp	oint molecules	· · · · · ·			
	PD-L1	6	545	0.891 (0.565-1.404)	0.989	68.5%	0.007
	Ethnicity	U U	010		0.202	00.070	0.007
	Asian	3	233	0.846 (0.492-1.455)	0.744	48.3%	0.144
	Caucasian	3	312	1.218 (0.372-3.993)	0.546	82.7%	0.003
	Tumor locati		512	1.210(0.372-3.773)	0.040	02.770	0.005
	NPC	2	227	0.762 (0.506-1.149)	0.195	0.0%	0.935
	OSCC	1	84	0.576 (0.308-1.076)	0.084	-	-
	HPSCC	1	83	1.350 (0.740-2.463)	0.328	-	-
	Sample size	1	05	1.330 (0.740-2.403)	0.520	-	-
	Large	1	161	0.770 (0.480-1.235)	0.279		
	-	1	384			-	0.005
	Small	5		1.067 (0.536-2.125)	0.853	73.0%	
DSS	Overall	5	699	0.779 (0.330-1.839)	0.569	84.7%	< 0.001
-	Overall	3	361	0.599 (0.346-1.035)	0.066	0.0%	0.604

# Table 2. Results of the meta-analysis on the prognostic effects of immune checkpoint molecules in HNC patients.

of these molecules was detected mainly at the protein level, except for three studies that evaluated *PD-L1* mRNA levels. Overexpression was defined based on cut-off criteria that differed among the studies (as presented in Table 1). When the data for all three immune checkpoint molecules were pooled, there was no significant relationship between the overexpression of these molecules and OS (hazard ratio [HR] = 0.964; 95% confidence interval [CI]: 0.791-1.175, P = 0.714; Table 2), and there was obvious overall heterogeneity ( $I^2 = 74.8\%$ ,  $P_h < 0.001$ ; Figure 2). Similar results were obtained for DFS, PFS, DSS and DMFS.

#### Subgroup analyses

Subgroup analyses stratified according to the immune checkpoint molecule, patient ethnicity, tumor location

and sample size were performed to detect potential sources of heterogeneity. In the stratification based on the immune checkpoint molecule (Figure 2), poorer OS was consistently found in patients with higher levels of IDO (Table 2), correlating with a poorer prognosis (HR = 2.197, 95% CI: 1.199-4.023, P = 0.011, Figure 2). However, no obvious trend in DFS was found according to IDO expression (Table 2).

The same hierarchical strategy was used to evaluate the studies of PD-L1 (Table 3). Among the immune checkpoint molecules, PD-L1 was the focus of the largest percentage of studies, as 32 studies with 4854 patients reported the relationship between PD-L1 expression and OS (Figure 3). There was a possible trend for a better prognosis in patients overexpressing PD-L1 (HR = 0.874; 95% CI: 0.711-1.073, P = 0.197).

ID	HR (95% CI)	Weight
oscc		
Ahn et al 2017	0.32 (0.11, 0.94)	2.82
Chen et al 2015	2.07 (1.33, 3.22)	5.95
Cho et al 2011	0.78 (0.22, 2.76)	2.26
De et al 2017	0.51 (0.29, 0.91)	5.11
Feng et al 2017	1.15 (0.66, 2.01)	5.24
Hanna et al 2018	0.58 (0.45, 0.74)	7.05
Hong et al 2016	0.54 (0.26, 1.12)	4.27
Kogashiwa et al 2017	0.26 (0.16, 0.43)	5.58
Lin et al 2015	1.21 (0.89, 1.64)	6.75
Straub et al 2016	<b>1.45</b> (0.28, 7.57)	1.50 46.55
Subtotal (I-squared = 84.7%, p = 0.000)	0.73 (0.47, 1.12)	46.55
NPC ! Chan et al 2017	0.81 (0.47, 1.40)	5.30
Chang et al 2017	0.49 (0.07, 3.49)	5.30 1.13
Larbcharoensub et al 2018	0.76 (0.43, 1.34)	5.18
Lee et al 2016	0.63 (0.22, 1.80)	2.90
Ono et al 2018	0.67 (0.29, 1.55)	3.75
Zheng et al 2017	◆ 5.61 (0.18, 172.93)	0.41
Zhu et al 2017	0.56 (0.34, 0.95)	5.48
Subtotal (I-squared = 0.0%, p = 0.855)	0.69 (0.52, 0.92)	24.16
. I OPSCC I		
Ogu et al 2017	0.83 (0.36, 1.94)	3,70
Solomon et al 2018	- 1.90 (0.68, 5.28)	2.99
Ukpo et al 2013	0.95 (0.74, 1.22)	7.04
Subtotal (I-squared = $0.0\%$ , p = $0.403$ )	0.98 (0.77, 1.23)	13.73
HPSCC I Ono et al 2017	1.30 (0.70, 2.42)	4.88
Subtotal (I-squared = .%, p = .)	1.30 (0.70, 2.42)	4.88
SSCC Riobello et al 2018	1.35 (0.74, 2.49)	4.96
Subtotal (I-squared = .%, p = .)	1.35 (0.74, 2.49)	4.96 4.96
. !	1.33 (0.74, 2.49)	4.50
LSCC Vassilakopoulou et al 2016	0.63 (0.39, 1.02)	5.73
Subtotal (I-squared = .%, p = .)	0.63 (0.39, 1.02) 0.63 (0.39, 1.02)	5.73 5.73
	0.03 (0.39, 1.02)	5.75
Overall (I-squared = 70.0%, p = 0.000)	0.80 (0.64, 1.01)	100.00
NOTE: Weights are from random effects analysis		
.00578 1	173	

Figure 3. Overall forest plot of stratified analysis based on the tumor location for the association between PD-L1 and OS.

	Variable	Study	Sample size	HR (95% CI)	D voluo	Hetero	geneity
	variable	no.	Sample size	пк (93% UI)	r value	I <sup>2</sup>	P value
OS	Overall	32	4854	0.874 (0.711-1.073)	0.197	72.8%	< 0.001
	Ethnicity				I         I         P value $1.073$ ) $0.197$ $72.8\%$ <0.00		
	Asian	14	2074	0.792 (0.537-1.168)		< 0.001	
	Caucasian	18	2780	0.91 (0.716-1.158)		< 0.001	
	Tumor locat	ion					
	OSCC	10	1198	0.726 (0.470-1.121)	0.148	84.7%	< 0.001
	NPC	7	795	0.692 (0.523-0.915)	0.01	0.0%	0.855
	OPSCC	3	495	0.975 (0.771-1.234)	0.835	0.0%	0.403
	HPSCC	1	83	1.300 (0.700-2.415)	0.407	-	-
	SSCC	1	53	1.355 (0.739-2.485)	0.326	-	-
	LSCC	1	260	0.635 (0.393-1.025)	0.063	-	-
	Sample size						
	Large	12	3132	1.022 (0.790-1.321)	0.87	71.4%	< 0.001
	Small	20	1722	0.77 (0.575-1.031)	0.08	66.6%	< 0.001
DFS	Overall	13	2011	0.874 (0.523-1.465)	0.607	93.9%	< 0.001
	Ethnicity						
	Asian	5	617	0.824 (0.290-2.338)	0.716	94.2%	< 0.001
	Caucasian	8	1394	0.883 (0.638-1.221)	0.451	62.4%	0.009
	Tumor locat	ion					0.009
	OSCC	3	247	0.610 (0.208-1.793)	0.369	70.5%	0.034
	NPC	4	549	1.042 (0.349-3.111)	0.941	94.9%	< 0.001
	SSCC	1	53	1.834 (0.955-3.522)	0.068	-	-
	OPSCC	1	181	1.090 (0.783-1.518)	0.610	-	-
	LSCC	1	260	0.591 (0.350-0.997)	0.048	-	-
	Sample size					72.8%       <0.00	
	Large	4	1167	0.829 (0.597-1.151)	0.263	57.5%	0.07
	Small	9	844	0.900 (0.454-1.785)	0.762	91.7%	< 0.001
PFS	Overall	7	630	0.996 (0.632-1.569)	0.986	62.1%	0.015
	Ethnicity						
	Asian	4	318	0.879 (0.585-1.321)	0.534	24.2%	0.266
	Caucasian	3	312	1.219 (0.372-3.997)	0.744	82.6%	0.003
	Tumor locat	ion					
	OSCC	2	169	0.706 (0.416-1.197)	0.196	7.8%	0.298
	HPSCC	1	83	1.350 (0.737-2.473)	0.331	-	-
	NPC	2	227	0.762 (0.503-1.154)	0.200	0.0%	0.935
	Sample size			. ,			
	Large	1	161	0.770 (0.476-1.246)	0.287	-	< 0.001
	Small	6	469	1.058 (0.600-1.876)	0.845	66.2%	0.011
DSS	Overall	5	699	0.779 (0.330-1.839)	0.569	84.7%	< 0.001
DMFS	Overall	3	361	0.599 (0.346-1.035)			0.604

# Table 3. Results of the meta-analysis on the prognostic effects of PD-L1 in HNC patients.

	Variable	S4 d a	Comula dina	UD (050/ CI)	Develope	Hetero	geneity
	Variable	Study no.	Sample size	HR (95% CI)	P value -	I <sup>2</sup>	P value
OS	Overall	17	2581	0.754 (0.568-1.002)	0.051	76.6%	< 0.001
	Ethnicity				P value $l^2$ P value           (0.568-1.002)         0.051         76.6%         <0.		
	Asian	7	1255	0.720 (0.385-1.348)		< 0.001	
	Caucasian	10	1326	0.742 (0.578-0.954)	0.020	46.1%	0.054
	Tumor locat	tion		$1^2$ $p_{V3}$ $0.754 (0.568-1.002)$ $0.051$ $76.6\%$ $<0.$ $0.720 (0.385-1.348)$ $0.305$ $88.1\%$ $<0.$ $0.742 (0.578-0.954)$ $0.020$ $46.1\%$ $0.020$ $0.653 (0.292-1.462)$ $0.300$ $90.7\%$ $<0.$ $0.649 (0.458-0.920)$ $0.015$ $0.0\%$ $0.7$ $0.975 (0.771-1.234)$ $0.835$ $0.0\%$ $0.4$ $0.635 (0.393-1.025)$ $0.063$ $  0.984 (0.659-1.468)$ $0.936$ $79.5\%$ $<0.$ $0.582 (0.426-0.796)$ $0.001$ $50.3\%$ $0.6$ $0.928 (0.618-1.392)$ $0.717$ $69.4\%$ $0.6$ $0.809 (0.241-2.720)$ $0.732$ $85.8\%$ $0.6$ $0.699 (0.114-4.263)$ $0.697$ $78.4\%$ $0.6$ $1.215 (0.288-5.133)$ $0.791$ $91.0\%$ $0.6$ $0.591 (0.351-0.996)$ $0.048$ $  0.753 (0.485-1.171)$ $0.208$ $66.8\%$ $0.6$			
	OSCC	5	531	0.653 (0.292-1.462)	0.300	90.7%	P value           P value           %         <0.001           %         <0.001           %         <0.001           %         <0.001           %         <0.001           %         <0.001           %         <0.001           %         <0.003           %         0.003           %         0.001           %         0.001           %         0.001           %         0.001           %         0.001           %         0.001           -         -           %         0.001           -         -           %         0.001           -         -           %         0.016
	NPC	3	389	0.649 (0.458-0.920)	0.015	0.0%	
	OPSCC	3	495	0.975 (0.771-1.234)	0.835	0.0%	0.403
	LSCC	1	260	0.635 (0.393-1.025)	0.063	-	P value           %         <0.001
	Sample size						
	Large	7	1715	0.984 (0.659-1.468)	0.936	79.5%	< 0.001
	Small	10	866	0.582 (0.426-0.796)	0.001	50.3%	0.034
DFS	Overall	7	1066	0.928 (0.618-1.392)	0.717	69.4%	0.744 0.403 - <0.001 0.034 0.003 0.001 0.150 0.032
	Ethnicity						
	Asian	3	416	0.809 (0.241-2.720)	0.732	85.8%	6%         <0.001           1%         <0.001
	Caucasian	4	650	0.938 (0.663-1.328)	0.719	43.6%	
	Tumor locat	tion					
	OSCC	2	148	0.699 (0.114-4.263)	0.697	78.4%	0.032
	NPC	2	348	1.215 (0.288-5.133)	0.791	91.0%	0.001
	OPSCC	1	181	1.090 (0.783-1.518)	720)       0.732       85.8%         328)       0.719       43.6%         263)       0.697       78.4%         .133)       0.791       91.0%         .518)       0.610       -	-	
	LSCC	1	260	0.591 (0.351-0.996)	0.048	-	-
	Sample size						
	Large	3	650	0.753 (0.485-1.171)	0.208	66.8%	0.049
	Small	4	416	1.146 (0.536-2.450)	0.725	71.0%	0.016
PFS	Overall	3	188	0.618 (0.388-0.985)	0.043	0.0%	0.867

Table 4. Sensitivity analysis results for high-quality studies on the prognostic effects of PD-L1 in HNC patients.

In nasopharyngeal carcinoma (NPC) patients, the OS was better for those expressing higher levels of PD-L1 (HR = 0.692, 95% CI: 0.523-0.915, P = 0.010). However, no obvious trend in DFS, PFS, DSS or DMFS was found according to PD-L1 expression. In laryngeal squamous cell carcinoma patients, higher PD-L1 expression was associated with better DFS (HR = 0.591, 95% CI: 0.350-0.997, P = 0.048).

#### Sensitivity analysis and publication bias

A sensitivity analysis of the association between the expression of PD-L1 and the prognosis of HNC patients was performed for high-quality studies (NOS score  $\geq$  7, Table 4). The overall HRs and 95% CIs followed the same trends as those in the previous analysis. Higher levels of PD-L1 exhibited a trend of correlation with better OS (HR = 0.754, 95% CI: 0.568-1.002, *P* = 0.051, Figure 4A) and were associated with better PFS (HR = 0.618, 95% CI: 0.388-0.985, *P* = 0.043, Figure

4B) in the high-quality studies. As in the previous analysis, the OS of NPC patients was better in the high-PD-L1 group (HR = 0.649, 95% CI: 0.458-0.920, P = 0.015, Figure 4A). The heterogeneity among the studies decreased slightly for OS, but it remained statistically significant (I<sup>2</sup> = 76.6%,  $P_h < 0.001$ ; Table 4). In addition, subgroup analyses revealed that higher PD-L1 levels were associated with better OS in Caucasian patients (HR = 0.742, 95% CI: 0.578-0.954, P = 0.020) and in studies with small sample sizes (HR = 0.582, 95% CI: 0.426-0.796, P < 0.001, Table 4).

Funnel plots of OS were created for all the studies (Figure 5A), for the studies on PD-L1 (Figure 5B) and for the high-quality studies on PD-L1 (Figure 5C). For all three plots, the studies were distributed uniformly around the axis, manifesting no obvious publication bias (P = 0.509, 0.876 and 0.868 for all the studies, the studies on PD-L1 and the high-quality studies on PD-L1, respectively).



Figure 4: Overall forest plots of sensitivity analysis. (A) Stratified analysis based on the tumor location for the association between PD-L1 and OS. (B) Overall forest plots of sensitivity analysis for the association between PD-L1 and PFS.





#### **DISCUSSION**

As immune checkpoint molecules could be involved in the immune surveillance of tumor development and progression and the clearance of tumors [11], antiimmune-checkpoint drugs such as pembrolizumab [3, 19], nivolumab [26, 27] and ipilimumab [28] have been approved to treat melanoma, non-small cell lung cancer, renal cell carcinoma, prostate cancer and HNC. Recent studies have examined how immune checkpoint molecules, especially PD-L1, influence the prognosis of cancer patients, and a large number of updated reports have been published in the past two years. However, no consensus has been reached on the effects of immune checkpoint molecules on the prognosis of HNC.

This meta-analysis on the prognostic value of immune checkpoint molecules included 52 studies with a total of 7127 patients. The expression of immune checkpoint molecules was found to be a controversial prognostic factor for the OS, DFS, PFS, DSS and DMFS of HNC patients. Although the current view is that immune checkpoint molecules may be important predictors of a poor prognosis in HNC [17-19, 22, 29-31], our subgroup analysis stratified according to the immune checkpoint molecule revealed that different molecules had different associations with the patient prognosis. Thus, our results require careful attention.

Higher IDO expression was associated with a poorer prognosis for HNC patients in our study. Similarly, high IDO expression has been reported to correlate with a poor prognosis in patients with melanoma, breast cancer and colon cancer [32-34]. However, in our study, higher expression of PD-L1 tended to be associated with better OS. Kogashiwa et al. [35] found that higher expression of PD-L1 was associated with a higher number of CD8+ tumor-infiltrating lymphocytes, leading to better OS for HNC patients. Balermapas et al. and Hanna et al. [36, 37] also reported higher levels of tumor-infiltrating lymphocytes in HNC patients expressing higher levels of PD-L1, which could explain the improved OS of these patients.

As PD-L1 attracted the most attention of the included immune checkpoint molecules, and a large number of updated studies reported the relationship between high levels of PD-L1 and the prognosis of HNC in 2017 and 2018, we considered it important to conduct a further meta-analysis solely on this molecule. We found that higher PD-L1 expression was associated with better OS in NPC patients, although for HNC overall there was only a positive trend, rather than a concrete link (Figure 3). A sensitivity analysis revealed the same trends in OS. In addition, higher PD-L1 expression was found to correlate with better PFS. The results of the sensitivity analysis may be more dependable than the former results, as all the included studies were of high quality. Furthermore, the same relationship between PD-L1 expression and OS was confirmed in Caucasian subjects, NPC patients and studies with small sample sizes.

Tumors can develop adaptive immune resistance, which is one of the two mechanisms regulating tumor PD-L1 expression (the second being intrinsic immune resistance) [38]. While the intrinsic mechanism leads to PD-L1 expression after oncogenic mutation [39], the adaptive mechanism causes tumor cells to express PD-L1 after they have been stimulated by interferon gamma secreted by CD8+ T cells [40, 41]. Therefore, tumormembranous PD-L1 levels could partly reflect the amount of tumor-infiltrating lymphocytes, especially cytotoxic T cells, accounting to some extent for the better survival of patients with higher PD-L1 levels.

There are several limitations to this meta-analysis. Firstly, the overall heterogeneity was high, so random effects models were required for the analysis, and there was less sensitivity to detect significant differences. Secondly, all the included studies were prospective, and the majority of studies did not have adequate random sequences or comparable cohorts, increasing the risk of bias. Thus, the quality of the included studies was not perfect. Lastly, the study populations were all of Asian or Caucasian ethnicity, which may have caused a population selection bias.

Our meta-analysis indicated that different immune checkpoint molecules correlated with different prognoses in HNC patients: higher IDO expression predicted a poorer prognosis, while higher PD-L1 expression was associated with a better prognosis. Furthermore, our study revealed that higher expression of PD-L1 was associated with significantly better OS in Caucasian subjects, NPC patients and studies with small sample sizes. In summary, our study suggested that the immune checkpoint molecules IDO and PD-L1 have potential prognostic value and applicability to immune therapy for HNC.

#### **METHODS**

#### Literature-search strategy

carcinoma[Title/Abstract]) OR head and neck neoplasm\$[Title/Abstract]) OR HNSCC[Title/Abstract] ) OR SCCHN[Title/Abstract]) OR HNC[Title /Abstract]) OR mouth neoplasms[Title /Abstract]) OR cancers of mouth[Title/Abstract]) OR oral[Title /Abstract]) OR laryn\*[Title/Abstract]) OR pharyn\* [Title/Abstract]) OR tongue[Title/Abstract]) OR oropharyn\*[Title/Abstract]) OR nasopharyn\* [Title/ Abstract]) OR hypopharyn\*[Title/Abstract]) OR trachea[Title/Abstract]) OR laryngopharyn\* [Title /Abstract]) OR cervical tracheal[Title/Abstract]) OR cervical esophagus[Title/Abstract]) OR lip [Title /Abstract])) OR sinonasal[Title/Abstract]) OR head and neck cutaneous squamous cell carcinoma [Title/ Abstract]) OR squamous cell carcinoma of the oral cavity[Title/Abstract]) OR salivary gland carcinoma OR SGC[Title/Abstract]) [Title/Abstract]) AND [Title/Abstract]) OR b7-h4[Title/Abstract]) OR vtcn1[Title/Abstract]) OR btla[Title/Abstract]) OR b and t lymphocyte attenuator) OR cd272[Title/Abstract]) OR ctla-4[Title/Abstract]) OR cytotoxic t-lymphocyteassociated protein 4[Title/Abstract]) OR cd152[Title /Abstract]) OR ido[Title/Abstract]) OR indoleamine 2,3-dioxygenase[Title/Abstract]) OR kir[Title/Abstract] ) OR killer-cell immunoglobulin-like receptor[Title/ Abstract]) OR lag3[Title/Abstract]) OR lymphocyte activation gene-3[Title/Abstract]) OR pd-1[Title/ Abstract]) OR programmed death 1 receptor [Title/Abstract]) OR pd-l1[Title/Abstract]) OR programmed death ligand 1[Title/Abstract]) OR pd-12[Title/Abstract]) OR tim-3[Title/Abstract]) OR t-cell immunoglobulin domain and mucin domain 3[Title/Abstract]) OR vista[Title/Abstract]) OR vdomain ig suppressor of t cell activation[Title/Abstract]) OR b7-h1[Title/Abstract]). Two reviewers (Y.Q.J. and B.Y.) inspected all candidate articles independently. Discrepancies were resolved by discussion with the senior authors (Z.W. and B.C.).

#### Inclusion and exclusion criteria

The available prospective comparative studies (cohort studies) were included in this study based on their conformance to the following inclusion criteria: 1) the association of immune checkpoint marker expression with OS/DFS/PFS/DSS/DMFS in HNC was reported; 2) the diagnosis of HNC was made based on pathological examination; 3) HRs and 95% CIs were provided or could be estimated from the text; 4) only the more recent or complete article was selected when multiple reports described the same population, to avoid the duplicate inclusion of data; and 5) articles were published as original research.

The exclusion criteria were: 1) reviews, meeting abstracts, letters; 2) animal model studies; 3) sample size < 30 patients; 4) insufficient data to estimate the HR and 95% CI; 5) the main type of tumor was not SCC; 6) the number of studies on a single molecule was less than three; and 7) the study design was not prospective.

#### Data extraction and quality assessment

Two reviewers (Y.Q.J. and B.Y.) extracted the following information independently from the included studies: author, year of publication, study country or region, sample ethnicity, tumor location, follow-up period, sample size, gender, cut-off values of immune checkpoint molecules, detection method, TNM stage, and survival data such as OS, DFS, PFS, DSS and DMFS. The HR and 95% CI were either reported or calculated from the P value or Kaplan-Meier survival curve [42, 43]. Disagreements were resolved by a senior reviewer (Z.W.).

Two reviewers (L.L.W. and W.X.M.) independently assessed the quality of the included studies by the NOS. A score of 0–9 was given to each study, and studies with NOS scores  $\geq$  7 were defined as high-quality. Consensus was reached by discussion with senior reviewers (B.C. and Z.W.) when there were inconsistent results. Importantly, the procedure of assessing the quality of the studies was blinded to the reviewers who extracted the data (Y.Q.J. and B.Y.).

#### Statistical analysis

This meta-analysis was performed in accordance with recommendations from the Cochrane Collaboration and the Quality of Reporting of Meta-analyses guidelines [44, 45]. The HR was used as a summary statistic for censored outcomes (OS, DFS, PFS, DSS and DMFS). HRs > 1 represented a poor prognosis in HNC.

Heterogeneity among the primary studies was evaluated by Cochrane's Q statistic and the I<sup>2</sup> statistic. A *P* value < 0.10 in Cochrane's Q test or an I<sup>2</sup> value > 50%indicates substantial heterogeneity among studies, so a random effects model was used to calculate the pooled HR and 95% CI in such cases. Otherwise, a fixed effects model was applied.

We used the mean sample size as the boundary between studies with large and small sample sizes. Subgroup analyses were carried out according to the immune checkpoint molecule, ethnicity, sample size and tumor location. Sensitivity analysis was applied to highquality studies (NOS  $\geq$  7). Begg's funnel plots were used to assess publication bias. All statistical analyses were conducted with STATA 12.0 statistical software (Stata Corporation, College Station, TX, USA). A two-tailed P value < 0.05 was considered statistically significant.

#### Abbreviations

PD-L1: programmed death ligand 1; PD-1: programmed death 1; LAG-3: lymphocyte activation gene 3; CTLA-4: cytotoxic T-lymphocyte-associated protein 4; PD-L2: programmed death ligand 2; IDO: indoleamine 2, 3dioxygenase; VISTA: V-domain Ig suppressor of T cell activation; HR: hazard ratio; M/F: male/female; NA: not available; HNC: head and neck cancer; HNSCC: head and neck squamous cell carcinoma; OSCC: oral squamous cell carcinoma; NPC: nasopharyngeal carcinoma; OPSCC: oropharyngeal squamous cell carcinoma; LSCC: laryngeal squamous cell carcinoma; cut-off value: the value that can be diagnosed as positive/high expression of an immune checkpoint molecule; AQUA: automated quantitative analysis; IHC: immunohistochemistry; IF: immunofluorescence; qRT-PCR: quantitative reverse transcription polymerase chain reaction; FACS: fluorescence-activated cell sorting; FISH: fluorescence in situ hybridization; OS: overall survival; PFS: progression-free survival; DFS: disease-free survival; DSS: disease-specific survival; DMFS: distant metastases-free survival; P: prospective; NOS: Newcastle-Ottawa Quality Assessment Scale; CI: confidence interval.

#### **AUTHOR CONTRIBUTIONS**

Y.Q.J. and B.Y. conceived and designed the research, analyzed the data and wrote the paper. Z.W. and B.C. reviewed drafts of the paper and participated in its design and coordination. L.L.W. and W.X.M. evaluated the quality of the literature and prepared the figures and tables. All the authors read and approved the final manuscript.

#### **CONFLICTS OF INTEREST**

The authors declare no conflict of interest.

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## SUPPLEMENTARY MATERIAL

Supplementary Table 1. Risk of bias in prospective studies based on the modified Newcastle-Ottawa Scale.

		SELE	CTION		COMPARABILITY		OUTCOME		
Study	Representativeness of the Exposed Cohort	Selection of the Non- Exposed Cohort	Ascertainment of Exposure	Demonstration That Outcome of Interest Was Not Present at Start of Study	Comparability of Cohorts on the Basis of the Design or Analysis	Assessment of Outcome	Long Enough Follow-Up for Outcomes to Occur	Adequacy of Follow- Up of Cohorts	Quality Score
Ahn et al. 2017	0	1	1	1	1	1	1	1	7
Badoual et al. 2013	0	1	1	1	1	0	1	1	6
Balempas et al. 2017	0	1	1	1	1	1	1	1	7
Ben-Haj-Ayed et al. 2016	0	1	1	1	1	1	1	1	7
Birtalan et al. 2017	0	1	1	1	1	1	1	0	6
Budczies et al. 2016	0	1	1	1	1	1	0	0	5
Chan et al. 2017	0	1	1	1	1	1	0	1	6
Chang et al. 2017	0	1	1	1	1	1	0	0	5
Chen et al. 2015	0	1	1	1	1	1	1	1	7
Chen et al. 2017	0	1	1	1	1	1	1	1	7
Cho et al. 2011	0	1	1	1	1	1	1	0	6
De et al. 2017	0	1	1	1	1	1	0	1	6
Fang et al. 2014	0	1	1	1	1	1	0	1	6
Feng et al. 2017	1	1	1	1	1	1	1	0	6
Fiedler et al. 2018	0	1	1	1	1	1	1	1	7
Hanna et al. 2018	0	1	1	1	1	1	1	1	7
Hong et al. 2016	0	1	1	1	1	1	0	1	6
Hsu et al. 2010	Ő	1	1	- 1	1	0	Õ	0	4
Kansy et al. 2017	0	1	1	1	1	0	1	1	6
Kim et al. 2016	0	1	1	1	1	1	1	0	6
Kim et al. 2016	0	1	1	1	1	1	1	1	7
Kogashiwa et al. 2017	0	1	1	1	1	1	1	1	7
Laimer et al.	0	1	1	1	1	1	1	1	7

2011									
Larbcharoensub	0	1	1	1	1	1	1	1	7
et al. 2018									
Lee et al. 2016	0	1	1	1	1	0	1	0	5
Li et al. 2017	0	1	1	1	1	0	1	0	5
Lin et al. 2015	0	1	1	1	1	0	1	1	6
Muller et al.	0	1	1	1	1	1	0	1	6
2017									
Ock et al. 2016	0	1	1	1	1	1	0	1	6
Oguejiofor et al.	0	1	1	1	1	1	1	1	7
2017									
Oliveira-Costa et	0	1	1	1	0	1	1	1	6
al. 2015									
Ono et al. 2017	0	1	1	1	1	1	1	1	6
Ono et al. 2018	0	1	1	1	1	1	1	1	7
Ou et al. 2017	0	1	1	1	1	1	1	1	7
Qu et al. 2017	0	1	1	1	1	1	0	1	6
Riobello et al.	0	1	1	1	1	1	0	1	5
2018									
Roper et al. 2017	0	1	1	0	1	1	1	1	6
Satgunaseelan et	0	1	1	1	1	1	0	1	6
al. 2016									
Schneider et al.	0	1	1	1	1	1	1	1	7
2018									
Seppälä et al.	0	1	1	1	0	1	1	1	6
2016	<u>^</u>							_	_
Solomon et al.	0	1	1	1	I	1	1	1	1
2018	0								-
Steuer et al.	0	1	1	1	1	1	1	1	
2018	0	1	1	1	1	1	0	0	~
Strati et al. 2017 Straub et al.	0	1	1	1	1	1	0	0	5
2016	0	1	1	1	1	1	1	1	/
Tang et al. 2017	0	1	1	1	1	1	1	0	6
Ukpo et al. 2013	0	1	1	1	1	1	1	0	0
Vassilakopoulou	0	1	1	1	1	1	1	1	7
et al. 2015	0	1	1	1	1	1	1	1	/
Ye et al. 2013	0	1	1	1	1	1	0	1	6
Zhang et al.	0	1	1	1	1	1	1	1	07
2015	0	1	1	1	1	1	1	1	/
Zheng et al.	0	1	1	1	1	1	1	0	6
2017	v	T	1	1	ī	1	1	0	0
Zhu et al. 2017	0	1	1	1	1	1	1	1	7
	\$	-	Ŧ	1	-	±	1		,