



Article Disentangling Population Level Differences in Juvenile Migration Phenology for Three Species of Salmon on the Yukon River

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Abstract: Migration phenology influences many important ecological processes. For juvenile Pacific salmon, the timing of the seaward migration from fresh to marine waters is linked to early marine survival and adult returns. Seaward migration phenology is determined by interactions between the intrinsic attributes of individual species and environmental factors that are acting upon them. Temperature and discharge are two factors of the freshwater environment that have been shown to influence intra- and interannual variation in juvenile salmon phenology, but these factors may affect the migrations of sympatric species differently. Understanding how variations in phenology change with environmental heterogeneity is a critical first step in evaluating how the future climate may affect salmon. This is especially crucial for high-latitude rivers, where the pace of climate change is nearly twice as rapid as it is for more temperate areas. This research investigates the influence of river conditions on the seaward migration phenology of Chinook, chum, and coho salmon in the Yukon River. The results identified species-specific differences in the factors affecting migration duration, concentration, and skew and provide a starting point for a more detailed examination of how phenological variability may affect the temporal matching of juvenile salmon with biological resources and environmental conditions for optimal survival.

Keywords: phenology; Pacific salmon; Yukon River; smolt; freshwater migration; environmental variables

1. Introduction

Pacific salmon have been an important means of subsistence and cultural resource for the Yukon River area for millennia. In the recent past, they have also supported robust commercial fisheries that provided employment in remote locations with limited economic opportunities. Chinook salmon (Oncorhynchus tshawytscha) have been in a prolonged period of low productivity since the early 1990s, resulting in the complete closure of commercial and subsistence fisheries in 2011 and 2020, respectively. Chum salmon (Oncorhynchus keta) and coho salmon (Oncorhynchus kisutch) suffered catastrophic declines in adult returns beginning in 2021 and have been closed to commercial and subsistence fishing since. The role of freshwater versus marine factors in these declines is unclear. Substantial effort has been invested in understanding the factors contributing to the survival and recruitment of Chinook salmon once they enter the marine environment [1,2]. In the Northern Bering Sea (NBS), research on juvenile Yukon River Chinook salmon at the end of the first marine summer identified a positive correlation between the abundance of Canadian-origin juvenile salmon in the NBS and adult returns to the Yukon River [3]. This relationship suggests that factors affecting recruitment likely occur earlier in the salmon life cycle.

The migration from freshwater to the marine environment is a critical life-history threshold for juvenile salmon, requiring changes in morphology, physiology, and behavior in response to environmental cues. Migration phenologies have evolved to exploit



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Copyright: © 2023 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). natural variations in local abiotic conditions and resource pulses that favor survival and growth. A substantial body of research on juvenile salmon indicates that the onset of seaward migration is linked to environmental conditions (see [4–8]). The timing of seaward migration has been linked with early marine survival and the subsequent return to spawn as adults. Survival is higher when migration coincides with optimal ocean prey and environmental conditions [9]. Seaward migration phenology is a population-level process that can vary between co-occurring salmonid species [7] as species with different life histories respond to the same environmental factors in different ways. Very few studies have examined synchrony in the downstream migration between sympatric species [10,11]. A better understanding of this phenological synchrony could identify whether relationships between migration timing and hydrology vary across species, which hydrologic factors are most important in determining seaward migration phenology for each species and enhance understanding of how natural and climate-induced environmental changes may alter migration patterns in the future.

The environmental factors affecting the downstream migration of anadromous species have been investigated by a number of researchers (see [10,12]). Some evidence suggests that water temperature and discharge are important factors in determining the timing, rate, and duration of migration [13–15], while other studies have found little to no relationship with these factors [16]. Many of the studies evaluating the role of both temperature and discharge have focused on small rivers [17] or highly altered systems with controlled flow regimes [18,19], and as such, may not be applicable to large natural rivers, particularly those in high latitudes with strong seasonal temperature extremes. Climate change is having a pronounced effect on arctic ecosystems. Over the last 60 years, climate warming in Alaska has been almost twice as rapid as in the rest of the United States [20], impacting both marine and freshwater systems can cause disruptions to the hydrological cycle [21] and the complex interactions between temperature, ice, precipitation, and permafrost. These changes are expected to affect river discharge and temperature during juvenile salmon rearing and seaward migration, with the potential to disrupt seaward migration cycles.

This research investigates the influence of river conditions on the seaward migration phenology of Chinook, chum, and coho salmon in a large, subarctic river. The primary objectives of the research are to evaluate synchrony in emigration timing and the shape of the migration phenology distributions between these sympatric species and to identify environmental covariates that may influence these differences.

2. Materials and Methods

2.1. Study Area

The Yukon River, located in northwest Canada and central Alaska, is the fourth-largest river in North America and one of its largest and most diverse intact ecosystems [22]. The Yukon River watershed encompasses over 832,000 km² (Figure 1) and is divided into 13 major hydrologic basins [22,23], all of which are underlain by varying amounts of permafrost [24]. The river is fed by eight major tributaries and numerous distributaries. Below St. Mary's, approximately 168 miles upstream from the southern river mouth, the Yukon splits into three primary distributaries, each discharging into the Bering Sea. These distributaries flow across the Yukon River Delta, a gently sloping subarctic coastal plain, and across the submerged delta platform that extends 10–30 km offshore. Free-floating and landfast ice occur over the delta platform and along the delta shoreline seasonally from late fall through early spring [25]. Sub-ice channels extend from major distributaries, transporting Yukon River flow offshore during ice-covered periods [26].



Figure 1. Map of the Yukon River Basin (gray) showing the location of the USGS gauge at Pilot Station and Fairbanks. Black box corresponds to the inset showing the study area. Black circles are the nine permanent biological sampling stations.

Yukon River hydrology changes seasonally. For a six-to-nine-month period, the region is snow-covered with minimal streamflow. Peak discharges occur concurrent with or shortly after river breakup in the early spring, and high river flow continues until ice-up in late fall. In the lower river, discharge ranges from as low as $1200 \text{ m}^3 \text{ s}^{-1}$ in winter to a springtime peak in excess of 19,800 m³ s⁻¹ [27]. The spring discharge pulse is primarily driven by snow melt [23,28], with additional contributions from glacial ice melt, precipitation, and melting permafrost [29–31]. The relationship between accumulated snow (snowpack), snow melt, and river ice breakup is complex [23,28]. As temperatures rise in the spring, the snowpack undergoes a period of melting and refreezing as the snowpack becomes fully saturated with water. This period is known as the "snow melt duration" [28] and is followed by the snow off date, which is the last day of significant snow cover [23] and a period of reduced surface albedo and enhanced solar energy loading [32]. Snow off and the onset of snow melt are correlated with the timing of river breakup and discharge and snow off is also correlated to spring air temperatures across much of the Yukon basin [23]. The timing of

the highest river discharge coincides with the start of annual migrations of juvenile salmon from the river to the Bering Sea; however, the extent to which earlier hydrological processes, such as the snow melt duration, affect seaward migration timing is unknown.

Chinook, chum, and coho salmon have different life history characteristics. Chinook and chum salmon spawn along the river and its tributaries up to the headwaters, over 3000 km from the coast [33]. Coho salmon spawning locations are not well documented, but they are believed to spawn throughout the lower river into the Tanana River, but not past the Yukon River's boundary with Canada [34]. Chinook salmon are composed of 36 spatially and genetically distinct populations with similar timings for adult spawning migrations [35]. Chum salmon can be genetically assigned to two populations: summer and fall [36]. Summer chum adults enter the river in June and spawn predominantly in the lower river, while fall chum adults enter the river in July and spawn higher in the watershed into Canada. Juvenile Chinook salmon spend a year rearing in freshwater before migrating but may move downstream from the spawning areas during this rearing period [37,38]. Juvenile chum salmon migrate to the ocean shortly after hatching from the locations where they were spawned, but Chinook spend a year rearing in freshwater before emigrating. Juvenile coho generally spend one to three years rearing in freshwater before migrating to the ocean and may move considerably during this time.

2.2. Biological Data

Since 2014, salmon and other juvenile fish species have been captured at nine permanent stations on the three main distributaries of the lower Yukon River (Figure 1) near the village of Emmonak. Stations were sampled three times per week from ice-out from the middle of May through the end of July using surface tow nets (6.8 m long \times 1.8 m² at the mouth, tapering to a 0.3 m \times 0.3 m cod end). Few juvenile salmon are captured after the end of July, and it is difficult to know whether these salmon are emigrating or relocating to winter nursery areas farther downstream. Therefore, the end of July was selected as the end of the migration. At each station, three replicate 15-min tows were performed during daylight hours by two small (i.e., 6 to 7 m) open skiffs towing against the current and holding the net open. Surface temperature measurements were taken at the start of sampling at each station. At the end of each tow, captured fish were sorted into species. All juvenile Chinook and coho salmon were retained for further analysis, while individuals of other species, including chum salmon, were enumerated, measured, and released. Retained salmon were placed in refrigerated bait coolers by station and transported alive to the field station in Emmonak. At the field station, salmon were photographed, euthanized, measured to 1 mm fork length, weighed to 0.001 g, and frozen. The caudal fin was clipped and placed in ethanol for genetic stock identification. The frozen samples were transported to the National Oceanic and Atmospheric Administration's (NOAA) laboratory in Juneau, Alaska.

2.3. Phenology Modeling

Seaward migration phenology in juvenile salmon can be thought of as a series of migration completions carried out by individuals in a population over time. While phenology is typically evaluated using discrete events, such as the onset, mid-point, and end of the migration (generally determined to be 5%, 50%, and 90–95% of the total run, respectively) [39,40], this approach loses information on the shape of the migration distribution throughout its duration [39,41,42]. These methods do not easily lend themselves to the comparison of phenological phenomena between species because they lack common model parameters to evaluate against environmental variables. Parameters, such as the time required to complete the migration, the spread of individuals over the migration duration, and the skew of the distribution, are all variables that can be affected by the migration environment and by changes in that environment over time. Phenology distributions are often leptokurtic or skewed, and their shapes may change from year to year. These factors limit the applicability of measures of central tendency such as the median and the

mean [42–44] for comparing differences in phenology between species and time periods. Recognizing these limitations, analytical methods that consider the full phenological distribution have been developed to more accurately compare the overlap or gaps between the distributions of sympatric species or the same species over time (see [42,44–49]). In this research, migration is modeled in terms of cumulative probabilities, making it possible to compare migrations of different durations and numbers of individuals. The model parameters describe important characteristics of the phenology distribution that can be used to compare models to one another and to environmental covariates.

The biological dataset used for this research comprised daily totals of Chinook, chum, and coho salmon smolts intercepted at the sampling stations for the periods 2014–2019 and 2021. In 2020, the start of sampling was delayed by two weeks because of the COVID-19 pandemic. Since the juvenile salmon emigration was underway by the time sampling started, these data were not included in this analysis. Daily totals for a given sampling day were pooled across all stations sampled that day. The study was designed for the effort to remain constant across years, but some differences in effort occurred due to equipment malfunctions and weather delays. To address this, catch per unit effort (CPUE) for each sampling day was calculated as the total catch divided by the number of tows. The use of CPUE instead of total count made no appreciable difference to the models, so the total count was selected for use due to its easier interpretability.

Daily catches of Chinook, chum, and coho salmon were modeled for each year as the cumulative probability of migration using the inverse logit, as modified by Franco [43]:

$$\frac{y}{y_{max}} = 1 - \left(1 - \frac{r_{max}}{1 + e^{-c(x-t)}}\right)^x$$

Using proportions as the response makes this model robust to sample size differences between years and species. Parameter r_{max} , the maximum proportional rate of migration completion, is a dimensionless variable that describes the rate at which the cumulative distribution function (*cdf*) increases to its maximum value. In this study, all intercepted salmon were defined as successful migrations, and the end of the sampling period was the end of July. Very small numbers of juvenile salmon are captured close to or at the end of the sampling period. As a result, in this analysis, r_{max} is nearly equivalent to the length of the sampling period. Parameter *c* is a measure of the concentration or spread of the distribution (units: time⁻¹); increasing values of *c* reduce the spread of the probability density function (*pdf*). Parameter *t* is a weighted measure of the migration duration.

The start of migration was defined as the number of days from the official ice breakup date in Emmonak, which changed each year [50]. Breakup date was chosen over "day of year" because field sampling indicates that the bulk of the migration for the species being evaluated occurs after ice leaves the river [51,52]. The objective of the research was to compare seaward migration phenologies between salmon species during the period in which that migration can occur each year. The day of the year on which migration becomes possible is determined by the ice breakup date; therefore, the day of year by itself has limited value in this analysis.

Models were fit using non-linear least squares regression employing the Levenberg– Marquardt algorithm in R [53] using the package *nlstimedist* [54]. Once all parameters were estimated, they were used to calculate *pdfs* for each species and year.

2.4. Environmental Data and Modeling

Subarctic hydrology and the long, downstream migration differentiate juvenile salmon phenology in the Yukon River from that of other North American rivers. Abrupt changes in both temperature and discharge during the transition from winter to spring, and considerable variations in temperature and flow throughout the migratory period, may affect seaward migration phenology at different time scales. The effects of these temporal variations in the environment may be different for each salmon species. Environmental covariates that may influence hydrology and phenology occur prior to and during the migration period.

Prior to migration, snow off and snow melt affect river breakup and discharge. These early hydrological changes may influence seaward migration. Observations of snow-off date and onset of snow melt are extremely sparse and temporally discontinuous in the Yukon River basin. Melt onset is moderately correlated to spring air temperatures [23], which may serve as proxies for direct measurements of these variables; however, continuous air temperature datasets are also sparse. The most complete air temperature data for the analysis period were obtained from the Natural Resources Conservation Service (NRCS) snow telemetry station at Little Chena Ridge near Fairbanks, AK (65.12 N, 146.12 W). This location is not on the mainstem of the Yukon River but falls within a catchment that includes the U.S. Geological Survey (USGS) gage at Pilot Station (61.93 N, 162.88 W) [27], which provides discharge data. The gage is located approximately 120 miles upstream of the river mouth and prior to the point where the river divides into three primary distributaries. This gage is considered an outflow point for the entire Yukon River drainage [55] and is the lowest downstream gaging station. The data from this gage provide discharge information once the spring runoff cycle begins and, in combination with early spring air temperatures, can be used to evaluate early hydrology on seaward migration.

Changes in discharge and water temperature once migration has started are hypothesized to affect the shape of the migration phenology distribution. High discharge may compress the distribution, shortening the migration duration. Warm water temperatures may also encourage earlier and faster migration [56]. River discharge during the migration period was obtained from the Pilot Station gage. Water temperature data have not been systematically collected on the Yukon River. The USGS maintains four gauges that record water temperatures on the mainstem of the Yukon River, but none of these had a complete time series of temperatures covering the analysis period. Water temperatures collected during biological sampling do not provide information on temperatures that may have influenced migration prior to interception since all salmon are collected near the river mouth near the end of the migration. Research has found that, at the monthly timescale, patterns between air and water temperatures are similar [57]. Therefore, air temperatures are used in this analysis as proxies for water temperatures during the seaward migration period.

To evaluate when temperature and discharge covariates are most strongly associated with phenological variations and at what scale, a range of summary statistics was evaluated for each month of the migratory period (Table 1). Maximum and mean air temperatures for each month were compiled from daily air temperature data from the snow telemetry station at Little Chena Ridge [58]. Maximum and mean air temperatures for April and May are included to represent solar loading influencing snow-off and melt duration periods. Maximum and mean air temperatures during the migration period are proxies for water temperatures. Monthly mean discharge, monthly range of discharge, and maximum monthly discharge were compiled from daily discharge records collected at Pilot Station.

Table 1. Response and explanatory variables included in correlation analysis of migration phenologies of juvenile Chinook, chum, and coho salmon from the Yukon River, 2014–2019 and 2021. pdf = probability density function, cdf = cumulative distribution function.

Variable Name	Meaning					
Response variables						
c	A measure of the temporal concentration of the phenology distribution. Increasing values of <i>c</i> reduce the spread of the <i>pdf</i> .					
t	A measure of the migration time lag. Higher values of <i>t</i> shift the <i>cdf</i> and <i>pdf</i> to the right.					
skew	A measure of distributional asymmetry in the <i>pdf</i> .					
Explanatory variables—Discharge						
Range of discharge	The difference between the maximum and minimum values of discharge for each month in the migration period (May, June, and July) and for April, which is at the beginning or prior to the onset of peak discharge in spring.					
Average discharge	for April.					
Maximum discharge	The maximum discharge for each month in the migration period (May, June, and July) and for April.					
Explanatory variables—Air Temperature						
Maximum temperature	The maximum air temperature for each month in the migration period (May, June, and July) and for April as a proxy for water temperature.					
Average temperature	Average monthly temperatures for the migration period (May, June, and July) and for April as proxies for water temperatures.					

Maximum values represent the effect of extreme discharge events on phenology, while mean values represent the overall discharge trend. The monthly discharge range is a metric of how abruptly discharge changes. The monthly timescale was selected for consistency with air temperatures. Discharge variables were plotted as anomalies from the long-term mean values. The reference data for discharge covered the years from 2002 to 2013. This represents the continuously available data up to the study period following a gap in discharge data from 1997 to 2001.

Pearson's correlation coefficients were used to examine the relationship between the parameters of the phenology distributions from the logit models for each species and year and each of the environmental variables for discharge and temperature. The size of the sample dataset, which has only seven years of data for each species, is small when viewed in the context of frequentist statistics, increasing the possibility that significant correlations between the model parameters and the environmental variables may be a chance occurrence. There is much current debate on the validity of *p*-values as measures of the significance of statistical relationships (see [59-61]). This research utilized the *p*-values of the correlations as a first-pass evaluation of the strength of the relationship. A p-value of 0.05 suggests a stronger linear relationship than a higher value, but no claims about the significance of this relationship are made. The intent is to identify covariates and time periods from a large potential range that may inform interpretation of the seaward phenology distributions for future consideration. Because environmental variables rarely act on species in isolation, the univariate correlations used here can only be a first step in evaluating potential relationships. Correlations were computed using the *psych* package in R [62].

3. Results

All study years occurred during a period of earlier-than-average ice breakup at Emmonak. Prior to 2013, breakup dates varied widely but did not demonstrate a trend toward earlier or later breakups (Figure 2). Although only the study years 2016 and 2019 had earlier breakup dates than previously recorded, the consistently earlier-than-average breakup is anomalous for this time series. The timing of ice breakup was uncorrelated with peak discharge (r = 0.23, p = 0.62), which occurred between 9 and 29 days later.



Figure 2. Annual river ice breakup dates for the Yukon River near Emmonak. Orange circles indicate the study years. Blue dashed line shows the trend from 1979–2013.

There was substantial monthly and intra-annual variation in discharge and in the timing and magnitude of the discharge peak during the study period (Figure 3). The discharge did not vary substantially before mid-April. Peak discharge occurred between 22 May and 7 June, and the three highest discharges occurred coincident with the latest peak discharges: 3 June 2015, 7 June 2018, and 6 June 2019. The earliest and steepest increases in discharge occurred in 2014 and 2017. In most years, the discharge declined abruptly following the peak and continued to decline throughout the study period. The exceptions were the years 2014 and 2020. In 2014, discharge declined slowly into July. In 2020, discharge declined initially but then increased to a second peak in July.

Monthly air temperatures during the study period declined slightly but non-significantly in April and May (p = 0.5 and 0.4) and increased slightly but non-significantly in June and July (p = 0.1 and 0.07) (Figure 4). Maximum and mean May air temperatures were highest in 2015 and lowest in 2018. The highest maximum and mean June air temperatures occurred in 2019.

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Figure 3. (**A**) Daily and (**B**) monthly discharge from the USGS gauge at Pilot Station for each sampling year and month. (**A**) Mean daily discharge by year. Discharge prior to 16 April shows the same trend, so is omitted from this figure. (**B**) Box plots of discharge by month. Bold lines represent median discharge. Blue dots represent maximum discharge and orange dots represent mean discharge.



Figure 4. Monthly average air temperatures (°C) for the study period from the Little Chena Ridge SNOTEL station on the Yukon River. (**A**) Gray area represents 95% confidence intervals. (**B**) Box plots of air temperature by month. Bold lines represent median temperature. Blue dots represent maximum temperature and orange dots represent mean temperature.

The dataset used for this analysis included records of over 75,000 juvenile salmon migrating from the Yukon River between 2014–2019 and 2021. Juvenile chum salmon comprised 91% of the total count, with Chinook and coho salmon contributing 5.1% and 3.4%, respectively. The total number of each species varied considerably over time, but all species exhibited a dramatic decline in 2021 (Table S1). The number of Chinook and coho encountered in 2021 was 25% of the annual high, and chum salmon was only 18% of the annual high. The nonlinear models fitted to the migration proportions of each salmon species and year successfully converged. The model parameter values demonstrated small standard errors and highly significant fits (p < 0.001, Table S1).

The migration distributions for all species and years were moderately to heavily rightskewed, and all but three were leptokurtic (Table S1). Juvenile Chinook salmon had the most variability in the migration timing and duration of the three species evaluated (Figure 5). All three model parameters for Chinook salmon were correlated to the maximum discharge in June (Table 2). The skewness increased, and the migration was more concentrated (higher *c*) with higher values of discharge. Total migration time also decreased with higher discharge (negative correlation with *t*). Migration was later and more spread out in 2014, 2016, 2017, and 2021 (c = 0.08, 0.09, 0.11, and 0.01, respectively) when the maximum June discharge was lowest (Table S1). Migration was earlier and more concentrated in 2015, 2018, and 2019 (c = 0.20, 0.16, and 0.30) when the maximum June discharge was highest. The maximum discharge in May and the range of discharge in June were associated with a higher skewness and a lower migration time but not with increased migration concentration. The maximum June temperatures were also associated with a higher skewness and a shorter migration time.



Figure 5. Modeled cumulative distributions (**A**) and probability densities (**B**) of migration probabilities for juvenile Chinook (*Oncorhynchus tshawytscha*), chum (*O. keta*), and coho salmon (*O. kisutch*) from the Yukon River, Alaska, 2014–2019 and 2021.

Maniah laa	Chinook Salmon		Chum Salmon			Coho Salmon			
variables	Skew	С	t	Skew	С	t	Skew	С	t
Discharge									
Range—Apr	0.19	0.22	0.23	-0.21	-0.24	0.46	0.41	0.10	0.43
	(0.68)	(0.63)	(0.62)	(0.66)	(0.61)	(0.30)	(0.33)	(0.67)	(0.34)
Range—May	0.61	0.37	-0.80	0.56	0.76	-0.57	0.11	0.27	-0.66
	(0.15)	(0.42)	(0.03 *)	(0.19)	(0.05 *)	(0.18)	(0.82)	(0.56)	(0.11)
Range—Jun	0.76	0.68	-0.76	0.67	0.62	-0.75	-0.44	-0.51	-0.06
	(0.05 *)	(0.09)	(0.05 *)	(0.10)	(0.14)	(0.05 *)	(0.32)	(0.24)	(0.90)
Range—Jul	0.52	0.36	-0.28	0.04	0.22	0.08	0.43	0.38	-0.15
	(0.24)	(0.42)	(0.55)	(0.93)	(0.64)	(0.86)	(.33)	(0.40)	(0.74)
Max—Apr	0.38	0.43	0.03	-0.10	-0.16	0.27	0.29	-0.03	0.43
	(0.400)	(0.34)	(0.96)	(0.82)	(0.72)	(0.56)	(0.53)	(0.96)	(0.34)
Max—May	0.74	0.52	-0.75	0.49	0.65	-0.39	0.30	0.29	-0.44
	(0.05 *)	(0.23)	(0.05 *)	(0.27)	(0.11)	(0.39)	(0.51)	(0.52)	(0.32)
Max—June	0.89	0.77	-0.83	0.40	0.59	-0.40	0.21	0.20	-0.43
	(0.01 *)	(0.04 *)	(0.02 *)	(0.38)	(0.17)	(0.38)	(0.64)	(0.66)	(0.33)
Max—July	-0.61	-0.50	0.62	-0.59	-0.57	0.77	0.69	0.65	0.06
	(0.08)	(0.25)	(0.14)	(0.17)	(0.19)	(0.04 *)	(0.08)	(0.11)	(0.89)
Average—Apr	0.67	0.76	-0.30	0.03	-0.03	0.00	0.10	-0.25	0.33
	(0.150)	(0.05 *)	(0.52)	(0.96)	(0.96)	(0.99)	(0.83)	(0.99)	(0.46)
Average—May	-0.47	-0.45	0.33	-0.26	-0.36	0.33	0.44	0.16	0.12
	(0.10)	(0.25)	(0.46)	(0.58)	(0.42)	(0.47)	(0.33)	(0.73)	(0.70)
Average—Jun	0.49	0.51	-0.39	-0.15	0.10	0.23	0.75	0.69	-0.38
	(0.29)	(0.32)	(0.39)	(0.75)	(0.87)	(0.62)	(0.05 *)	(0.09)	(0.40)
Average—Jul	-0.73	-0.60	0.64	-0.20	-0.55	0.73	0.67	0.71	-0.01
	(0.06)	(0.15)	(0.12)	(0.20)	(0.23)	(0.07)	(0.10)	(0.07)	(0.98)
Air Temperature									
Max—Apr	-0.09	0.18	-0.02	-0.56	-0.62	0.38	0.49	-0.01	0.0-4
1	(0.84)	(0.70)	(0.96)	(0.19)	(0.14)	(0.40)	(0.26)	(0.99)	(0.93)
Max—May	0.52	0.62	-0.45	-0.16	0.17	-0.45	-0.32	-0.69	0.34
,	(0.24)	(0.14)	(0.32)	(0.37)	(0.72)	(0.31)	(0.48)	(0.90)	(0.45)
Max—Jun	0.77	0.75	-0.66	-0.71	0.44	-0.56	-0,24	-0.35	-0.28
	(0.04 *)	(0.05 *)	(0.10)	(0.50)	(0.32)	(0.19)	(0.60)	(0.43)	(0.55)
Max—Jul	0.80	(0.10)	-0.82	-0.84	0.81	-0.83	-0.17	-0.08	-0.13
	(0.03 *)	(0.83)	(0.02 *)	(0.02 *)	(0.03 *)	(0.02 *)	(0.72)	(0.87)	(0.78)
Avg—Apr	0.46	0.58	-0.45	-0.17	0.10	-0.44	-0.29	-0.71	0.30
0 1	(0.30)	(0.17)	(0.31)	(0.47)	(0.84)	(0.33)	(0.53)	(0.07)	(0.51)
Average—May	0.46	0.51	-0.50	-0.25	0.30	-0.57	-0.37	-0.66	0.24
8 ,	(0.30)	(0.24)	(0.26)	(0.22)	(0.51)	(0.18)	(0.41)	(0.10)	(0.61
Average—Jun	0.43	0.52	-0.38	-0.47	-0.07	-0.12	-0.12	-0.18	-0.29
0.	(0.34)	(0.23)	(0.40)	(0.57)	(0.88)	(0.78)	(0.79)	(0.71)	(0.52)
Average—Jul	0.66	0.38	-0.24	-0.40	0.54	-0.33	-0.44	-0.17	0.06
U .	(0.11)	(0.40)	(0.60)	(0.41)	(0.21)	(0.48)	(0.32)	(0.72)	(0.89)
Average—May Average—Jun Average—Jul	$\begin{array}{c} (0.30) \\ 0.46 \\ (0.30) \\ 0.43 \\ (0.34) \\ 0.66 \\ (0.11) \end{array}$	$(0.17) \\ 0.51 \\ (0.24) \\ 0.52 \\ (0.23) \\ 0.38 \\ (0.40)$	$\begin{array}{c} (0.51) \\ -0.50 \\ (0.26) \\ -0.38 \\ (0.40) \\ -0.24 \\ (0.60) \end{array}$	$\begin{array}{c} (0.47) \\ -0.25 \\ (0.22) \\ -0.47 \\ (0.57) \\ -0.40 \\ (0.41) \end{array}$	$\begin{array}{c} (0.04)\\ 0.30\\ (0.51)\\ -0.07\\ (0.88)\\ 0.54\\ (0.21)\end{array}$	$\begin{array}{c} (0.33) \\ -0.57 \\ (0.18) \\ -0.12 \\ (0.78) \\ -0.33 \\ (0.48) \end{array}$	$\begin{array}{c} (0.33) \\ -0.37 \\ (0.41) \\ -0.12 \\ (0.79) \\ -0.44 \\ (0.32) \end{array}$	$\begin{array}{c} (0.07) \\ -0.66 \\ (0.10) \\ -0.18 \\ (0.71) \\ -0.17 \\ (0.72) \end{array}$	$\begin{array}{c} (0.31) \\ 0.24 \\ (0.61 \\ -0.29 \\ (0.52) \\ 0.06 \\ (0.89) \end{array}$

Table 2. Pearson's correlation coefficients (and significance values) between parameters from nonlinear models fitted to juvenile Yukon River salmon migration proportions and variables of discharge and temperature for the years 2014–2019 and 2021. c = spread of the distribution, t = weighted measure of migration duration. Bolded values and asterisks indicate a *p*-value of 0.05 or lower.

For juvenile chum salmon, the skewness was not correlated with any of the discharge variables. The increased chum migration concentration was associated with a higher range in discharge in May. Since discharge was increasing throughout May in all sampling years, this equates to a higher concentration of migrating chum salmon as the slope of the discharge curve increased. The range of May discharge was highest in 2015 (c = 0.19), 2017 (c = 0.22), 2018 (c = 0.24), and 2019 (c = 0.16). A higher range of discharge in June was associated with a slower outmigration (larger *t*). Discharge generally decreased from peak values in June, but it was still increasing at the start of June in 2015, 2018, and 2019. The maximum July air temperatures negatively affected the skew and delays in migration and positively affected the concentration of the migration.

Coho salmon exhibited poor correlation with the environmental variables. The skewness was positively correlated with higher-than-average June discharges that occurred in 2018, 2019, and 2021. Migration was delayed in 2014, 2016, and 2019 by an average of 10 days compared to the other years, but this delay was not correlated with any of the environmental variables. Overall, the variation in migration concentration was lowest for coho ($c_{range} = 0.11$) compared to Chinook ($c_{range} = 0.22$) and chum ($c_{range} = 0.15$). Coho also had the least variation in migration duration ($t_{range} = 15$) compared to Chinook ($t_{range} = 38$) and chum ($t_{range} = 29$).

4. Discussion

River thermal and hydrologic conditions influence biological processes at a variety of scales. In high latitudes, seasonally ice-covered rivers, snow cover dynamics, and spring air temperatures strongly influence river breakup and discharge, while water temperatures and discharge impact the formation, thickness, and melting of river ice [56]. Hydrology in arctic and subarctic river systems exhibits high interannual variation resulting from changes in large-scale atmospheric circulation. Feedbacks associated with snow albedo and sea ice make these systems highly sensitive to climate change, with climate models projecting that average Arctic air temperatures will increase by as much as 3°C by 2040 [63]. Seasonally migratory species, such as salmon, can adjust their migration phenologies in response to altered hydrology and temperatures, but changes in migration timing may have adverse consequences if they result in mismatches between the migrants and suitable environmental and biological conditions [64]. This research identified differences in seaward migration phenologies, and environmental factors influencing these phenologies, between sympatric species of salmon in the Yukon River, which can be used to better understand how natural and climate-induced environmental changes may alter migration patterns in the future.

Differences in the correlation between the environmental variables and distributional aspects of the phenological time course for the species evaluated in this research have plausible explanations in their life histories. Juvenile Chinook salmon migration distributions were most strongly influenced by discharge variables in May and June, which affected at least two of the distribution parameters. Higher May and June discharge encouraged a more strongly skewed, concentrated, and faster migration. Chinook salmon move downriver as a mixed stock of the different populations [65] that were spawned throughout the watershed. The higher discharge measured in May and June at Pilot Station likely reflects the downstream accumulation of high discharges from farther upriver, thereby transporting juveniles from upriver rearing locations. The *cdfs* and *pdfs* from the Chinook migration models illustrate the high interannual variability in discharge distributions.

Coho life history may partially explain the low correlation between coho migration phenology and environmental variables. For coho, the increased average discharge in June was associated with an earlier outmigration (larger skew) but not with migration timings or concentrations. Discharge, at the scale at which it was measured in this study, may have less of an effect on coho because they spend their freshwater juvenile stage in low-velocity pools and off-channel habitats [66,67]. Juvenile coho also spend between one and three years rearing in freshwater before migrating from the river, and during this period they may relocate long distances to rearing areas, potentially closer to the river mouth [68,69]. The coho migration duration was the shortest of all the salmon species examined. These factors combined may work to limit the amount of time juvenile coho are exposed to discharge regimes. Coho may time their movement from low-flow habitats into the mainstem of the river for optimal timing. This could explain the correlation with average but not maximal river flows.

July air temperatures had contrary relationships with migration parameters for Chinook and chum salmon. For Chinook salmon, maximum July temperatures encouraged a higher skew and shorter migration durations. Temperature increases metabolism and, in the presence of sufficient food, can enhance growth. Juvenile Chinook salmon demonstrate a large range of adaptability to warmer temperatures, including the ability to maintain or enhance their swimming speed [70,71]. The maximum temperatures exceeded 22 °C during July 2015, but given the high heat capacity of water, it is likely that river temperatures were much lower and well within the adaptive thermal range of Chinook salmon. Energetically fit juvenile Chinook salmon may have been able to increase their swimming speed in response to higher temperatures, while others may have struggled with the higher metabolic demand, thus drawing out the tail of the distribution. For chum salmon, higher maximum air temperatures in July decreased the skew, increased the concentration, and decreased the migration duration. Over 80% of the chum salmon migration was complete by day 45, or the last week of June, in all study years (Figure 5). Straggler chum may have

been less abundant in years with higher maximum July temperatures, thereby decreasing migration duration. But any relationship with temperature seems spurious.

By quantifying essential attributes of the phenology distributions, the models used in this research can identify meaningful and species-specific relationships between juvenile migration distributions and environmental covariates. However, they represent only a beginning in the analysis to understand migration variability. The use of summary statistics at a monthly increment may not adequately capture variation in environmental parameters at the level at which salmon respond to them. For example, while the range in discharge during May was a good approximation of the speed of increasing discharge, this was not true for June, where in some years discharge was still increasing while in others it was decreasing from its maximum level in May. Using the results of the models as a starting point, a more detailed investigation can be made on how changes in discharge in May and June affect migration at smaller time scales.

Incorporating genetic stock information into the phenology models could provide a better understanding of the influence of spatial differences in the timing and magnitude of temperature and discharge on migration. Although genetic samples have been collected from individual Chinook and coho salmon since the start of the research in 2014, these data have only been analyzed for a few of the years. The analyses that have been performed were limited to mixed stock analyses providing stock proportions for three reporting groups (upper, middle, and lower Yukon) during three periods corresponding to stock composition variations in adult returns. These analyses demonstrated that the contribution of the three reporting groups changes during the course of the migration, with stocks from more distant spawning areas increasing in proportion later in the process. Unfortunately, the periods used in the analysis do not correspond to the observed variations in juvenile Chinook migration phenology. In 2022, the Alaska Department of Fish and Game agreed to use a newly developed genetic baseline for Chinook salmon to assign individual Chinook samples from 2014 to 2022 to six population reporting groups. When completed, population-level phenology models could increase understanding of the environmental complexities influencing migration while also helping to untangle interannual variations for Chinook salmon.

Genetic variation in Yukon chum salmon is lower than for Chinook. Chum salmon have been assigned to fall and spring stocks that have a relatively high level of spatial distinction. Preliminary data suggest that chum migrate as a mixed stock group, but proportions of each phenotype vary throughout the migration period [72]. The individual assignment of chum salmon to spring and fall stocks is not currently possible at a level of probability that meets management requirements. However, current research investigating temporal variations in the stock composition may still provide additional insight into the influence of hydrology and temperature on migration.

Water temperature data are critically important to address questions of the phenology, growth, and health of juvenile salmon, but have not been systematically collected in the Yukon River Basin at the spatial and temporal resolution needed for analysis. The collection of in situ temperature measurements is challenging due to the Basin's large size, remoteness, and relatively small human habitation footprint. For future analyses, the recent success in extrapolating relatively accurate river water temperatures from remotely sensed thermal data may provide a promising option for obtaining more accurate temperature data [73].

5. Conclusions

Phenological distributions contain a wealth of information on how populations respond to environmental heterogeneity. For juvenile Pacific salmon, the way that seaward migration unfolds with time is determined by the interaction between the intrinsic attributes of the individual species and the environmental factors that are acting upon them. This research focused on examining differences in seaward migration phenologies for three species of salmon whose migrations occur at similar times in the Yukon River. The results identified species-specific differences in the factors affecting migration duration, concentration, and skewness and provide a starting point for a more detailed examination of how phenological variability may affect the temporal matching of juvenile salmon with biological resources and environmental conditions for optimal survival.

Supplementary Materials: The following supporting information can be downloaded at: https: //www.mdpi.com/article/10.3390/jmse11030589/s1, Table S1. Summary statistics from juvenile outmigration phenology models for Yukon River Chinook, chum, and coho salmon for the years 2014–19 and 2021. N = number of sampled fish, r = maximum proportional rate of migration completion, c = spread of the distribution, *t* = weighted measure of migration duration.

Author Contributions: K.B.M. is the principal investigator of this research. She designed the study, participated in field data collection, analyzed the data, and is the lead author in writing the manuscript. C.M.W. is the lead field biologist responsible for overseeing the work of the vessel captains and crew; leading data field data collection, species identification, and sample management; managing the project data and implementing quality controls; and managing the grants and contractors. She prepared the figures and tables for the manuscript and provided the technical review. All authors have read and agreed to the published version of the manuscript.

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