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Propensity Score Matched Analysis on Survival Profile of Head and Neck Cancer in the U.S. Hispanic Population

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2015

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Abstract<br>Propensity Score Matched Analysis on Survival Profile of Head and Neck Cancer in the U.S. Hispanic Population<br>By Xiting Zhu

Introduction: Head and neck cancer (HNC) ranks as the $10^{\text {th }}$ most common cancer worldwide and the $7^{\text {th }}$ most common cancer inducing mortality. It causes 300,000 deaths each year worldwide among 600,000 newly diagnosed cases. A better understanding of potential risk factors for the incidence of HNC should be completed as well as thorough survival analyses for each racial and ethnic population. Despite the continuous increase in the U.S. Hispanic population in the recent decade, however, limited data was published on the survival of HNC in the Hispanic population. The purpose of this paper is to compare overall survival (OS) and disease-specific survival between Hispanic and the general HNC patients of the United States and to identify prognostic factors that might affect the survival outcomes of HNC patients.

Methods: The Surveillance, Epidemiology, and End Results (SEER) database was queried for HNC patients diagnosed between 2000 and 2013. Kaplan-Meier and log-rank test were performed to compare OS and disease-specific survival by racial-ethnical groups. A multivariable Cox proportional hazard model was fit to select prognostic factors. In addition, a multivariable model was fit to test for the impact of interaction between race-ethnic groups and age on survival outcomes. A method of propensity score matching was also implemented.

Results: A total of 113885 patients were identified in the study; 8132 (7.1\%) were Hispanic, 12669 ( $11.1 \%$ ) were non-Hispanic Black, 85995 ( $75.5 \%$ ) were non-Hispanic White, and 7089 $(6.2 \%)$ were non-Hispanic other. 31720 ( $27.9 \%$ ) patients were diagnosed with laryngeal cancer, 49850 ( $43.8 \%$ ) were oral cavity cancer, and 32315 ( $28.4 \%$ ) were pharyngeal cancer. In patients with HNC, Hispanic race was associated with poor median OS and 5 -year overall survival rate compared with non-Hispanic White ( 72 vs. 80 months; $53 \%$ vs. $56 \% ; P<0.0001$ ). In the propensity score matched sample, a poorer OS for Hispanic patients versus non-Hispanic White patients was obtained (HR, $1.09 ; 95 \%$ CI, 1.01-1.16; $P=0.021$ ). Inferior median OS and 5-year overall survival rate remained associated with Hispanic race (compared with non-Hispanic White: 64 months vs. 78 months; $51 \%$ vs. $55.6 \% ; P<0.0001$ ).

Conclusions: Hispanic race was associated with poor OS and disease-specific survival in HNC patients. A series of prognostic factors resulting in inferior survival were found in Hispanic HNC patients.

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## 1. INTRODUCTION

Head and neck cancer (HNC) ranks as the $10^{\text {th }}$ most common cancer worldwide and the $7^{\text {th }}$ most common cancer inducing mortality (World Health Organization, 2008). It causes 300,000 deaths each year worldwide among 600,000 newly diagnosed cases, with the most common cancer site being the larynx, the oral cavity, and the pharynx (Boyle \& Levin, 2008). Smoking, alcohol use, genetic factors, and viral infection are known as the most prominent risk factors identified so far to induce the incidence of HNC, with the most important prognostic factors being cancer site and tumor TNM stage (Mehanna et al., 2010a). The standard treatment which has been widely accepted for HNC is combined surgery and radiotherapy along with rehabilitation (Mehanna et al., 2010a). However, the racial disparities in HNC and other socioeconomically disadvantaged factors of the American population adversely affect the treatment and its outcome (Goodwin et al., 2007). The high prevalence and mortality rates of HNC urge the need of thorough survival analyses for each racial and ethnic population as well as better understanding of potential risk factors to the incidence of HNC.

### 1.1 Problem Statement

In the United States, 50.5 million people (or 16 percent) among the total population selfreported their ethnicity as Hispanic or Latino according to the 2010 census with an increase of 15.2 million people since 2000 (Ennis et al., 2011). This accounts for over half of the 27.3 million increase in the total population of the United States (Ennis et al., 2011). However, there is little information published regarding the survival of HNC in the Hispanic population given the fact
that the Hispanic population is less likely to be involved in clinical trial studies (Murthy et al., 2004).

### 1.2 Purpose Statement

The purpose of this paper is to compare overall survival (OS) and disease-specific survival between four race/ethnicity groups: Hispanic, non-Hispanic White, non-Hispanic Black and nonHispanic other patients of the United States and to identify prognostic factors that are associated with the survival outcomes of HNC patients. To reduce potential bias from its retrospective design, a propensity score analysis was performed by balancing the baseline demographic and socioeconomic factors among different racial groups.

### 1.3 Significance Statement

This paper will provide understanding of survival in HNC in the Hispanic population and how the survival varies by site. This will assist in diagnosis and treatment for these particular cancers in the Hispanic population. Moreover, this project could investigate different risk factors in this population to help with developing new preventive interventions, evidence of early detection or innovative therapies for HNC .

## 2. BACKGROUND/LITERATURE REVIEW

## Study Topic: Head and Neck Cancer in Hispanic Population

### 2.1 Head and Neck Cancer Overview

Head and neck cancers include cancers of the upper aerodigestive tract (including oral cavity, nasopharynx, oropharynx, hypopharynx and larynx), the paranasal sinuses, and the salivary glands (Mehanna et al., 2010b). In the United States, the estimated incidence and mortality in 2015 were 59,340 new cases and 12,290 deaths, of which more than $90 \%$ are squamous cell carcinomas (Wyss et al., 2016). The epidemic of HNC demonstrates a trend that men and elders have higher incidence rates compared with women and younger population. Large scale epidemiological studies and national cancer registries reported that the incidence rate for male versus female varies from 2:1 to 15:1 depending on the site of cancer (Mehanna et al., 2010b). In Europe, $98 \%$ and $50 \%$ newly diagnosed patients are middle-aged and elderly people over 40 and 60 years of age, respectively (Mehanna et al., 2010b). The Indian subcontinent, Australia, France, Brazil, and Southern Africa suffer from a higher incidence of HNC while the incidence of oral cavity, laryngeal and other tobacco use related cancers is declining in North America and western Europe likely due to decreased exposure to tobacco (Boyle \& Levin, 2008). Some developing countries which have the highest tobacco prevalence due to lagged tobacco free policy or regulation, have relative high incidence of HNC (Mehanna et al., 2010b). Although each particular cancer site has different courses and histopathological types of HNC, squamous cell carcinoma is the most common one (Mehanna et al., 2010b). The cancers and their treatments
heavily affect the corresponding anatomical sites on functions such as speech, swallowing, taste, and smell so that the patients' life quality cannot be assured (Mehanna et al., 2010b).

Known risk factors include tobacco and alcohol use, HPV infection of the mouth and throat, and genetic factors such as family history of HNC (Mehanna et al., 2010b). Many studies suggested a synergistic relationship between smoking and alcohol use that results in a 30 -fold increased risk for individuals who both smoke and drink heavily (American Cancer Society, 2017). Non-smokers who have three or more alcoholic drinks per day have double the risk of developing the disease compared with non-drinkers, reported by pooled analyses of 15 casecontrol studies (odds ratio: 2.04, 95\% confidence interval: 1.29 to 3.21) (Mehanna et al., 2010b). Individuals who have family history of HNC and who are infected by HPV have 1.7-fold (95\% CI, 1.2 to 2.3 ) and 63 -fold ( $95 \%$ CI, 14 to 480 ) increased risk of developing HNC, respectively. (Mehanna et al., 2010b).

Patients with HNC usually present a variety of symptoms, depending on the function of the cancer original site (Mehanna et al., 2010a). Laryngeal cancers usually present with hoarseness, while pharyngeal and oral cavity cancers normally present lesion or lump in the throat or mouth, ear pain, a neck mass or coughing up blood (American Cancer Society, 2017). Prevention of HNC is closely linked to the success in control of tobacco use. Previous studies demonstrated that quitting from smoking for 1-4 years reduces the risk of developing HNC ( $0.70,95 \% \mathrm{CI} 0.61$ to 0.81 ) compared with current smoking, and further reduces the risk for 20 years or more ( 0.23 ,
$95 \%$ CI 0.18 to 0.31 ) (Oxford Cancer Intelligence Unit, 2010). Benefit of quitting alcohol use, however, was only seen for 20 years or more of quitting time (Marron et al., 2010).

### 2.2 Prognosis and Treatment

The most important prognostic factors are site and tumor TNM (tumor, node, metastasis) stage (Mehanna et al., 2010a). The average survival rate for HNC at 5 -years is $34.4 \%$, with a survival rate of $44.9 \%$ for women and $31.0 \%$ for men, respectively (Gatta et al., 2015). The survival rates for HNC on each of the three major cancer sites (oral cavity, larynx and pharynx), however, vary study by study. Generally, the overall 5-year survival rate was in the range of 20$43 \%$ for oral cancer, $8-25 \%$ for pharyngeal cancers and $25-62 \%$ for laryngeal cancer (Rao et al., 1998). Another prominent prognostic factor is comorbid illness, which was proved to be associated with poor prognosis (Mehanna et al., 2010a).

Combined surgery and radiotherapy along with rehabilitation have now become the standard treatment for HNC, often encouraged by national guidelines and protocols (National Institute for Health and Clinical Excellence, 2004). Radiotherapy and cancer-directed surgery are two common and widely accepted treatments for HNC. The choice of treatment depends on individual factors related to the site of the tumor and cancer stage, but also patient preference (Mehanna et al., 2010a). For early stage tumors, radiotherapy is more superior than cancer-directed surgery for better organ preservation (Scottish Intercollegiate Guidelines Network). For advanced tumors, on the other hand, combined use of surgery and postoperative radiotherapy, or combined
chemotherapy and radiotherapy, ensure the highest chance of achieving the best outcome (Bhalavat et al., 2003).

### 2.3 Influence of Race/Ethnicity

According to the 2010 Census, 308.7 million people resided in the United States, of which 50.5 million people (or 16 percent) were of Hispanic or Latino origin (Ennis et al., 2011). The Hispanic population increased by 15.2 million from 35.3 million (13 percent) in 2000 , which accounted for over half of the increase in the total population of the United States (Ennis et al., 2011). Between 2000 and 2010, the Hispanic population grew by 43 percent, which was four times the growth of the total population (Ennis et al., 2011).

Given the huge increase in the U.S. Hispanic population, however, there is very little data published on the survival of cancer in the Hispanic population, particularly in HNC. Compared with a $1.8 \%$ cancer research enrollment proportion among White patients, a relatively low enrollment rate was seen in Hispanic patients ( $1.3 \%$; odds ratio vs. whites, $0.72 ; 95 \% \mathrm{CI}, 0.68$ to 0.77) (Murthy et al., 2004). Moreover, Shavers et al studied patterns of care, socioeconomic status and comorbidity in 471 patients with HNC and found racial disparities for the receipt of cancer treatment among patients with particular types of HNC (Shavers et al., 2003). Shavers et al also found that the differences in cancer stage and the lower rates of cancer screening for Hispanic patients compared with White patients result in different rates of early detection, which in turn contributes to racial differences in survival (Shavers et al., 2003).

## Statistical Method: Propensity Score

### 2.4 Overview

Propensity score methods are widely used to analyze causal association between treatment and outcome in observational studies to reduce confounding that may affect the estimated effects of treatment. The propensity score is the probability of treatment assignment conditional on observed baseline covariates: $e_{i}=\operatorname{Pr}\left(Z_{i}=1 \mid X_{i}\right)$ where $Z_{i}$ is an indicator variable denoting whether or not the $i$ th subject was treated and $e_{i}$ denotes the propensity score of the $i$ th subject given his/her set of baseline covariates $X_{i}$ (Austin, 2011). The propensity score allows one to design and analyze an observational (nonrandomized) study since it mimics some particular characteristics of a randomized control trial (Austin, 2011). After matching based on the propensity scores, the distributions of measured baseline covariates are similar between treatment and control subjects (Austin, 2011). Consequently, the distribution of the observed baseline covariates will be the same between the treatment and control groups in a set of subjects who have the same propensity score (Austin, 2011). The implementation is based on a logistic regression model, in which treatment status is regressed on selected observed baseline covariates. The predicted probability of treatment derived from the logistic regression model is the propensity score.

### 2.5 Methodological decisions

There are four different propensity score methods used for removing the effects of confounding: propensity score matching, stratification (or subclassification) on the propensity
score, inverse probability of treatment weighting (IPTW) using the propensity score, and covariate adjustment using the propensity score (Rosenbaum \& Rubin, 1983). Among these methods, propensity score matching and inverse probability of treatment weighting using propensity score allow one to estimate the marginal effect of treatment on survival when estimating the effect of treatment on time-to-event outcomes using observational or nonrandomized data (Austin, 2012). However, IPTW outperforms propensity score matching in estimates with lower MSE, proving that the estimates resulting from IPTW have improved precision than those resulting from propensity score matching (Austin, 2012).

Matching on the propensity score generates matched pairs of treated and untreated subjects whose values of propensity score are similar (Rosenbaum \& Rubin, 1985). Similarity in the values of propensity score results in homogeneous distribution of the observed baseline covariates. Once the matched sample has been formed, the treatment effects can be directly estimated by comparing the outcomes from treated and untreated subjects in the matched sample (Austin, 2011). Several decisions must be made prior to forming matched pairs of treated and untreated subjects.

First, a decision of matching without replacement or matching with replacement needs to be considered. When using matching without replacement, once an untreated subject has been selected to be matched to a treated subject, that untreated subject is no longer available as a potential match for subsequent treated subjects (Austin, 2011). In contrast, matching with
replacement allows a given untreated subject to be included in more than one sets of matching (Austin, 2011).

A second choice is between greedy and optimal matching (Rosenbaum, 2002). The philosophy of greedy matching is that a treated subject is first selected at random. The untreated subject whose propensity score is closest to that of this randomly selected treated subject is chosen for matching to this treated subject (Austin, 2011). This process is then repeated until untreated subjects have been matched to all treated subjects or until the list of treated subjects for whom a list of untreated subjects can be found has been exhausted (Austin, 2011). An alternative to greedy matching is optimal matching, in which matches are formed in order to minimize the total within-pair difference of propensity score (Austin, 2011). However, $\mathrm{Gu} \&$ Rosenbaum found that optimal matching did not perform better than greedy matching in producing balanced matched sample (Gu \& Rosenbaum, 1993).

Third, a choice between nearest neighbor matching (NNM) and nearest neighbor matching within a specified caliper distance is then raised (Rosenbaum \& Rubin, 1985). NNM selects for matching to a given treated subject that untreated subject whose propensity score is closest to that of the treated subject (Austin, 2011). One important setting here is that there are no restrictions upon the maximum acceptable distance between the propensity scores of two matched subjects (Austin, 2011). Nearest neighbor matching within a specified caliper distance resembles NNM with the further restriction that the absolute difference in the propensity scores of matched subjects must be below the pre-specified arbitrary caliper distance (Austin, 2011). If no untreated
subjects whose propensity scores lay within the pre-specified caliper distance of the propensity score of the treated subject, that treated subject would not be matched with any untreated subject (Austin, 2011). Then unmatched treated subject would be then excluded from the matched sample so that bad matches are avoided and the matching quality is ensured.

Inverse probability of treatment weighting (IPTW) using propensity score uses weights calculated by propensity score to create a synthetic sample in which the distribution of measured baseline covariates is independent of treatment groups (Austin, 2011). Rosenbaum first proposed that weights can be defined as $w_{i}=\frac{z_{i}}{e_{i}}+\frac{\left(1-z_{i}\right)}{1-e_{i}}$ (Rosenbaum, 1987). Then the weights based on the inverse probability of treatment can be used on regression models to estimate causal effects of treatments (Joffe et al., 2004).

In terms of the balance diagnostics, comparing the similarity of baseline covariates for treated and untreated subjects in the propensity score sample should begin with a comparison of the means or medians of continuous covariates and the distribution of categorical covariates between treated and untreated subjects (Austin, 2011). The standardized difference is widely used in comparing the mean of continuous and binary variables between treatment groups (multilevel categorical variables can be represented as several binary indicator variables) (Flury \& Riedwyl, 1986). For a continuous covariate, the standard difference is given by:

$$
d=\frac{\left(\bar{x}_{\text {treatment }}-\bar{x}_{\text {control }}\right)}{\sqrt{\frac{s^{2}{ }_{\text {treatment }}+s^{2}{ }_{\text {control }}}{2}}}
$$

where $\bar{x}_{\text {treatment }}$ and $\bar{x}_{\text {control }}$ denote the sample mean of the covariate in treated and untreated subjects, whereas $s^{2}{ }_{\text {treatment }}$ and $s^{2}{ }_{\text {control }}$ denote the sample variance of the covariate in
treated and untreated subjects (Austin, 2011). For dichotomous variables, the standardized difference is defined as:

$$
d=\frac{\left(\hat{p}_{\text {treatment }}-\hat{p}_{\text {control }}\right)}{\sqrt{\frac{\hat{p}_{\text {treatment }}\left(1-\hat{p}_{\text {treatment }}\right)+\hat{p}_{\text {control }}\left(1-\hat{p}_{\text {control }}\right)}{2}}}
$$

where $\hat{p}_{\text {treatment }}$ and $\hat{p}_{\text {control }}$ denote the prevalence of mean of the dichotomous variable in treated and untreated subjects (Austin, 2011). The standardized difference compares the difference in means in units of pooled sample variance. Moreover, it ignores the influence of sample size and allows for comparison between variables measured in different units (Austin, 2011). Even though there is no universal agreement upon what threshold of standardized difference can be used to indicate significant imbalance, a standardized difference that is less than 0.1 is considered as a criteria indicating a negligible difference in the mean or prevalence of a covariate between treatment and control groups (Austin, 2011).

### 2.6 Propensity Score Estimation for Multiple Treatments

The method of propensity score has been mostly used for 2 treatment settings. However, there are many cases in clinical research that one might be interested in comparison of more than 2 treatments. For nominal treatments, Imbens proposed the use of generalized propensity score, defined as the conditional probability of receiving a particular level of the treatment given a set of measured baseline covariates (Imbens, 2000). The generalized propensity score can be estimated with a multinomial logistic or probit regression (Spreeuwenberg et al., 2010). It predicts the probability that an outcome (e.g., treatment assignments, $Z_{i}$ ) equals each of its possible values $(t=1, \ldots, M)$ as a function of a linear combination of the covariates and their products and cross
products:

$$
\begin{gathered}
P\left(Z_{i}=t \mid X_{i}\right)=\frac{e^{\beta_{t}^{\prime} X_{i}}}{1+\sum_{t^{\prime}=1}^{M-1} e^{\beta_{t^{\prime}}^{\prime} X_{i}}}, t=1, \ldots, M-1, \\
P\left(Z_{i}=M \mid X_{i}\right)=\frac{1}{1+\sum_{t^{\prime}=1}^{M-1} e^{\beta_{t^{\prime}}^{\prime} X_{i}}}
\end{gathered}
$$

where $\beta_{t}$ are unknown coefficients and estimated from the data (McCaffrey, 2013). For each subject, the probability of receiving each treatment given the observed set of baseline covariates is estimated (Spreeuwenberg et al., 2010). Imbens proved that, the generalized propensity score, like the simple propensity score, is a balancing score which can be used for sample matching (Imbens, 2000). Therefore, the generalized propensity score can be used to balance the measured baseline differences and allows for valid estimates in multiple treatment comparisons (Spreeuwenberg et al., 2010).

Once generalized propensity score is estimated for each patient on $M$ levels of treatment assignments, the matching is generalized from a one-dimenstional line to a $M-1$ dimentional space formed by propensity score vectors. The algorithm tries to search for clusters of subjectsone receiving each of treatments $t=1, \ldots, M$-with the smallest within-cluster distance (Rassen et al., 2013). The goal is to find an optimal within-cluster distance to limit required computations for all possible combinations of subjects (Rassen et al., 2013). Pairwise standard differences are used for balance check for baseline covariates across treatment groups after matching (Austin, 2009).

## 3. MATERIALS \& METHODS

### 3.1 Study Design

The present study utilized the Surveillance, Epidemiology, and End Results (SEER) database, a National Cancer Institute program that provides information on the incidence and survival rates of cancer in the United States and that includes population-based cancer registries covering approximately $28 \%$ of the population of the United States.

Eligible patients were those who were diagnosed with HNC between 2000 and 2013 and whose cancer diagnosis is the only one or the first during the patient's life time. This helps reduce the confounding effect from the treatment for previous cancer diagnosis. Furthermore, patients with unknown survival information were excluded from the study. From the SEER database, a total of 113,994 patients were diagnosed with HNC and 109 patients were excluded from the study because of history of multiple synchronous cancers. As a result, 113,885 patients (Hispanic, $\mathrm{n}=8132$; non-Hispanic White, $\mathrm{n}=85995$; non-Hispanic Black, $\mathrm{n}=12669$; non-Hispanic other, n $=7089)$ were included and analyzed for this study.

Basic demographics, socioeconomic status, characteristics of disease, survival information and underlying prognostic factors were examined, including age, race, gender, marital status, insurance status and type, education, unemployment rate, area of residence (rural or urban county), movement history (moved from outside US), smoking history, year of diagnosis (20002006 or 2007-2013), cancer-directed surgery status (accepted or not), site of cancer, tumor grade, tumor size and combined SEER summary stage. Continuous variables education, ever smoker,
moved from outside U.S., unemployed and urban were defined as the proportion of adults in the patient's area of residence that did not graduate high school, that used to smoke or currently smoking, that moved from outside U.S., that were unemployed, and that lived in urban area. All continuous variables were collapsed based on their quartiles and only collapsed variables were analyzed in the study. Sites of HNC were classified as larynx, oral cavity (lip, tongue, floor of mouth, gum and other mouth) and pharynx (nasopharynx, tonsil, oropharynx, hypopharynx). Due to the small sample size in certain levels in the "race and origin code" variable, levels of "nonHispanic American Indian/Alaska Native", "non-Hispanic Asian or Pacific Islander" and "nonHispanic unknown race" were collapsed in "non-Hispanic other".

The primary outcome of the study was overall survival, which was defined as months from date of registration in SEER to death or last follow-up. Secondary outcome was disease-specific survival, which was defined as months from date of registration in SEER to death of any types of head and neck cancers or last follow-up. Death due to other reason was censored at date of death.

### 3.2 Statistical Analysis

Statistical analysis was conducted using SAS version 9.4 and SAS macros or software developed at the Biostatistics and Bioinformatics at Winship Cancer Institute (Nickleach et al., 2013). Descriptive statistics for each variable were reported. The univariate association with racial subgroups (Hispanic, non-Hispanic White, non-Hispanic Black, non-Hispanic other) was assessed using the chi-square test for categorical covariates and analysis of variance for continuous covariates. The univariate association between each covariate including race groups
and study outcomes (overall survival and disease-specific survival) were assessed using Cox proportional hazards models and log-rank tests. A multivariable Cox proportional hazard model was fit by a backward variable selection method applying an alpha $=.05$ removal criteria in order to select covariates. Insurance status was removed from multivariate analysis due to missing information for all subjects who were diagnosed before 2007. Kaplan-Meier plots were produced to compare the survival curves by subgroups along with log-rank $P$-value. In addition, a multivariable model was fit to test for the impact of interaction between race and age on survival outcomes to explore how the impact of race on survival outcomes varies by different age groups. The survival information of each race/ethnicity subgroup stratified by age were estimated along with $P$-value. All $P$-values were two-sided, and $P<.05$ was considered statistically significant. One thing to note is that the $P$-values are very significant in this study due to large sample size. Thus, interpretation with focusing on clinical relevance is highly recommended.

Propensity score matching between Hispanic, non-Hispanic White, non-Hispanic Black and non-Hispanic other groups based on the estimated propensity score for each subject was performed to reduce heterogeneity in distributions of measured baseline covariates. A multinomial logistic regression model predicting Hispanic, non-Hispanic Black and non-Hispanic other groups, which together determine the propensity score of non-Hispanic White, was used to calculate propensity scores including the covariates age, gender, marital status, year of diagnosis, cancer-directed surgery status, site of cancer, tumor grade, tumor size, combined SEER summary stage, education, unemployment rate, area of residence, movement history, and smoking history.

Cases from the four race groups were matched to each other based on generalized propensity scores using a caliper-based greedy matching without replacement algorithm (Austin, 2011). The sums of within-cluster distance were calculated and ploted versus their corresponding matching ID. The caliper for the matching algorithm was identified as the caliper for the matching ID as to when the sum of within-cluster distance began increasing dramatically. Then the matching process was repeated for each of three types (laryngeal, oral cavity, and pharyngeal) of HNC. The effectiveness of the matching was evaluated by calculating the standardized differences of the covariates on the matched sample.

## 4. RESULTS

A total of 113885 patients were registered between 2000 and 2013 and met study entry criteria. 8132 ( $7.1 \%$ ) patients were Hispanic (all races), 12669 (11.1\%) were non-Hispanic Black, $85995(75.5 \%)$ were non-Hispanic White, and $7089(6.2 \%)$ were non-Hispanic other. A total of 3092 (2.7\%) patients were 20-39 years of age, $47943(42.1 \%)$ were $40-59$ years of age, 52617 ( $46.2 \%$ ) were $60-79$ years of age, and $10233(9.0 \%)$ were 80 and above years of age. The majority of patients had oral cavity cancer ( $n=49850 ; 43.8 \%$ ), while 31720 ( $27.9 \%$ ) patients had laryngeal cancer and 32315 ( $28.4 \%$ ) patients had pharyngeal cancer. The median tumor size was 2.7 cm (Table 1).

A comparison of patient, disease, and prognostic characteristics by race is presented in Table 2. Younger Hispanic population (20-39 years of age and 40-59 years of age) was more likely to be involved in this study compared with non-Hispanic White. Hispanic population was more likely to be not covered by insurance compared with non-Hispanic White. Furthermore, low education level and high unemployment rate were found in Hispanic population compared with non-Hispanic White. Finally, proportion of smokers in Hispanic population was significantly lower than that in non-Hispanic White population.

Factors associated with improved OS on univariate analysis were younger age, non-Hispanic other races, male gender, married status, being covered by insurance, later years of diagnosis, cancer-directed surgery received, lower tumor grade, smaller tumor size, localized and regional combined SEER summary stage, higher education level, lower unemployment rate, and lower
proportion of smokers in community (Table 3.1). In addition, the result showed that Hispanic population had a poorer OS compared with non-Hispanic White (HR, 1.07; 95\% CI, 1.03-1.10; $P$ $<0.001$ ). However, for patients with laryngeal cancer, the OS for Hispanic population was better compared with non-Hispanic White (HR, $0.93 ; 95 \%$ CI, $0.87-0.99 ; P=0.015$; Table 3.2). In terms of the disease-specific survival, factors associated with improved performance were similar to that of OS except for the exclusion of non-Spanish-Hispanic-Latino population and larynx and oral cavity cancer sites (Table 3.3).

Factors that remained associated with improved OS on multivariate analysis were younger age, non-Hispanic other races, male gender, married status, later years of diagnosis, cancerdirected surgery received, pharynx cancer site, tumor grade II and III, smaller tumor size, localized combined SEER summary stage, higher education level, unemployment rate $\geq 1 \%, \leq$ $5 \%$, moved from outside U.S. $>5 \%, \leq 10 \%$, ever smoker $\geq 29 \%, \leq 45 \%$ (Table 4.1). Again, a poorer OS was obtained for Hispanic population versus non-Hispanic White (HR, 1.06; 95\% CI, 1.03-1.10; $P<0.001$ ) while a better OS for Hispanic laryngeal cancer patients was found compared with non-Hispanic White laryngeal cancer patients (HR, 0.91; 95\% CI, 0.85-0.97; $P=$ 0.004; Table 4.2). For disease specific survival, similar prognostic factors of significance were found (Table 4.3). There was a significant interaction between age and race ( $P<0.001$ ). When the range of patient's age was in 40-59 or 60-79, OS for Hispanic population was significantly inferior to the for non-Hispanic White or non-Hispanic other (Table 5.1). The same interaction effect was also found in the multivariate analysis for disease-specific survival (Table 5.2).

The median OS for Hispanic population was 72 months, compared with 80 months for nonHispanic White and 102 months for non-Hispanic other races ( $P<0.0001$; Figure 1.1). The overall 5 -year survival rate was $53 \%$ for Hispanic population, $56 \%$ for non-Hispanic White, $38.2 \%$ for non-Hispanic Black and $59.2 \%$ for non-Hispanic other races. Similar trends were obtained from the survival analysis for oral cavity and pharyngeal cancer. A comparable survival trend was obtained for disease-specific survival ( $P<0.0001$; Figure 1.2). For oral cavity cancer, the 5-year survival rate was $52 \%$ for Hispanic patients, $56.8 \%$ for non-Hispanic White, $34.2 \%$ for non-Hispanic Black and 58.1 for non-Hispanic other races ( $P<0.0001$; Figure 1.3). And for pharyngeal cancer, 5-year survival rates of $51.8 \%$ for Hispanic, $56 \%$ for non-Hispanic White, $32 \%$ for non-Hispanic Black and $59.2 \%$ for non-Hispanic other were seen ( $P<0.0001$; Figure 1.4). However, for patients with laryngeal cancer, the median OS was 80 months for Hispanic population versus 73 months for non-Hispanic White ( $P<0.0001$; Figure 1.5). Furthermore, the 5-year survival rate for laryngeal cancer was also higher for Hispanic patients than that for nonHispanic White, with $55.6 \%$ for Hispanic versus $54.6 \%$ for non-Hispanic White.

As demonstrated in Table 2, there was statistical difference in all covariates included in the multivariable models across the four groups. To adjust the unequal distributions of measured baseline covariates and to mimic the setting in a randomized study, a propensity score matching was performed to generate subgroups of 3230 patients from each group (Table 6). A caliper was applied for the matching algorithm to optimize the within-cluster distances and to limit the required computations (Figure 2.1). The result showed no difference in the distributions of some
important factors such as age, gender, marital status, cancer-directed surgery status, oral cavity and pharynx of cancer site, tumor grade, tumor size, combined SEER summary stage, education, and etc., in the four subgroups. Propensity score matching was also implemented for 3 major sites of HNC to balance the baseline covariates, with 969 laryngeal cancer patients, 1220 oral cavity cancer patients, and 1016 pharyngeal cancer patients from each subgroup. Calipers for each of the three matching algorithms were determined based on the plots of sums of within-cluster distance versus matching ID (Figure 2.2-2.4).

The impact of race on OS was investigated within the propensity score matched sample (Table 7.1). A poorer OS for Hispanic patients versus non-Hispanic White patients was obtained, which was consistent with the result from multivariate analysis before matching (HR, 1.09; 95\% CI, 1.01-1.16; $P=0.021$ ). The result of disease-specific survival analysis for Hispanic patients compared with non-Hispanic White patients was similar to that of OS (HR, $1.13 ; 95 \% \mathrm{CI}, 1.04-$ 1.23; $P=0.005$; Table 7.2). The median OS was 64 months for Hispanic patients versus 78 months for non-Hispanic White patients, and the 5-year survival rate was $51 \%$ for Hispanic patients and was $55.6 \%$ for non-Hispanic White ( $P<0.0001$; Figure 3.1). A similar trend was seen in terms of the disease-specific survival, with 5-year survival rate of $60.5 \%$ for Hispanic patients and $65.3 \%$ for non-Hispanic White patients ( $P<0.0001$; Figure 3.2). In terms of oral cavity ( $P=0.0001$; Figure 3.3) and pharyngeal cancer $(P<0.0001$; Figure 3.4), poor median OS (oral cavity cancer, 63 months; pharyngeal cancer, 64 months) and 5-year survival rate (oral cavity cancer, $51 \%$; pharyngeal cancer, $64 \%$ ) were found for Hispanic population. In contrast, for
laryngeal cancer patients, OS for Hispanic population was found to be better than that for nonHispanic White (HR, $0.84 ; 95 \%$ CI, $0.74-0.96 ; P=0.01$; Table 7.3). Similarly, the median OS was higher for Hispanic laryngeal cancer patients than that for non-Hispanic White laryngeal cancer patients, with 91 months for Hispanic and 71 months for non-Hispanic White ( $P<0.0001$;

Figure 3.5).

## 5. DISCUSSION

The purpose of this analysis was to compare overall survival and disease-specific survival between Hispanic (all races) and non-Hispanic HNC patients in the United States. Another purpose of this study was to identify prognostic factors that might affect the survival of HNC patients. Although the multidisciplinary care of combined surgery and radiotherapy along with rehabilitation has gained significant improvement in patient's prognosis and survival time, its accessibility and reliability for minor races/ethnicities such as Hispanic patients remain unknown due to the lack of relevant clinical evidences. Given this uncertainty, it is necessary to compare the survival between Hispanic HNC patients and the major race, non-Hispanic White.

It was hypothesized that the survival for Hispanic HNC patients would vary significantly from the general population in the U.S. given the socioeconomic and other demographic disparities in Hispanic population. In the present study, survival data from the SEER database was utilized and it was demonstrated that the median survival and the 5 -year survival rate for Hispanic HNC patients were significantly lower than those for non-Hispanic White and for nonHispanic other races. Furthermore, the trends for survival data for each cancer group were also collected and analyzed. In respect to the oral cavity and the pharyngeal cancers, Hispanic patients demonstrated poorer OS as well as disease-specific survival than those for non-Hispanic White patients. These results coincided with our hypothesis and were consistent with literature which states that racial disparities result in inferior OS in patients with HNC (Shin \& Truong, 2015).

However, one surprising finding was that, for laryngeal cancer, Hispanic patients showed better

OS than that for non-Hispanic White patients, even though this superiority disappeared when comparing disease-specific survival between two racial groups. This illustrates that the Hispanic population outperforms the non-Hispanic White population in some specific prognostic factors such as tobacco use, especially smoking.

A multivariate analysis was implemented to identify the impact of important prognostic factors on both OS and disease-specific survival in Hispanic and non-Hispanic patients. It was found that Hispanic race is associated with inferior OS and disease-specific survival in patients with HNC. Hispanic patients were associated with relatively lower prevalence of being married, lower insurance cover rate, advanced tumor size, lower education, and higher unemployment rate, all of which were predictors of inferior OS and disease-specific survival in both univariate analyses and multivariable models. These socioeconomic and demographic factors account for the poor survival in Hispanic HNC patients. Besides, elder age, non-Hispanic Black race, advanced tumor grade, and higher prevalence of smoking were independent prognostic factors for worse survival rate, which have been published by previous study (Mehanna et al., 2010b).

Apart from the prognostic factors that have been discovered, the interaction between race and age has been proved to have significant impact on two survival outcomes (OS and diseasespecific survival). It can be seen that among patients with 40-59 and 60-79 years of age, Hispanic race had significant poorer survival versus non-Hispanic White and non-Hispanic other races. However, this significant effect of interaction no longer existed in the youngest group (20-39 years of age) and in the oldest group (80+ years of age). For the youngest group, having a better
prognosis of HNC was conjectured based on the finding. On the other hand, the fact that elderly Hispanic patients were less likely to be enrolled in the study compared with non-Hispanic White and non-Hispanic other races accounted for the absence of significant interaction in the oldest group.

The results of propensity score matching based on complete sample and samples for each cancer site further illustrate that Hispanic HNC patients have poorer OS and disease-specific survival when comparing with non-Hispanic White or non-Hispanic other races. Hispanic patients presented lower hazard ratio in OS compared with non-Hispanic White for laryngeal cancer and higher hazard ratio in disease-specific survival for pharyngeal cancer. However, unlike the hazard ratio $P$-values in multivariate analyses before propensity score matching, there were many $P$ values calculated based on matched sample being not statistically significant as sample size of each racial sub-group decreased. From Kaplan-Meier curves generated from matched sample, it can be seen that the survival of Hispanic patients fell behind that of non-Hispanic White patients during the earlier phase of prognosis while it caught up during the later phase. This "catching-up" property of survival in Hispanic patients can be found in both OS and disease-specific survival Kaplan-Meier curves. A pathological reason for the presence of the "catching-up" property needs to be explored in future studies.

### 5.1 Implications

The results of this study illustrate that OS and disease-specific survival in Hispanic HNC patients were inferior to those in the general population in the U.S. The prognostic factors that
resulted in higher hazard ratio in Hispanic patients' survival reflect a variety of disparities in socioeconomic status and accessibility to public health well being among Hispanic population. These results also support the revision of standard treatment for HNC in Hispanic population and the improvement of public health welfare such as insurance coverage for Hispanic population in the U.S. These findings also identify the impact of particular behavior such as smoking on the prognosis of HNC, which can help with policy making in public health.

### 5.2 Strengths and Limitations

This study included data from a population-based cancer registry, which can be generalized to the entire U.S. population. Furthermore, the implementation of propensity score matching which allows for balanced baseline covariates between subgroups censored some drawbacks of a retrospective study and made the direct comparison of the effect of race on survival outcomes feasible. Despite the strengths, there are also some limitations and defects to the study. For example, there are still some measured baseline covariates that did not achieve a satisfactory balance between subgroups in the matched sample. This would make imprecision estimates for the impact of race on survival outcomes. However, the results of this analysis can still be considered valid because most covariates with levels were balanced perfectly. In addition, inadequate tumor grade information made it impossible to use a traditional TNM stage in the analysis. A combined SEER summary stage was used instead. Furthermore, insurance coverage was taken out from the multivariable models due to limited insurance coverage information
before 2007. These two are both important prognostic factors which could affect survival outcomes significantly.

### 5.3 Recommendations

These results can provide information on future studies regarding the survival of Hispanic HNC patients in the U.S. and the impact of prognostic factors. The implementation of prospective randomized studies is recommended to further validate the results of this study and to investigate more precise estimates for the impact of race on OS and disease-specific survival of HNC patients. In terms of the utilization of propensity score methods, a dataset with more comprehensive information at all longitudinal points could be used to further eliminate confounding from analysis. Additionally, more interaction effects between race and other baseline covariates such as gender, marital status, and year of diagnosis could be included in multivariate analysis in future studies to explore the effect of race on survival outcomes at different levels.

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## Appendix: TABLES AND FIGURES

Table 1 Patient and Treatment Characteristics

| Demographic or Prognostic Characteristics | Level | $\begin{gathered} \mathrm{N}=113885 \\ \mathrm{~N}(\%) \end{gathered}$ |
| :---: | :---: | :---: |
| Year of diagnosis | 2000-2006 | 53691 (47.1) |
|  | 2007-2013 | 60194 (52.9) |
| Cancer site | Larynx | 31720 (27.9) |
|  | Oral cavity | 49850 (43.8) |
|  | Pharynx | 32315 (28.4) |
| Race | Hispanic (All Races) | 8132 (7.1) |
|  | Non-Hispanic Black | 12669 (11.1) |
|  | Non-Hispanic Others | 7089 (6.2) |
|  | Non-Hispanic White | 85995 (75.5) |
| Origin recode NHIA (Hispanic, Non-Hisp) | Non-Spanish-Hispanic- | 105753 (92.9) |
|  | Latino |  |
|  | Spanish-Hispanic-Latino | 8132 (7.1) |
| Sex | Female | 27723 (24.3) |
|  | Male | 86162 (75.7) |
| Age | 20-39 years | 3092 (2.7) |
|  | 40-59 years | 47943 (42.1) |
|  | 60-79 years | 52617 (46.2) |
|  | 80+ years | 10233 (9.0) |
| Tumor grade | Moderately | 45915 (40.3) |
|  | differentiated; Grade II |  |
|  | Poorly differentiated; | 28173 (24.7) |
|  | Grade III |  |
|  | Undifferentiated; anaplastic; Grade IV | 1441 (1.3) |
|  | Unknown | 23626 (20.7) |
|  | Well differentiated; | 14730 (12.9) |
|  | Grade I |  |
| Marital status at diagnosis | Married | 58812 (51.6) |
|  | Other | 55073 (48.4) |


| Demographic or Prognostic Characteristics | Level | $\begin{gathered} \mathrm{N}=113885 \\ \mathrm{~N}(\%) \\ \hline \end{gathered}$ |
| :---: | :---: | :---: |
| Insurance status | Any Medicaid | 9002 (15.0) |
|  | Insurance status | 2688 (4.5) |
|  | unknown |  |
|  | Insured | 35700 (59.3) |
|  | Insured/No specifics | 9620 (16.0) |
|  | Uninsured | 3184 (5.3) |
|  | Missing | 53691 |
| Combined SEER summary stage | Distant | 18885 (16.6) |
|  | Localized | 41940 (36.8) |
|  | Regional | 48222 (42.3) |
|  | Unknown | 4838 (4.2) |
| Cancer-directed surgery accepted | No | 57882 (50.8) |
|  | Unknown | 1194 (1.0) |
|  | Yes | 54809 (48.1) |
| CS tumor size in cm | $>=0,<=2$ | 13440 (11.8) |
|  | $>2,<=3$ | 13852 (12.2) |
|  | $>3,<=4$ | 15097 (13.3) |
|  | $>4,<=99$ | 10672 (9.4) |
|  | Unknown | 60824 (53.4) |
| \% Ever Smoker (males age 18+) | $>=29,<=45$ | 28500 (25.0) |
|  | $>45,<=51$ | 29320 (25.7) |
|  | $>51,<=56$ | 28012 (24.6) |
|  | $>56,<=73$ | 28040 (24.6) |
|  | Unknown | 13 (0.0) |
| \% < High school education | $>=3,<=16$ | 28695 (25.2) |
|  | $>16,<=19$ | 28500 (25.0) |
|  | $>19,<=27$ | 28810 (25.3) |
|  | $>27,<=51$ | 27867 (24.5) |
|  | Unknown | 13 (0.0) |
| \% Moved from outside US | $>=0,<=2$ | 28698 (25.2) |
|  | $>2,<=3$ | 28399 (24.9) |
|  | $>3,<=5$ | 29194 (25.6) |
|  | $>5,<=10$ | 27581 (24.2) |
|  | Unknown | 13 (0.0) |
| \% Unemployed | $>=1,<=5$ | 26621 (23.4) |
|  | $>5,<=6$ | 33112 (29.1) |
|  | $>6,<=9$ | 32733 (28.7) |
|  | $>9,<=18$ | 21406 (18.8) |
|  | Unknown | 13 (0.0) |


| Demographic or Prognostic <br> Characteristics | Level | $\mathbf{N}=\mathbf{1 1 3 8 8 5}$ <br> $\mathbf{N}(\%)$ |
| :--- | :--- | ---: |
| \% Urban | $>=0,<=76$ | $28578(25.1)$ |
|  | $>76,<=95$ | $29757(26.1)$ |
|  | $>95,<=99$ | $27285(24.0)$ |
|  | $>99,<=100$ | $28252(24.8)$ |
|  | Unknown | $13(0.0)$ |

Table 2 Univariate Association with Race

| Demographic or Prognostic Characteristic | Level | Hispanic <br> (All Races) $\begin{gathered} \mathrm{N}=\mathbf{8 1 3 2} \\ \mathrm{N}(\%) \end{gathered}$ | Non-Hispanic <br> Black $\mathbf{N}=12669$ $\mathbf{N}$ (\%) | Non-Hispanic <br> Others $\mathbf{N}=7089$ N(\%) | Non-Hispanic <br> White $\mathbf{N}=\mathbf{8 5 9 9 5}$ $\mathbf{N}(\%)$ | Parametric P-value* |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Age | 20-39 years | 412 (5.07) | 296 (2.34) | 466 (6.57) | 1918 (2.23) | <. 001 |
|  | 40-59 years | 3411 (41.95) | 6444 (50.86) | 2881 (40.64) | 35207 (40.94) |  |
|  | 60-79 years | 3656 (44.96) | 5368 (42.37) | 3017 (42.56) | 40576 (47.18) |  |
|  | $80+$ years | 653 (8.03) | 561 (4.43) | 725 (10.23) | 8294 (9.64) |  |
| Sex | Female | 1799 (22.12) | 2901 (22.9) | 2008 (28.33) | 21015 (24.44) | <. 001 |
|  | Male | 6333 (77.88) | 9768 (77.1) | 5081 (71.67) | 64980 (75.56) |  |
| Marital status at | Married | 4188 (51.5) | 4040 (31.89) | 4211 (59.4) | 46373 (53.93) | <. 001 |
| diagnosis | Other | 3944 (48.5) | 8629 (68.11) | 2878 (40.6) | 39622 (46.07) |  |
| Insurance status | Any | 1147 (24.95) | 1896 (29.96) | 751 (18.75) | 5208 (11.51) | <. 001 |
|  | Medicaid unknown | 190 (4.13) | 177 (2.8) | 482 (12.03) | 1839 (4.06) |  |
|  | Insured | 2206 (47.98) | 2621 (41.42) | 2103 (52.5) | 28770 (63.56) |  |
|  | Insured/No specifics | 694 (15.09) | 1059 (16.74) | 513 (12.81) | 7354 (16.25) |  |
|  | Uninsured | 361 (7.85) | 575 (9.09) | 157 (3.92) | 2091 (4.62) |  |
| Year of | 2000-2006 | 3534 (43.46) | 6341 (50.05) | 3083 (43.49) | 40733 (47.37) | <. 001 |
|  | 2007-2013 | 4598 (56.54) | 6328 (49.95) | 4006 (56.51) | 45262 (52.63) |  |
| Cancer-directed | No | 4123 (50.7) | 8102 (63.95) | 3697 (52.15) | 41960 (48.79) | <. 001 |
| surgery | Unknown | 71 (0.87) | 146 (1.15) | 74 (1.04) | 903 (1.05) |  |
| accepted | Yes | 3938 (48.43) | 4421 (34.9) | 3318 (46.8) | 43132 (50.16) |  |
| Cancer site | Larynx | 2461 (30.26) | 4850 (38.28) | 1272 (17.94) | 23137 (26.91) | <. 001 |
|  | Oral cavity | 3335 (41.01) | 3822 (30.17) | 3196 (45.08) | 39497 (45.93) |  |
|  | Pharynx | 2336 (28.73) | 3997 (31.55) | 2621 (36.97) | 23361 (27.17) |  |
| Tumor grade | Grade II | 3257 (40.05) | 5768 (45.53) | 2503 (35.31) | 34387 (39.99) | <. 001 |
|  | Grade III | 2120 (26.07) | 3097 (24.45) | 1791 (25.26) | 21165 (24.61) |  |
|  | Grade IV | 92 (1.13) | 151 (1.19) | 364 (5.13) | 834 (0.97) |  |
|  | Unknown | 1521 (18.7) | 2610 (20.6) | 1514 (21.36) | 17981 (20.91) |  |
|  | Grade I | 1142 (14.04) | 1043 (8.23) | 917 (12.94) | 11628 (13.52) |  |


| Demographic or Prognostic Characteristic | Level | Hispanic <br> (All Races) $\mathrm{N}=\mathbf{8 1 3 2}$ <br> (\%) | Non-Hispanic <br> Black N=12669 <br> $\mathbf{N}$ (\%) | Non-Hispanic <br> Others $\mathbf{N}=\mathbf{7 0 8 9}$ <br> $\mathbf{N}(\%)$ | Non-Hispanic <br> White $\mathbf{N}=\mathbf{8 5 9 9 5}$ <br> N(\%) | Parametric P-value* |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| CS tumor size | $>=0,<=2$ | 834 (10.26) | 722 (5.7) | 875 (12.34) | 11009 (12.8) | <. 001 |
| in cm (quartile) | $>2,<=3$ | 995 (12.24) | 1173 (9.26) | 877 (12.37) | 10807 (12.57) |  |
|  | $>3,<=4$ | 1140 (14.02) | 1682 (13.28) | 930 (13.12) | 11345 (13.19) |  |
|  | $>4,<=99$ | 953 (11.72) | 1722 (13.59) | 702 (9.9) | 7295 (8.48) |  |
|  | Unknown | 4210 (51.77) | 7370 (58.17) | 3705 (52.26) | 45539 (52.96) |  |
| Combined | Distant | 1568 (19.28) | 3293 (25.99) | 1342 (18.93) | 12682 (14.75) | <. 001 |
| SEER summary | Localized | 2847 (35.01) | 3405 (26.88) | 2510 (35.41) | 33178 (38.58) |  |
| stage | Regional | 3373 (41.48) | 5554 (43.84) | 2749 (38.78) | 36546 (42.5) |  |
|  | Unknown | 344 (4.23) | 417 (3.29) | 488 (6.88) | 3589 (4.17) |  |
| \% < High | $>=3,<=16$ | 987 (12.14) | 1450 (11.45) | 2285 (32.23) | 23973 (27.88) | <. 001 |
| school | $>16,<=19$ | 1780 (21.89) | 2788 (22.01) | 2034 (28.69) | 21898 (25.46) |  |
| education | $>19,<=27$ | 2036 (25.04) | 4957 (39.13) | 1085 (15.31) | 20732 (24.11) |  |
| (quartile) | $>27,<=51$ | 3329 (40.94) | 3474 (27.42) | 1681 (23.71) | 19383 (22.54) |  |
|  | Unknown | 0 (0) | 0 (0) | 4 (0.06) | 9 (0.01) |  |
| \% Unemployed | $>=1,<=5$ | 850 (10.45) | 1176 (9.28) | 1260 (17.77) | 23335 (27.14) | <. 001 |
| (quartile) | $>5,<=6$ | 2120 (26.07) | 2894 (22.84) | 2186 (30.84) | 25912 (30.13) |  |
|  | $>6,<=9$ | 3230 (39.72) | 4076 (32.17) | 2748 (38.76) | 22679 (26.37) |  |
|  | $>9,<=18$ | 1932 (23.76) | 4523 (35.7) | 891 (12.57) | 14060 (16.35) |  |
|  | Unknown | 0 (0) | 0 (0) | 4 (0.06) | 9 (0.01) |  |
| \% Urban | $>=0,<=76$ | 513 (6.31) | 2526 (19.94) | 632 (8.92) | 24907 (28.96) | <. 001 |
| (quartile) | $>76,<=95$ | 2335 (28.71) | 2656 (20.96) | 1280 (18.06) | 23486 (27.31) |  |
|  | $>95,<=99$ | 1926 (23.68) | 2490 (19.65) | 2473 (34.89) | 20396 (23.72) |  |
|  | $>99,<=100$ | 3358 (41.29) | 4997 (39.44) | 2700 (38.09) | 17197 (20) |  |
|  | Unknown | 0 (0) | 0 (0) | 4 (0.06) | 9 (0.01) |  |
| \% Moved from | $>=0,<=2$ | 547 (6.73) | 3303 (26.07) | 376 (5.3) | 24472 (28.46) | <. 001 |
| outside US | $>2,<=3$ | 1498 (18.42) | 3579 (28.25) | 1049 (14.8) | 22273 (25.9) |  |
| (quartile) | $>3,<=5$ | 2285 (28.1) | 2015 (15.9) | 2509 (35.39) | 22385 (26.03) |  |
|  | $>5,<=10$ | 3802 (46.75) | 3772 (29.77) | 3151 (44.45) | 16856 (19.6) |  |
|  | Unknown | 0 (0) | 0 (0) | 4 (0.06) | 9 (0.01) |  |


| Demographic or Prognostic Characteristic | Level | Hispanic <br> (All Races) $\mathrm{N}=\mathbf{8 1 3 2}$ <br> N(\%) | Non-Hispanic <br> Black N=12669 <br> $\mathbf{N}$ (\%) | Non-Hispanic <br> Others $\mathbf{N}=7089$ <br> $\mathbf{N}$ (\%) | Non-Hispanic <br> White $\mathbf{N}=85995$ <br> $\mathbf{N}(\%)$ | Parametric <br> P-value* |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| \% Ever Smoker | $>=29,<=45$ | 3876 (47.66) | 3050 (24.07) | 2888 (40.74) | 18686 (21.73) | <. 001 |
| (males age 18+) | $>45,<=51$ | 2607 (32.06) | 2632 (20.78) | 2062 (29.09) | 22019 (25.6) |  |
| (quartile) | $>51,<=56$ | 1266 (15.57) | 3262 (25.75) | 1736 (24.49) | 21748 (25.29) |  |
|  | $>56,<=73$ | 383 (4.71) | 3725 (29.4) | 399 (5.63) | 23533 (27.37) |  |
|  | Unknown | 0 (0) | 0 (0) | 4 (0.06) | 9 (0.01) |  |

Table 3.1 Univariate Association with Overall Survival

| Covariate | Level | N | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio (95\% <br> CI) | HR P- <br> value | $\begin{aligned} & \text { Log- } \\ & \text { rank } P \text { - } \\ & \text { value } \end{aligned}$ |
| Age | 40-59 years | 47943 | 1.72 (1.60-1.85) | <. 001 | $<.001$ |
|  | 60-79 years | 52617 | 2.65 (2.47-2.84) | <. 001 |  |
|  | 80+ years | 10233 | 5.14 (4.77-5.53) | $<.001$ |  |
|  | 20-39 years | 3092 | - | - |  |
| Race | Hispanic (All Races) | 8132 | 1.07 (1.03-1.10) | <. 001 | $<.001$ |
|  | Non-Hispanic Black | 12669 | 1.63 (1.59-1.67) | <. 001 |  |
|  | Non-Hispanic Others | 7089 | 0.87 (0.84-0.91) | <. 001 |  |
|  | Non-Hispanic White | 85995 | - | - |  |
| Origin recode NHIA <br> (Hispanic, Non- | Non-Spanish- <br> Hispanic-Latino | 105753 | 0.99 (0.96-1.02) | 0.592 | 0.591 |
| Hisp) | Spanish-Hispanic- <br> Latino | 8132 | - | - |  |
| Sex | Female | 27723 | 1.10 (1.08-1.12) | <. 001 | <. 001 |
|  | Male | 86162 | - | - |  |
| Marital status at |  | 58812 | 0.60 (0.59-0.61) | <. 001 | <. 001 |
| diagnosis | Other | 55073 | - | - |  |
| Insurance status | Any Medicaid | 9002 | 1.28 (1.20-1.37) | <. 001 | $<.001$ |
|  | Insurance status unknown | 2688 | 0.78 (0.72-0.86) | <. 001 |  |
|  | Insured | 35700 | 0.62 (0.58-0.66) | <. 001 |  |
|  | Insured/No specifics | 9620 | 0.90 (0.85-0.96) | 0.002 |  |
|  | Uninsured | 3184 | - | - |  |
| Year of diagnosis | 2000-2006 | 53691 | 1.14 (1.11-1.16) | <. 001 | <. 001 |
|  | 2007-2013 | 60194 | - | - |  |
| Cancer-directed | No | 57882 | 1.74 (1.71-1.77) | <. 001 | <. 001 |
|  | Unknown | 1194 | 1.75 (1.62-1.89) | <. 001 |  |
|  | Yes | 54809 | - | - |  |


| Cancer site | Larynx | 31720 | 1.00 (0.98-1.02) | 0.912 | <.001 |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  | Oral cavity | 49850 | 0.95 (0.93-0.97) | <. 001 |  |
|  | Pharynx | 32315 | - | - |  |
| Tumor grade | Moderately | 45915 | 1.37 (1.34-1.41) | <. 001 | $<.001$ |
|  | differentiated; Grade II |  |  |  |  |
|  |  | 28173 | 1.35 (1.31-1.39) | <. 001 |  |
|  | Grade III |  |  |  |  |
|  | Undifferentiated; anaplastic; Grade IV | 1441 | 1.02 (0.94-1.11) | 0.583 |  |
|  | Unknown | 23626 | 1.27 (1.23-1.31) | <. 001 |  |
|  |  | 14730 | - | - |  |
|  | Grade I |  |  |  |  |
| CS tumor size in cm (quartile) | $>2,<=3$ | 13852 | 1.64 (1.57-1.72) | <. 001 | $<.001$ |
|  | $>3,<=4$ | 15097 | 2.16 (2.07-2.25) | <. 001 |  |
|  | $>4,<=99$ | 10672 | 3.46 (3.31-3.61) | <. 001 |  |
|  | Unknown | 60824 | 2.11 (2.03-2.19) | <. 001 |  |
|  | $>=0,<=2$ | 13440 | - | - |  |
| Combined SEER summary stage | Distant | 18885 | 1.59 (1.53-1.66) | <. 001 | $<.001$ |
|  | Localized | 41940 | 0.55 (0.53-0.57) | <. 001 |  |
|  | Regional | 48222 | 0.87 (0.84-0.91) | <. 001 |  |
|  | Unknown | 4838 | - | - |  |
| \% < High school education (quartile) | $>16,<=19$ | 28500 | 1.12 (1.09-1.14) | <. 001 | $<.001$ |
|  | $>19,<=27$ | 28810 | 1.24 (1.21-1.27) | <. 001 |  |
|  | $>27,<=51$ | 27867 | 1.26 (1.23-1.29) | <. 001 |  |
|  | Unknown | 13 | 0.88 (0.37-2.11) | 0.776 |  |
|  | $>=3,<=16$ | 28695 | - | - |  |
| \% Unemployed (quartile) | $>5,<=6$ | 33112 | 1.07 (1.05-1.10) | <. 001 | $<.001$ |
|  | $>6,<=9$ | 32733 | 1.18 (1.15-1.21) | <. 001 |  |
|  | $>9,<=18$ | 21406 | 1.32 (1.28-1.35) | <. 001 |  |
|  | Unknown | 13 | 0.86 (0.36-2.07) | 0.741 |  |
|  | $>=1,<=5$ | 26621 | - | - |  |
| \% Urban (quartile) | $>76,<=95$ | 29757 | 0.97 (0.94-0.99) | 0.006 | <. 001 |
|  | $>95,<=99$ | 27285 | 0.93 (0.91-0.95) | <. 001 |  |
|  | >99, <=100 | 28252 | 1.03 (1.01-1.06) | 0.006 |  |
|  | Unknown | 13 | 0.75 (0.31-1.81) | 0.523 |  |


|  | $>=0,<=76$ | 28578 | - | - |  |
| :--- | :--- | :---: | :---: | :---: | :---: |
| \% Moved from | $>2,<=3$ | 28399 | $0.99(0.96-1.01)$ | 0.219 | $<.001$ |
| outside US | $>3,<=5$ | 29194 | $0.90(0.88-0.92)$ | $<.001$ |  |
| (quartile) | $>5,<=10$ | 27581 | $0.96(0.94-0.99)$ | $\mathbf{0 . 0 0 2}$ |  |
|  | Unknown | 13 | $0.74(0.31-1.77)$ | 0.492 |  |
|  | $>=0,<=2$ | 28698 | - | - |  |
| \% Ever Smoker | $>45,<=51$ |  |  |  |  |
| (males age 18+) | $>51,<=56$ | 29320 | $1.00(0.97-1.02)$ | 0.770 | $<.001$ |
| (quartile) | $>56,<=73$ | 28012 | $1.07(1.04-1.09)$ | $<.001$ |  |
|  | Unknown | 28040 | $1.11(1.08-1.13)$ | $<.001$ |  |
|  | $>=29,<=45$ | 13 | $0.80(0.33-1.91)$ | 0.611 |  |

Table 3.2 Univariate Association with Overall Survival for Laryngeal Cancer

| Covariate | Level | N | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio (95\% CI) | HR Pvalue | Log-rank <br> $P$-value |
| Age | 40-59 years | 11370 | 2.93 (2.33-3.68) | <. 001 | <. 001 |
|  | 60-79 years | 17132 | 3.93 (3.13-4.93) | <. 001 |  |
|  | 80+ years | 2812 | 7.37 (5.85-9.28) | <. 001 |  |
|  | 20-39 years | 406 | - | - |  |
| Race | Hispanic (All Races) | 2461 | 0.93 (0.87-0.99) | 0.015 | <. 001 |
|  | Non-Hispanic Black | 4850 | 1.26 (1.21-1.31) | <. 001 |  |
|  | Non-Hispanic Others | 1272 | 0.80 (0.73-0.87) | <. 001 |  |
|  | Non-Hispanic White | 23137 | - | - |  |
| Origin recode NHIA <br> (Hispanic, Non-Hisp) | Non-Spanish-Hispanic- | 29259 | 1.11 (1.05-1.18) | <. 001 | <. 001 |
|  | Latino |  |  |  |  |
|  | Spanish-Hispanic-Latino | 2461 | - | - |  |
| Sex | Female | 6017 | 1.06 (1.02-1.10) | 0.007 | 0.006 |
|  | Male | 25703 | - | - |  |
| Marital status at diagnosis | Married | 16499 | 0.64 (0.62-0.67) | <. 001 | <. 001 |
|  | Other | 15221 | - | - |  |
| Insurance status | Any Medicaid | 2863 | 1.28 (1.14-1.45) | <. 001 | <. 001 |
|  | Insurance status unknown | 411 | 0.85 (0.69-1.04) | 0.116 |  |
|  | Insured | 8718 | 0.74 (0.66-0.83) | <. 001 |  |
|  | Insured/No specifics | 2798 | 0.99 (0.88-1.13) | 0.924 |  |
|  | Uninsured | 899 | - | - |  |
| Year of diagnosis | 2000-2006 | 16031 | 1.02 (0.98-1.05) | 0.308 | 0.304 |
|  | 2007-2013 | 15689 | - | - |  |
| Cancer-directed surgery accepted | No | 19020 | 1.33 (1.29-1.38) | <.001 | <. 001 |
|  | Unknown | 413 | 1.26 (1.11-1.44) | <. 001 |  |
|  | Yes | 12287 | - | - |  |


| Covariate | Level | Survival months |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | N | Hazard Ratio (95\% <br> CI) | HR P- <br> value | Log-rank P-value |
| Tumor grade | Moderately differentiated; Grade II | 15051 | 1.44 (1.37-1.52) | <. 001 | <. 001 |
|  | Poorly differentiated; Grade III | 5685 | 2.06 (1.95-2.19) | <. 001 |  |
|  | Undifferentiated; anaplastic; Grade IV | 146 | 1.94 (1.56-2.41) | <. 001 |  |
|  | Unknown | 6431 | 1.37 (1.29-1.45) | <. 001 |  |
|  | Well differentiated; Grade I | 4407 | - | - |  |
| CS tumor size in cm (quartile) | $>2,<=3$ | 2330 | 1.84 (1.67-2.03) | <. 001 | <. 001 |
|  | $>3,<=4$ | 2443 | 2.24 (2.04-2.46) | $<.001$ |  |
|  | $>4,<=99$ | 1572 | 2.89 (2.62-3.19) | <. 001 |  |
|  | Unknown | 23049 | 1.46 (1.35-1.57) | <. 001 |  |
|  | $>=0,<=2$ | 2326 | - | - |  |
| Combined SEER summary stage | Distant | 5740 | 1.81 (1.66-1.97) | <. 001 | <. 001 |
|  | Localized | 18147 | 0.65 (0.60-0.71) | <. 001 |  |
|  | Regional | 6692 | 1.38 (1.27-1.50) | <. 001 |  |
|  | Unknown | 1141 | - | - |  |
| \% < High school education (quartile) | $>16,<=19$ | 7660 | 1.08 (1.03-1.13) | 0.002 | <. 001 |
|  | $>19,<=27$ | 8515 | 1.12 (1.07-1.17) | <. 001 |  |
|  | $>27,<=51$ | 8437 | 1.15 (1.10-1.20) | <. 001 |  |
|  | Unknown | 3 | 2.22 (0.31-15.73) | 0.426 |  |
|  | $>=3,<=16$ | 7105 | - | - |  |
| \% Unemployed (quartile) | $>5,<=6$ | 9040 | 1.06 (1.01-1.11) | 0.014 | <. 001 |
|  | $>6,<=9$ | 9099 | 1.10 (1.05-1.15) | <. 001 |  |
|  | $>9,<=18$ | 6532 | 1.22 (1.17-1.28) | <. 001 |  |
|  | Unknown | 3 | 2.22 (0.31-15.72) | 0.426 |  |
|  | $>=1,<=5$ | 7046 | - | - |  |
| \% Urban (quartile) | $>76,<=95$ | 7892 | 1.00 (0.96-1.04) | 0.897 | 0.650 |
|  | $>95,<=99$ | 6948 | 1.01 (0.96-1.05) | 0.734 |  |
|  | >99, <=100 | 7832 | 1.03 (0.98-1.07) | 0.246 |  |
|  | Unknown | 3 | 2.05 (0.29-14.53) | 0.473 |  |
|  | $>=0,<=76$ | 9045 | - | - |  |


| Covariate | Level | N | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio (95\% CI) | HR P- <br> value | Log-rank <br> P-value |
| \% Moved from | $>2,<=3$ | 8112 | 1.03 (0.98-1.07) | 0.233 | 0.442 |
| outside US (quartile) | $>3,<=5$ | 7204 | 0.99 (0.94-1.03) | 0.534 |  |
|  | $>5,<=10$ | 7073 | 1.01 (0.97-1.05) | 0.673 |  |
|  | Unknown | 3 | 2.04 (0.29-14.50) | 0.474 |  |
|  | $>=0,<=2$ | 9328 | - | - |  |
| \% Ever Smoker | $>45,<=51$ | 7549 | 0.96 (0.92-1.01) | 0.129 | 0.077 |
| (males age 18+) | $>51,<=56$ | 8140 | 1.00 (0.95-1.04) | 0.877 |  |
| (quartile) | $>56,<=73$ | 9268 | 1.03 (0.98-1.07) | 0.267 |  |
|  | Unknown | 3 | 2.03 (0.29-14.40) | 0.479 |  |
|  | $>=29,<=45$ | 6760 | - | - |  |

Table 3.3 Univariate Association with Disease-Specific Survival

| Covariate | Level | N | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio (95\% <br> CI) | HR P- <br> value | Log-rank $P$-value |
| Age | 40-59 years | 47943 | 1.44 (1.34-1.56) | <. 001 | <. 001 |
|  | 60-79 years | 52617 | 1.84 (1.70-1.99) | <. 001 |  |
|  | 80+ years | 10233 | 2.95 (2.72-3.21) | <. 001 |  |
|  | 20-39 years | 3092 | - | - |  |
| Race | Hispanic (All Races) | 8132 | 1.18 (1.13-1.22) | $<.001$ | <. 001 |
|  | Non-Hispanic Black | 12669 | 1.83 (1.78-1.88) | $<.001$ |  |
|  | Non-Hispanic Others | 7089 | 0.99 (0.95-1.04) | 0.732 |  |
|  | Non-Hispanic White | 85995 | - | - |  |
| Origin recode | Non-Spanish-Hispanic-Latino | 105753 | 0.92 (0.89-0.96) | <. 001 | <. 001 |
| NHIA <br> (Hispanic, Non- | Spanish-Hispanic-Latino | 8132 | - | - |  |
| Hisp) |  |  |  |  |  |
| Sex | Female | 27723 | 1.10 (1.08-1.13) | $<.001$ | <. 001 |
|  | Male | 86162 | - | - |  |
|  | Married | 58812 | 0.59 (0.58-0.60) | <. 001 | <. 001 |
| diagnosis | Other | 55073 | - | - |  |
| Insurance status | Any Medicaid | 9002 | 1.20 (1.12-1.29) | <. 001 | <. 001 |
|  | Insurance status unknown | 2688 | 0.69 (0.62-0.76) | $<.001$ |  |
|  | Insured | 35700 | 0.55 (0.52-0.59) | <. 001 |  |
|  | Insured/No specifics | 9620 | 0.81 (0.75-0.87) | <. 001 |  |
|  | Uninsured | 3184 | - | - |  |
| Year of | 2000-2006 | 53691 | 1.13 (1.11-1.16) | <. 001 | <. 001 |
|  | 2007-2013 | 60194 | - | - |  |
| Cancer-directed | No | 57882 | 2.02 (1.97-2.06) | $<.001$ | <. 001 |
| surgery | Unknown | 1194 | 2.04 (1.86-2.24) | <. 001 |  |
| accepted | Yes | 54809 | - | - |  |


| Covariate | Level | N | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio (95\% <br> CI) | HR Pvalue | Log-rank <br> $P$-value |
| Cancer site | Larynx | 31720 | 0.85 (0.83-0.87) | <. 001 | <. 001 |
|  | Oral cavity | 49850 | 0.85 (0.83-0.88) | $<.001$ |  |
|  | Pharynx | 32315 | - | - |  |
| Tumor grade | Moderately differentiated; <br> Grade II | 45915 | 1.63 (1.57-1.69) | <. 001 | <. 001 |
|  | Poorly differentiated; Grade III | 28173 | 1.71 (1.64-1.78) | $<.001$ |  |
|  | Undifferentiated; anaplastic; | 1441 | 1.33 (1.21-1.47) | <. 001 |  |
|  | Grade IV |  |  |  |  |
|  | Unknown | 23626 | 1.48 (1.42-1.54) | <. 001 |  |
|  | Well differentiated; Grade I | 14730 | - | - |  |
| CS tumor size | $>2,<=3$ | 13852 | 2.04 (1.93-2.17) | <. 001 | <. 001 |
| in cm (quartile) | $>3,<=4$ | 15097 | 2.95 (2.79-3.12) | <. 001 |  |
|  | $>4,<=99$ | 10672 | 5.11 (4.83-5.41) | <. 001 |  |
|  | Unknown | 60824 | 2.78 (2.65-2.93) | <. 001 |  |
|  | $>=0,<=2$ | 13440 | - | - |  |
| Combined | Distant | 18885 | 1.80 (1.71-1.90) | <. 001 | <. 001 |
| SEER summary | Localized | 41940 | 0.39 (0.37-0.41) | <. 001 |  |
| stage | Regional | 48222 | 0.91 (0.87-0.96) | $<.001$ |  |
|  | Unknown | 4838 | - | - |  |
| $\%$ < High | $>16,<=19$ | 28500 | 1.14 (1.11-1.18) | <. 001 | <. 001 |
| school | $>19,<=27$ | 28810 | 1.28 (1.24-1.32) | <. 001 |  |
| education | $>27,<=51$ | 27867 | 1.30 (1.26-1.33) | $<.001$ |  |
| (quartile) | Unknown | 13 | 1.15 (0.43-3.05) | 0.786 |  |
|  | $>=3,<=16$ | 28695 | - | - |  |
| \% Unemployed | $>5,<=6$ | 33112 | 1.10 (1.07-1.13) | <. 001 | <. 001 |
| (quartile) | $>6,<=9$ | 32733 | 1.24 (1.20-1.28) | <. 001 |  |
|  | $>9,<=18$ | 21406 | 1.37 (1.33-1.42) | <. 001 |  |
|  | Unknown | 13 | 1.13 (0.43-3.02) | 0.801 |  |
|  | $>=1,<=5$ | 26621 | - | - |  |


| Covariate |  | N | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  | Level |  | Hazard Ratio (95\% <br> CI) | HR P- <br> value | Log-rank <br> $P$-value |
| \% Urban | $>76,<=95$ | 29757 | 1.00 (0.97-1.03) | 0.971 | <. 001 |
| (quartile) | >95, <=99 | 27285 | 0.98 (0.95-1.01) | 0.228 |  |
|  | >99, <=100 | 28252 | 1.10 (1.07-1.13) | <. 001 |  |
|  | Unknown | 13 | 0.99 (0.37-2.65) | 0.991 |  |
|  | $>=0,<=76$ | 28578 | - | - |  |
| \% Moved from | $>2,<=3$ | 28399 | 1.01 (0.98-1.04) | 0.633 | <. 001 |
| outside US | $>3,<=5$ | 29194 | 0.92 (0.90-0.95) | <. 001 |  |
| (quartile) | $>5,<=10$ | 27581 | 1.02 (0.99-1.05) | 0.117 |  |
|  | Unknown | 13 | 0.96 (0.36-2.56) | 0.938 |  |
|  | $>=0,<=2$ | 28698 | - | - |  |
| \% Ever Smoker | $>45,<=51$ | 29320 | 0.99 (0.96-1.02) | 0.433 | <. 001 |
| (males age | $>51,<=56$ | 28012 | 1.05 (1.02-1.08) | 0.001 |  |
| 18+) (quartile) | $>56,<=73$ | 28040 | 1.04 (1.01-1.07) | 0.006 |  |
|  | Unknown | 13 | 0.99 (0.37-2.65) | 0.989 |  |
|  | $>=29,<=45$ | 28500 | - | - |  |

Table 4.1 Multivariate Association with Overall Survival

| Covariate | Level | Survival months |  |
| :---: | :---: | :---: | :---: |
|  |  | Hazard Ratio ( $95 \% \mathrm{CI}$ ) | HR P- <br> value |
| Age | 40-59 years | 1.57 (1.47-1.69) | <. 001 |
|  | 60-79 years | 2.65 (2.47-2.85) | <. 001 |
|  | 80+ years | 5.73 (5.32-6.17) | <. 001 |
|  | 20-39 years | - | - |
| Race | Hispanic (All Races) | 1.06 (1.03-1.10) | <. 001 |
|  | Non-Hispanic Black | 1.34 (1.30-1.37) | <. 001 |
|  | Non-Hispanic Others | 0.91 (0.88-0.95) | <. 001 |
|  | Non-Hispanic White | - | - |
| Marital status at diagnosis | Married | 0.68 (0.66-0.69) | $<.001$ |
|  | Other | - | - |
| Year of diagnosis | 2000-2006 | 1.19 (1.17-1.22) | <. 001 |
|  | 2007-2013 | - | - |
| Cancer-directed surgery accepted | No | 1.53 (1.50-1.55) | <. 001 |
|  | Unknown | 1.51 (1.39-1.63) | $<.001$ |
|  | Yes | - | - |
| Cancer site | Larynx | 1.03 (1.00-1.05) | 0.045 |
|  | Oral cavity | 1.12 (1.09-1.14) | <. 001 |
|  | Pharynx | - | - |
| Tumor grade | Moderately differentiated; Grade II | 1.19 (1.16-1.23) | <. 001 |
|  | Poorly differentiated; Grade III | 1.08 (1.05-1.11) | <. 001 |
|  | Undifferentiated; anaplastic; Grade IV | 0.86 (0.79-0.94) | $<.001$ |
|  | Unknown | 1.03 (1.00-1.07) | 0.042 |
|  | Well differentiated; Grade I | - | - |


| Covariate | Level | Survival months |  |
| :---: | :---: | :---: | :---: |
|  |  | $\begin{gathered} \text { Hazard Ratio } \\ \text { (95\% CI) } \end{gathered}$ | HR P- <br> value |
| CS tumor size in cm (quartile) | $>2,<=3$ | 1.25 (1.20-1.31) | <. 001 |
|  | $>3,<=4$ | 1.40 (1.34-1.46) | <. 001 |
|  | $>4,<=99$ | 1.95 (1.87-2.04) | <. 001 |
|  | Unknown | 1.54 (1.48-1.60) | <. 001 |
|  | $>=0,<=2$ | - | - |
| Combined SEER summary stage | Distant | 1.86 (1.78-1.95) | <. 001 |
|  | Localized | 0.67 (0.65-0.70) | <. 001 |
|  | Regional | 1.13 (1.08-1.18) | <. 001 |
|  | Unknown | - | - |
| $\%<$ High school education (quartile) | $>16,<=19$ | 1.05 (1.02-1.08) | <. 001 |
|  | $>19,<=27$ | 1.07 (1.04-1.10) | <. 001 |
|  | $>27,<=51$ | 1.12 (1.08-1.16) | <. 001 |
|  | Unknown | 1.22 (0.51-2.94) | 0.653 |
|  | $>=3,<=16$ | - | - |
| \% Unemployed (quartile) | $>5,<=6$ | 1.00 (0.98-1.03) | 0.888 |
|  | $>6,<=9$ | 1.05 (1.02-1.08) | <. 001 |
|  | $>9,<=18$ | 1.08 (1.05-1.12) | <. 001 |
|  | Unknown | - | - |
|  | $>=1,<=5$ | - | - |
| \% Urban (quartile) | $>76,<=95$ | 0.97 (0.94-1.00) | 0.055 |
|  | $>95,<=99$ | 0.97 (0.94-1.01) | 0.112 |
|  | $>99,<=100$ | 0.99 (0.95-1.02) | 0.437 |
|  | Unknown | - | - |
|  | $>=0,<=76$ | - | - |
| \% Moved from outside US (quartile) | $>2,<=3$ | 1.01 (0.98-1.04) | 0.436 |
|  | $>3,<=5$ | 0.97 (0.94-1.01) | 0.130 |
|  | $>5,<=10$ | 0.95 (0.91-0.99) | 0.014 |
|  | Unknown | - | - |
|  | $>=0,<=2$ | - | - |


| Covariate | Level | Survival months |  |
| :---: | :---: | :---: | :---: |
|  |  | Hazard Ratio (95\% CI) | HR P- <br> value |
| \% Ever Smoker (males age | $>45,<=51$ | 1.01 (0.98-1.04) | 0.465 |
| 18+) (quartile) | $>51,<=56$ | 1.06 (1.03-1.10) | <. 001 |
|  | $>56,<=73$ | 1.06 (1.02-1.10) | 0.001 |
|  | Unknown | - | - |
|  | $>=29,<=45$ | - | - |

* Number of observations in the original data set $=113885$. Number of observations used $=113885$.
** Backward selection with an alpha level of removal of 0.05 was used. The following variables were removed from the model: Origin recode NHIA (Hispanic, Non-Hisp).

Table 4.2 Multivariate Association with Overall Survival for Laryngeal Cancer Survival months

| Covariate | Level | $\begin{gathered} \text { Hazard Ratio } \\ (\mathbf{9 5 \%} \% \mathbf{C I}) \\ \hline \end{gathered}$ | HR P- <br> value |
| :---: | :---: | :---: | :---: |
| Age | 40-59 years | 2.72 (2.17-3.42) | <. 001 |
|  | 60-79 years | 4.27 (3.40-5.36) | <. 001 |
|  | 80+ years | 9.20 (7.30-11.59) | <. 001 |
|  | 20-39 years | - | - |
| Race | Hispanic (All Races) | 0.91 (0.85-0.97) | 0.004 |
|  | Non-Hispanic Black | 1.08 (1.03-1.12) | 0.001 |
|  | Non-Hispanic Others | 0.79 (0.72-0.86) | <. 001 |
|  | Non-Hispanic White | - | - |
| Sex | Female | 0.92 (0.89-0.96) | <. 001 |
|  | Male | - | - |
| Marital status at diagnosis | Married | 0.71 (0.69-0.73) | <. 001 |
|  | Other | - | - |
| Year of diagnosis | 2000-2006 | 1.12 (1.08-1.16) | <. 001 |
|  | 2007-2013 | - | - |
| Cancer-directed surgery accepted | No | 1.46 (1.41-1.51) | <. 001 |
|  | Unknown | 1.31 (1.15-1.50) | <. 001 |
|  | Yes | - | - |
| Tumor grade | Moderately differentiated; Grade II | 1.21 (1.15-1.28) | <. 001 |
|  | Poorly differentiated; Grade III | 1.44 (1.36-1.53) | <. 001 |
|  | Undifferentiated; anaplastic; Grade IV | 1.38 (1.11-1.72) | 0.004 |
|  | Unknown | 1.21 (1.14-1.28) | <. 001 |
|  | Well differentiated; Grade I | - | - |


| Covariate | Level | Survival months |  |
| :---: | :---: | :---: | :---: |
|  |  | $\begin{gathered} \text { Hazard Ratio } \\ \text { (95\% CI) } \end{gathered}$ | HR P- <br> value |
| CS tumor size in cm (quartile) | $>2,<=3$ | 1.21 (1.09-1.33) | <.001 |
|  | $>3,<=4$ | 1.28 (1.16-1.40) | <. 001 |
|  | $>4,<=99$ | 1.54 (1.39-1.71) | <. 001 |
|  | Unknown | 1.24 (1.15-1.34) | <. 001 |
|  | $>=0,<=2$ | - | - |
| Combined SEER summary stage | Distant | 2.13 (1.95-2.33) | <. 001 |
|  | Localized | 0.74 (0.68-0.80) | <. 001 |
|  | Regional | 1.57 (1.44-1.72) | <. 001 |
|  | Unknown | - | - |
| $\%<$ High school education (quartile) | $>16,<=19$ | 1.00 (0.94-1.05) | 0.863 |
|  | $>19,<=27$ | 1.03 (0.98-1.09) | 0.217 |
|  | $>27,<=51$ | 1.07 (1.01-1.13) | 0.030 |
|  | Unknown | 1.63 (0.23-11.61) | 0.624 |
|  | $>=3,<=16$ | - | - |
| \% Unemployed (quartile) | $>5,<=6$ | 1.03 (0.98-1.08) | 0.315 |
|  | $>6,<=9$ | 1.05 (1.00-1.11) | 0.063 |
|  | $>9,<=18$ | 1.11 (1.04-1.17) | <. 001 |
|  | Unknown | - | - |
|  | $>=1,<=5$ | - | - |
| \% Urban (quartile) | $>76,<=95$ | 1.01 (0.96-1.06) | 0.785 |
|  | $>95,<=99$ | 1.01 (0.95-1.07) | 0.727 |
|  | $>99,<=100$ | 0.94 (0.89-1.00) | 0.035 |
|  | Unknown | - | - |
|  | $>=0,<=76$ | - | - |
| \% Ever Smoker (males age | $>45,<=51$ | 0.95 (0.90-1.00) | 0.058 |
| 18+) (quartile) | $>51,<=56$ | 0.99 (0.94-1.05) | 0.781 |
|  | $>56,<=73$ | 1.01 (0.95-1.06) | 0.815 |
|  | Unknown | - | - |
|  | $>=29,<=45$ | - | - |


|  |  | Survival months |  |
| :---: | :---: | :---: | :---: |
| Covariate | Level | $\begin{gathered} \text {---- } \\ \text { Hazard Ratio } \\ (95 \% \text { CI) } \\ \hline \end{gathered}$ | HR P- <br> value |
| * Number of observations in the original data set $=31720$. Number of observations used $=31720$. <br> ** Backward selection with an alpha level of removal of 0.05 was used. The following variables were removed from the model: Origin recode NHIA (Hispanic, Non-Hisp). |  |  |  |

Table 4.3 Multivariate Association with Disease-Specific Survival

| Covariate | Level | Survival months |  |
| :---: | :---: | :---: | :---: |
|  |  | Hazard Ratio (95\% CI) | HR Pvalue |
| Age | 40-59 years | 1.30 (1.20-1.41) | <. 001 |
|  | 60-79 years | 1.87 (1.73-2.02) | <. 001 |
|  | 80+ years | 3.48 (3.20-3.78) | <. 001 |
|  | 20-39 years | - | - |
| Race | Hispanic (All Races) | 1.12 (1.08-1.17) | <. 001 |
|  | Non-Hispanic Black | 1.39 (1.34-1.43) | <. 001 |
|  | Non-Hispanic Others | 1.00 (0.95-1.04) | 0.866 |
|  | Non-Hispanic White | - | - |
| Sex | Female | 1.06 (1.03-1.08) | <. 001 |
|  | Male | - | - |
| Marital status at diagnosis | Married | 0.68 (0.67-0.70) | <. 001 |
|  | Other | - | - |
| Year of diagnosis | 2000-2006 | 1.20 (1.17-1.23) | <. 001 |
|  | 2007-2013 | - | - |
| Cancer-directed surgery accepted | No | 1.65 (1.62-1.69) | <. 001 |
|  | Unknown | 1.68 (1.53-1.84) | <. 001 |
|  | Yes | - | - |
| Cancer site | Larynx | 1.02 (0.99-1.05) | 0.150 |
|  | Oral cavity | 1.14 (1.11-1.17) | <. 001 |
|  | Pharynx | - | - |


| Covariate | Level | Survival months |  |
| :---: | :---: | :---: | :---: |
|  |  | Hazard Ratio （95\％CI） | HR P－ <br> value |
| Tumor grade | Moderately differentiated；Grade II | 1.26 （1．21－1．31） | ＜． 001 |
|  | Poorly differentiated； Grade III | 1.14 （1．09－1．19） | $<.001$ |
|  | Undifferentiated； anaplastic；Grade IV | 0.86 （0．77－0．95） | 0.003 |
|  | Unknown | 1.05 （1．01－1．10） | 0.018 |
|  | Well differentiated； Grade I | － | － |
| CS tumor size in cm （quartile） | $>2,<=3$ | 1.38 （1．30－1．46） | ＜． 001 |
|  | $>3,<=4$ | 1.60 （1．51－1．70） | ＜． 001 |
|  | $>4,<=99$ | 2.33 （2．20－2．47） | ＜． 001 |
|  | Unknown | 1.86 （1．76－1．95） | ＜． 001 |
|  | $>=0,<=2$ | － | － |
| Combined SEER summary stage | Distant | 2.09 （1．98－2．20） | ＜． 001 |
|  | Localized | 0.51 （0．49－0．54） | ＜． 001 |
|  | Regional | 1.18 （1．12－1．24） | ＜． 001 |
|  | Unknown | － | － |
| \％＜High school education （quartile） | $>16,<=19$ | 1.07 （1．03－1．10） | $<.001$ |
|  | $>19,<=27$ | 1.08 （1．04－1．12） | ＜． 001 |
|  | $>27,<=51$ | 1.14 （1．09－1．18） | ＜． 001 |
|  | Unknown | 1.57 （0．59－4．19） | 0.368 |
|  | $>=3,<=16$ | － | － |
| \％Unemployed（quartile） | $>5,<=6$ | 1.01 （0．98－1．05） | 0.390 |
|  | $>6,<=9$ | 1.07 （1．04－1．11） | ＜． 001 |
|  | $>9,<=18$ | 1.08 （1．04－1．13） | ＜． 001 |
|  | Unknown | － | － |
|  | $>=1,<=5$ | － | － |


| Covariate | Level | Survival months |  |
| :---: | :---: | :---: | :---: |
|  |  | Hazard Ratio (95\% CI) | HR P- <br> value |
| \% Moved from outside US (quartile) | >2, <=3 | 1.00 (0.97-1.03) | 0.966 |
|  | $>3,<=5$ | 0.95 (0.92-0.99) | 0.007 |
|  | $>5,<=10$ | 0.95 (0.91-0.99) | 0.018 |
|  | Unknown | - | - |
|  | $>=0,<=2$ | - | - |
| \% Ever Smoker (males age 18+) (quartile) | $>45,<=51$ | 1.02 (0.98-1.05) | 0.288 |
|  | $>51,<=56$ | 1.07 (1.02-1.11) | 0.002 |
|  | $>56,<=73$ | 1.03 (0.99-1.08) | 0.167 |
|  | Unknown | - | - |
|  | $>=29,<=45$ | - | - |

* Number of observations in the original data set $=113885$. Number of observations used $=113885$.
** Backward selection with an alpha level of removal of 0.05 was used. The following variables were removed from the model: Origin recode NHIA (Hispanic, Non-Hisp).

Table 5.1 Interaction Effect Between Race and Age on Overall Survival

| Covariate | Level | N | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio (95\% CI) | HR P- <br> value | $\begin{gathered} \text { Type3 } \\ \text { P-value } \end{gathered}$ |
| Comparisons | Race : |  | - | - | <. 001 |
| Stratified by |  |  |  |  |  |
| Age : |  |  |  |  |  |
| 20-39 years | Non-Hispanic Black vs. Hispanic (All Races) | 296 vs. 412 | 1.17 (0.91-1.51) | 0.215 | - |
|  | Non-Hispanic Others vs. Hispanic (All Races) | 466 vs. 412 | 0.84 (0.65-1.09) | 0.202 |  |
|  | Non-Hispanic White vs. Hispanic (All Races) | 1918 vs. 412 | 0.73 (0.59-0.90) | 0.003 |  |
| 40-59 years | Non-Hispanic Black vs. Hispanic (All Races) | 6444 vs. <br> 3411 | 1.43 (1.34-1.53) | <. 001 | - |
|  | Non-Hispanic Others vs. Hispanic (All Races) | $\begin{gathered} 2881 \text { vs. } \\ 3411 \end{gathered}$ | 0.80 (0.73-0.87) | <. 001 |  |
|  | Non-Hispanic White vs. Hispanic (All Races) | $\begin{aligned} & 35207 \text { vs. } \\ & 3411 \end{aligned}$ | 0.89 (0.84-0.94) | <. 001 |  |
| 60-79 years | Non-Hispanic Black vs. Hispanic (All Races) | $\begin{gathered} 5368 \text { vs. } \\ 3656 \end{gathered}$ | 1.13 (1.07-1.20) | <. 001 | - |
|  | Non-Hispanic Others vs. Hispanic (All Races) | $\begin{gathered} 3017 \text { vs. } \\ 3656 \end{gathered}$ | 0.86 (0.80-0.93) | <. 001 |  |
|  | Non-Hispanic White vs. Hispanic (All Races) | $\begin{gathered} 40576 \text { vs. } \\ 3656 \end{gathered}$ | 0.95 (0.91-1.00) | 0.042 |  |
| 80+ years | Non-Hispanic Black vs. Hispanic (All Races) | 561 vs. 653 | 0.95 (0.83-1.09) | 0.466 | - |
|  | Non-Hispanic Others vs. Hispanic (All Races) | 725 vs. 653 | 0.90 (0.79-1.02) | 0.103 |  |
|  | Non-Hispanic White vs. Hispanic (All Races) | 8294 vs. 653 | 1.01 (0.92-1.11) | 0.821 |  |


|  |  |  | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
| Covariate | Level | N | Hazard Ratio $(95 \% \text { CI) }$ | $\begin{aligned} & \text { HR P- } \\ & \text { value } \end{aligned}$ | Type3 <br> P-value |
| * Number of observations in the original data set $=113885$. Number of observations used $=$ 113885. <br> ** Backward selection with an alpha level of removal of .05 was used. The following variables were removed from the model: Origin recode NHIA (Hispanic, Non-Hisp). <br> *** The estimated stratified treatement effect was controlled by: \% Ever Smoker (males age 18+ - sae 2000-) (quartile), \% Moved from outside US 2000 (quartile), \% Unemployed 2000 (quartile), \% Urban 2000 (quartile), Age_80*RaceandoriginrecodeNHWNHBNHAIA, CS tumor size in $\mathrm{cm}(2004+$ ) (quartile), Cancer-directed surgery, Combined SEER summary stage, Grade, Marital status at diagnosis, Sex, Site recode ICD-O-3/WHO 2008, Year of diagnosis |  |  |  |  |  |
|  |  |  |  |  |  |
|  |  |  |  |  |  |

Table 5.2 Interaction Effect Between Race and Age on Disease-Specific Survival

| Covariate | Level | N | Survival months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio <br> (95\% CI) | HR P- <br> value | Type3 P- <br> value |
| Comparisons | Race : |  | - | - | <. 001 |
| Stratified by |  |  |  |  |  |
| Age : |  |  |  |  |  |
| 20-39 years | Non-Hispanic Black vs. Hispanic (All Races) | 296 vs. 412 | 1.06 (0.81-1.39) | 0.686 | - |
|  | Non-Hispanic Others vs. Hispanic (All Races) | 466 vs. 412 | 0.85 (0.64-1.11) | 0.232 |  |
|  | Non-Hispanic White vs. Hispanic (All Races) | 1918 vs. 412 | 0.76 (0.61-0.95) | 0.014 |  |
| 40-59 years | Non-Hispanic Black vs. Hispanic (All Races) | 6444 vs. 3411 | 1.40 (1.30-1.50) | <. 001 | - |
|  | Non-Hispanic Others vs. Hispanic (All Races) | 2881 vs. 3411 | 0.87 (0.78-0.96) | 0.004 |  |
|  | Non-Hispanic White vs. Hispanic (All Races) | $\begin{gathered} 35207 \text { vs. } \\ 3411 \end{gathered}$ | 0.88 (0.83-0.94) | <. 001 |  |
| 60-79 years | Non-Hispanic Black vs. Hispanic (All Races) | 5368 vs. 3656 | 1.11 (1.04-1.19) | 0.003 | - |
|  | Non-Hispanic Others vs. Hispanic (All Races) | 3017 vs. 3656 | 0.87 (0.80-0.95) | 0.001 |  |
|  | Non-Hispanic White vs. Hispanic (All Races) | $\begin{gathered} 40576 \text { vs. } \\ 3656 \end{gathered}$ | 0.89 (0.84-0.94) | <. 001 |  |
| 80+ years | Non-Hispanic Black vs. Hispanic (All Races) | 561 vs. 653 | 1.03 (0.88-1.22) | 0.695 | - |
|  | Non-Hispanic Others vs. Hispanic (All Races) | 725 vs. 653 | 1.02 (0.87-1.21) | 0.788 |  |
|  | Non-Hispanic White vs. Hispanic (All Races) | 8294 vs. 653 | 0.97 (0.86-1.11) | 0.692 |  |



Table 6 Demographics of the Patient After Propensity Score Matching

| Covariate | Level | $\begin{gathered} \text { Hispanic (All } \\ \text { Races) } \mathbf{N}=\mathbf{3 2 3 0} \\ \mathbf{N}(\%) \end{gathered}$ | Non-Hispanic <br> Black $\mathbf{N}=3230$ N (\%) | Non-Hispanic <br> Others $\mathbf{N}=\mathbf{3 2 3 0}$ (\%) | Non-Hispanic <br> White $\mathbf{N}=\mathbf{3 2 3 0}$ <br> (\%) | Standardiz <br> ed <br> Difference |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Age | 20-39 years | 95 (2.94) | 89 (2.76) | 91 (2.82) | 76 (2.35) | 0.037 |
|  | 40-59 years | 1428 (44.21) | 1442 (44.64) | 1444 (44.71) | 1427 (44.18) | 0.011 |
|  | 60-79 years | 1478 (45.76) | 1479 (45.79) | 1450 (44.89) | 1494 (46.25) | 0.027 |
|  | 80+ years | 229 (7.09) | 220 (6.81) | 245 (7.59) | 233 (7.21) | 0.030 |
| Sex | Female | 756 (23.41) | 712 (22.04) | 798 (24.71) | 712 (22.04) | 0.063 |
|  | Male | 2474 (76.59) | 2518 (77.96) | 2432 (75.29) | 2518 (77.96) | 0.063 |
| Marital <br> status at <br> diagnosis | Married | 1653 (51.18) | 1637 (50.68) | 1629 (50.43) | 1675 (51.86) | 0.028 |
|  | Other | 1577 (48.82) | 1593 (49.32) | 1601 (49.57) | 1555 (48.14) | 0.028 |
| Year of diagnosis | 2000-2006 | 1515 (46.9) | 1454 (45.02) | 1513 (46.84) | 1715 (53.1) | 0.162 |
|  | 2007-2013 | 1715 (53.1) | 1776 (54.98) | 1717 (53.16) | 1515 (46.9) | 0.162 |
| Cancerdirected surgery accepted | No | 1800 (55.73) | 1750 (54.18) | 1777 (55.02) | 1719 (53.22) | 0.050 |
|  | Unknown | 30 (0.93) | 34 (1.05) | 28 (0.87) | 16 (0.5) | 0.064 |
|  | Yes | 1400 (43.34) | 1446 (44.77) | 1425 (44.12) | 1495 (46.28) | 0.059 |
| Cancer site | Larynx | 816 (25.26) | 955 (29.57) | 801 (24.8) | 953 (29.5) | 0.107 |
|  | Oral cavity | 1317 (40.77) | 1238 (38.33) | 1370 (42.41) | 1254 (38.82) | 0.083 |
|  | Pharynx | 1097 (33.96) | 1037 (32.11) | 1059 (32.79) | 1023 (31.67) | 0.049 |
| Tumor grade | Grade II | 1261 (39.04) | 1251 (38.73) | 1285 (39.78) | 1327 (41.08) | 0.048 |
|  | Grade III | 914 (28.3) | 921 (28.51) | 838 (25.94) | 880 (27.24) | 0.058 |
|  | Grade IV | 28 (0.87) | 22 (0.68) | 23 (0.71) | 24 (0.74) | 0.021 |
|  | Unknown | 691 (21.39) | 649 (20.09) | 729 (22.57) | 634 (19.63) | 0.072 |
|  | Grade I | 336 (10.4) | 387 (11.98) | 355 (10.99) | 365 (11.3) | 0.050 |
| CS tumor size in cm (quartile) | $>=0,<=2$ | 278 (8.61) | 285 (8.82) | 289 (8.95) | 226 (7) | 0.072 |
|  | $>2,<=3$ | 352 (10.9) | 391 (12.11) | 367 (11.36) | 326 (10.09) | 0.064 |
|  | $>3,<=4$ | 422 (13.07) | 462 (14.3) | 420 (13) | 388 (12.01) | 0.068 |
|  | $>4,<=99$ | 357 (11.05) | 381 (11.8) | 374 (11.58) | 330 (10.22) | 0.050 |
|  | Unknown | 1821 (56.38) | 1711 (52.97) | 1780 (55.11) | 1960 (60.68) | 0.156 |


|  |  | Hispanic (All | Non-Hispanic | Non-Hispanic | Non-Hispanic | Standardiz |
| :--- | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Covariate |  | Races) $\mathbf{N}=\mathbf{3 2 3 0}$ | Black $\mathbf{N}=\mathbf{3 2 3 0}$ | Others $\mathbf{N}=\mathbf{3 2 3 0}$ | White $\mathbf{N}=\mathbf{3 2 3 0}$ | ed |
|  |  | $\mathbf{N}(\%)$ | $\mathbf{N}(\%)$ | $\mathbf{N}(\%)$ | $\mathbf{N}(\%)$ | Difference |


| Combined | Distant | $613(18.98)$ | $591(18.3)$ | $623(19.29)$ | $594(18.39)$ | 0.025 |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |

SEER
summary
stage

|  | Localized | $1061(32.85)$ | $1099(34.02)$ | $1075(33.28)$ | $1119(34.64)$ | 0.038 |
| :---: | :--- | :---: | :---: | :---: | :---: | :---: |
|  | Regional | $1399(43.31)$ | $1392(43.1)$ | $1344(41.61)$ | $1378(42.66)$ | 0.034 |
|  | Unknown | $157(4.86)$ | $148(4.58)$ | $188(5.82)$ | $139(4.3)$ | 0.069 |
| $\%<$ High | $>=3,<=16$ | $547(16.93)$ | $603(18.67)$ | $526(16.28)$ | $601(18.61)$ | 0.063 |

school
education
(quartile)

|  | $>16,<=19$ | $959(29.69)$ | $977(30.25)$ | $993(30.74)$ | $1025(31.73)$ | 0.044 |
| :--- | :--- | :---: | :---: | :---: | :---: | :---: |
|  | $>19,<=27$ | $690(21.36)$ | $611(18.92)$ | $705(21.83)$ | $587(18.17)$ | 0.091 |
|  | $>27,<=51$ | $1034(32.01)$ | $1039(32.17)$ | $1006(31.15)$ | $1017(31.49)$ | 0.022 |
| $\%$ | $>=1,<=5$ | $438(13.56)$ | $426(13.19)$ | $500(15.48)$ | $427(13.22)$ | 0.065 |

Unemploye
d (quartile)

|  | $>5,<=6$ | $1004(31.08)$ | $942(29.16)$ | $1038(32.14)$ | $1037(32.11)$ | 0.064 |
| :--- | :--- | :---: | :---: | :---: | :---: | :---: |
|  | $>6,<=9$ | $1328(41.11)$ | $1388(42.97)$ | $1174(36.35)$ | $1293(40.03)$ | $\mathbf{0 . 1 3 6}$ |
|  | $>9,<=18$ | $460(14.24)$ | $474(14.67)$ | $518(16.04)$ | $473(14.64)$ | 0.050 |
| \% Urban | $>=0,<=76$ | $191(5.91)$ | $221(6.84)$ | $182(5.63)$ | $153(4.74)$ | 0.090 |
| (quartile) |  |  |  |  |  |  |
|  | $>76,<=95$ | $711(22.01)$ | $703(21.76)$ | $709(21.95)$ | $748(23.16)$ | 0.033 |
|  | $>95,<=99$ | $946(29.29)$ | $960(29.72)$ | $840(26.01)$ | $973(30.12)$ | 0.092 |
|  | $>99,<=100$ | $1382(42.79)$ | $1346(41.67)$ | $1499(46.41)$ | $1356(41.98)$ | 0.096 |
| \% Moved | $>=0,<=2$ | $184(5.7)$ | $134(4.15)$ | $155(4.8)$ | $116(3.59)$ | $\mathbf{0 . 1 0 0}$ |

from
outside US
(quartile)

| $>2,<=3$ | $539(16.69)$ | $553(17.12)$ | $551(17.06)$ | $525(16.25)$ | 0.023 |
| :--- | :---: | :---: | :---: | :---: | :---: |
| $>3,<=5$ | $938(29.04)$ | $1035(32.04)$ | $827(25.6)$ | $956(29.6)$ | $\mathbf{0 . 1 4 3}$ |
| $>5,<=10$ | $1569(48.58)$ | $1508(46.69)$ | $1697(52.54)$ | $1633(50.56)$ | $\mathbf{0 . 1 1 7}$ |


|  |  | Hispanic (All | Non-Hispanic | Non-Hispanic | Non-Hispanic | Standardiz |
| :--- | :---: | :---: | :---: | :---: | :---: | :---: |
| Covariate | Level | Races) $\mathbf{N}=\mathbf{3 2 3 0}$ | Black $\mathbf{N = 3 2 3 0}$ | Others $\mathbf{N}=\mathbf{3 2 3 0}$ | White $\mathbf{N}=\mathbf{3 2 3 0}$ | ed |
|  |  | $\mathbf{N}(\%)$ | $\mathbf{N}(\%)$ | $\mathbf{N}(\%)$ | $\mathbf{N})$ | (\%) |
| 0 Difference |  |  |  |  |  |  |

Smoker
(males age
18+)
(quartile)

| $>45,<=51$ | $1060(32.82)$ | $1123(34.77)$ | $1060(32.82)$ | $1145(35.45)$ | 0.056 |
| :--- | :---: | :---: | :---: | :---: | :---: |
| $>51,<=56$ | $548(16.97)$ | $508(15.73)$ | $486(15.05)$ | $467(14.46)$ | 0.069 |
| $>56,<=73$ | $169(5.23)$ | $203(6.28)$ | $182(5.63)$ | $185(5.73)$ | 0.045 |

Table 7.1 The Effect of Race on Overall Survival in Matched Sample

| Covariate | Level | N | Survival Months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio (95\% CI) | HR P- <br> value | $\begin{gathered} \text { Type3 P- } \\ \text { value } \end{gathered}$ |
| Race | Hispanic (All | 3230 | 1.09 (1.01-1.16) | 0.021 | <. 001 |
|  | Races) |  |  |  |  |
|  | Non-Hispanic | 3230 | 1.26 (1.18-1.35) | <. 001 |  |
|  | Black |  |  |  |  |
|  | Non-Hispanic | 3230 | 0.90 (0.84-0.97) | 0.005 |  |
|  | Others |  |  |  |  |
|  | Non-Hispanic | 3230 | - | - |  |
|  | White |  |  |  |  |

Analysis was taken the clustering effect within Patient_ID into account, and N represented number of Patient_ID-times.

Table 7.2 The Effect of Race on Disease-Specific Survival in Matched Sample

| Covariate | Level | N | Survival Months |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  |  |  | Hazard Ratio (95\% CI) | HR P- <br> value | Type3 P- <br> value |
| Race | Hispanic (All | 3230 | 1.13 (1.04-1.23) | 0.005 | <.001 |
|  | Races) |  |  |  |  |
|  | Non-Hispanic | 3230 | 1.31 (1.21-1.42) | $<.001$ |  |
|  | Black |  |  |  |  |
|  | Non-Hispanic | 3230 | 0.98 (0.90-1.07) | 0.648 |  |
|  | Others |  |  |  |  |
|  | Non-Hispanic | 3230 | - | - |  |
|  | White |  |  |  |  |

Analysis was taken the clustering effect within Patient_ID into account, and N represented number of Patient_ID-times.

Table 7.3 The Effect of Race on Overall Survival in Laryngeal Cancer Matched Sample
Survival Months

| Covariate | Level | N | Hazard Ratio <br> $(\mathbf{9 5 \%} \mathbf{C I})$ | HR P- <br> value | Type3 P- <br> value |
| :--- | :--- | :---: | :---: | :---: | :---: |
| Race | Hispanic (All <br> Races) | 969 | $0.84(0.74-0.96)$ | $\mathbf{0 . 0 1 0}$ | $<.001$ |
|  | Non-Hispanic <br> Black | 969 | $1.02(0.90-1.15)$ | 0.812 |  |
| Non-Hispanic <br> Others | 969 | $0.75(0.66-0.86)$ | $<.001$ |  |  |
| Non-Hispanic <br> White | 969 | - | - |  |  |

Analysis was taken the clustering effect within Patient_ID into account, and N represented number of Patient_ID-times.

Figure 1.1 Kaplan-Meier Overall Survival Estimate of HNC Patients by Race


Figure 1.2 Kaplan-Meier Disease-Specific Survival Estimate of HNC Patients by Race


Figure 1.3 Kaplan-Meier Overall Survival Estimate of Oral Cavity Cancer Patients by Race


Figure 1.4 Kaplan-Meier Overall Survival Estimate of Pharyngeal Cancer Patients by Race


Figure 1.5 Kaplan-Meier Overall Survival Estimate of Laryngeal Cancer Patients by Race


Figure 2.1 The Sum of Within-cluster Distance vs. Matching ID


Figure 2.2 The Sum of Within-cluster Distance vs. Matching ID for Laryngeal Cancer Sample


Figure 2.3 The Sum of Within-cluster Distance vs. Matching ID for Oral Cavity Cancer Sample


Figure 2.4 The Sum of Within-cluster Distance vs. Matching ID for Pharyngeal Cancer Sample


Figure 3.1 Kaplan-Meier Overall Survival Estimate of HNC Patients by Race in Matched Sample


Figure 3.2 Kaplan-Meier Disease-Specific Survival Estimate of HNC Patients by Race in Matched Sample


Figure 3.3 Kaplan-Meier Overall Survival Estimate of Oral Cavity Cancer Patients by Race in Matched Sample


Figure 3.4 Kaplan-Meier Overall Survival Estimate of Pharyngeal Cancer Patients by Race in Matched Sample


Figure 3.5 Kaplan-Meier Overall Survival Estimate of Laryngeal Cancer Patients by Race in Matched Sample


