Oxygenation of offshore Southern California marine basins through the Holocene

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November 30, 2022

Abstract

In the face of ongoing marine deoxygenation, understanding timescales and drivers of past oxygenation change is of critical importance. Marine sediment cores from tiered silled basins provide a natural laboratory to constrain timing and implications of oxygenation changes across multiple depths. Here, we reconstruct oxygenation and environmental change over time using benthic foraminiferal assemblages from sediment cores from three basins across the Southern California Borderlands: Tanner Basin (EW9504-09PC, 1194 m water depth), San Nicolas Basin (EW9504-08PC, 1442 m), and San Clemente Basin (EW9504-05PC ,1818 m). We utilize indicator taxa, community ecology, and an oxygenation transfer function to reconstruct past oxygenation, and we directly compare reconstructed dissolved oxygen to modern measured dissolved oxygen. We generate new, higher resolution carbon and oxygen isotope records from planktic (Globigerina bulloides) and benthic foraminifera (Cibicides mckannai) from Tanner Basin. Geochemical and assemblage data indicate limited ecological and environmental change through time in each basin across the intervals studied. Early to mid-Holocene (11.0-4.7 ka) oxygenation below 1400 m (San Clemente and San Nicolas) was relatively stable and reduced relative to modern. San Nicolas Basin experienced a multi-centennial oxygenation episode from 4.7-4.3 ka and oxygenation increased in Tanner Basin gradually from 1.7-0.8 ka. Yet across all three depths and time intervals studied, dissolved oxygen is consistently within a range of intermediate hypoxia (0.5-1.5 ml L-1 [O2]). Variance in reconstructed dissolved oxygen was similar to decadal variance in modern dissolved oxygen and reduced relative to Holocene-scale changes in shallower basins.

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2	Mari	ne Basins through the Holocene
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16	Key Po	pints:
17	•	In the Southern California Borderlands, oxygenation below 1400 m was stable and reduced
18		relative to modern from 11.0-4.7 ka
19	•	San Nicolas Basin experienced an oxygenation episode from 4.7-4.3 ka and oxygenation in
20		Tanner Basin increased at 1.7 ka relative to 5.4-1.7 ka
21	•	Variance in reconstructed Holocene dissolved oxygen concentration is similar to decadal scale
22		variance in modern dissolved oxygen
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Abstract

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42 43 In the face of ongoing marine deoxygenation, understanding timescales and drivers of past oxygenation change is of critical importance. Marine sediment cores from tiered silled basins provide a natural laboratory to constrain timing and implications of oxygenation changes across multiple depths. Here, we reconstruct oxygenation and environmental change over time using benthic foraminiferal assemblages from sediment cores from three basins across the Southern California Borderlands: Tanner Basin (EW9504-09PC, 1194 m water depth), San Nicolas Basin (EW9504-08PC, 1442 m), and San Clemente Basin (EW9504-05PC ,1818 m). We utilize indicator taxa, community ecology, and an oxygenation transfer function to reconstruct past oxygenation, and we directly compare reconstructed dissolved oxygen to modern measured dissolved oxygen. We generate new, higher resolution carbon and oxygen isotope records from planktic (Globigerina bulloides) and benthic foraminifera (Cibicides mckannai) from Tanner Basin, Geochemical and assemblage data indicate limited ecological and environmental change through time in each basin across the intervals studied. Early to mid-Holocene (11.0-4.7 ka) oxygenation below 1400 m (San Clemente and San Nicolas) was relatively stable and reduced relative to modern. San Nicolas Basin experienced a multi-centennial oxygenation episode from 4.7-4.3 ka and oxygenation increased in Tanner Basin gradually from 1.7-0.8 ka. Yet across all three depths and time intervals studied, dissolved oxygen is consistently within a range of intermediate hypoxia (0.5-1.5 ml L⁻¹ [O₂]). Variance in reconstructed dissolved oxygen was similar to decadal variance in modern dissolved oxygen and reduced relative to Holocene-scale changes in shallower basins.

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Plain Language Summary

Globally, marine oxygenation is declining with detrimental impacts to ecosystems and economies. To better understand the drivers and consequences of ocean oxygen change, we can examine the fossil record to identify how oxygenation changed in the past. Specifically, we use the relative abundance and chemistry of microfossils (i.e., foraminifera) to reconstruct past oxygenation. Here, we examined microfossils from three sediment cores in three basins (Tanner, San Nicolas, San Clemente) off the coast of Southern California. Marine dissolved oxygen (below 1400 m water depth) was relatively stable and lower than modern from 11,000 to 4,700 years before present. San Nicolas Basin experienced a multicentennial oxygenation episode from 4,700-4,300 years before present and oxygenation increased in Tanner Basin gradually from 2,000-800 years before present. When compared to modern, the range of values of reconstructed oxygen through the entire time studied (thousands of years) is similar to the range of values of modern oxygen at the same depths, indicating that the changes in the last ten thousand years were similar to the amount of change occurring on annual and decadal timescales in the modern ocean.

60	1 Introduction
61	1.1 Marine oxygenation in the past and present
62	At present, global marine oxygenation is declining due to anthropogenic climate change, with important
63	implications for benthic and pelagic ecosystems (Breitburg et al., 2018; Oschlies et al., 2018; Schmidtko
64	et al., 2017). Ocean oxygenation, particularly at depth, is an important driver of ecosystem zonation, and
65	expansion of oxygen minimum zones (OMZ) is a current threat to global marine ecosystems (Breitburg et
66	al., 2018; Helly & Levin, 2004; Stramma, Schmidtko, et al., 2010). Deoxygenation at depth can be driven
67	by several processes: increased export of organic matter from surface to depth leading to increased
68	respiration below the photic zone, increased stratification (often due to surface warming) reducing
69	ventilation at depth, warming surface temperatures reducing rates of diffusion of atmospheric oxygen into
70	surface waters, and changes in the source, current velocity, or oxygenation of intermediate waters (Levin
71	et al., 2009; Oschlies et al., 2018; Stramma, Schmidtko, et al., 2010).
72	
73	Understanding drivers of changes to marine oxygenation in the past is critical for understanding and
74	predicting future change (Jaccard et al., 2014). Paleorecords provide an important archive to investigate
75	temporal and spatial scales of changes to marine oxygenation. In general, warm intervals during the late
76	Quaternary were associated with decreased global ocean oxygenation and cool intervals were associated
77	with increased oxygenation (Cannariato & Kennett, 1999; Cardich et al., 2019; Erdem et al., 2019;
78	Jaccard et al., 2014; Praetorius et al., 2015). The Holocene is an ideal epoch to investigate marine
79	oxygenation because it has well documented intervals of oceanographic change and provides an
80	opportunity to investigate ecological response to stress, including changes to temperature, oxygenation,
81	carbon cycling and ocean circulation (Addison et al., 2017; Barron et al., 2003; Fisler & Hendy, 2008;
82	Moffitt et al., 2015; Praetorius et al., 2015). Over millennial timescales, significant work along the
83	California margin has documented paleoceanographic changes in coastal basins and within the bounds of
84	the modern OMZ (Balestra et al., 2018; Cannariato & Kennett, 1999; Moffitt et al., 2014; Taylor et al.,
85	2015). However, additional work is needed to constrain the timing and extent of oxygenation change
86	below 1000 m, the relative impacts of surface processes and source waters on seafloor oxygenation, and
87	how oxygenation at these depths responds to global and regional environmental change.
88	
89	1.2 Southern California modern and paleoceanography
90	In the modern California Current eastern boundary upwelling system, an OMZ exists at approximately
91	500 - 1000 m water depth and is an important driver of ecosystem zonation, impacting species
92	distributions in pelagic and seafloor communities (Helly & Levin, 2004; Stramma, Johnson, et al., 2010).
93	The combination of high productivity, high export of organic matter, and age of water masses entering the

94 North Pacific at depth make this a particularly thick and laterally extensive OMZ (Bograd et al., 2008, 95 2019; Evans et al., 2020). The California Margin OMZ is expanding in intensity (decreasing dissolved 96 oxygen), horizontal extent, and vertical thickness (shoaling) (Schmidtko et al., 2017; Stramma, Johnson, 97 et al., 2010). Changes to the OMZ are caused by reductions in the oxygenation of source water and 98 reduced ventilation due to surface warming and stratification (Bograd et al., 2008; Evans et al., 2020). Off 99 the coast of California, dissolved oxygen decreased at a rate of up to 0.047 ml/L/year from 1984-2006 in 100 the upper 500 m of the water column (Bograd et al., 2008, 2015). In the Southern California Current 101 System, 81% of observed change in oxygenation in the upper 400 m of the water column from 1993-2018 102 can be attributed to changes in oxygenation of source waters (Evans et al., 2020). 103 104 In the modern system, Southern California Borderlands surface water flow is counterclockwise (Southern 105 California Eddy), with the western margin dominated by the southward flowing California Current and 106 the eastern margin driven by the northward-flowing California Countercurrent and Davidson Current. At 107 depth, the California Undercurrent flows south to north and is comprised of a mix of Southern 108 Component Intermediate Water from the eastern tropical Pacific and northerly sourced North Pacific 109 Intermediate Water (Balestra et al., 2018; Bograd et al., 2019; Checkley & Barth, 2009; Stott et al., 2000; 110 Talley, 1993). Offshore Southern California, a series of submarine basins, generally deepening from north 111 to south, structure intermediate and deep-water flow (Berelson, 1991; Berelson & Stott, 2003). 112 Oxygenation in basinal environments is impacted by the export of organic matter from overlying surface 113 waters, advection of intermediate and deep waters that spill into the basin, and within basin processes, 114 including sediment and pelagic biogeochemical cycles. By examining the environments of multiple silled 115 basins, the effects of water advection can be separated from surface processes and the depths of 116 significant biogeochemical change can be determined. 117 118 The California margin OMZ fluctuated throughout the Holocene (Balestra et al., 2018; Christensen et al., 1994; McGann, 2011; Moffitt et al., 2014; Palmer et al., 2020). Previous analyses of marine sediment 119 120 cores from the Santa Barbara Basin (SBB) document intervals of hypoxia in the early Holocene (11.5-10 ka) followed by oscillations in the strength of the OMZ from 10-6 ka with several intervals of hypoxia 121 122 (less than 0.5 ml L⁻¹ [O₂]) and an increase in oxygenation in the last 6 ka within SBB, yet the OMZ persists throughout the Holocene (Moffitt et al., 2014; Ohkushi et al., 2013; Wang et al., 2020). In Santa 123 Monica Basin (SMB), severe hypoxia (less than 0.3 ml L⁻¹ [O₂]) was present from the start of the 124 Holocene to 9 ka, and the mid to late Holocene (9-0 ka) had weaker hypoxia (0.3-1.5 ml L⁻¹ [O₂]) than the 125 early Holocene (Balestra et al., 2018). The modern OMZ, with oxygen levels at 0-1.5 ml L⁻¹ [O₂], 126 127 developed in the mid to late Holocene (by 6-4 ka) across the broader North Pacific (Addison et al., 2017;

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McGann, 2011; Ohkushi et al., 2013; Praetorius et al., 2015). Over the past several centuries the SMB experienced variable degrees of dysoxia at interannual to interdecadal time scales (Christensen et al., 1994). These changes are attributable to variable biological carbon flux and respiration at depth (Berelson and Stott, 2003; Stott et al, 2000), underscoring how sensitive the shallow-silled basins are to small changes in biological productivity. Previous analysis of ecosystem responses (benthic foraminiferal and invertebrate) to oxygenation change through the Holocene from the SBB indicate that intervals within the Holocene exhibit distinct phases of ecosystems that do not repeat or overlap, as the oxygen minimum zone and carbon maximum zone fluctuate (Moffitt et al., 2015; Myhre et al., 2017). Here, we utilize records from three offshore basins to constrain changes in ocean oxygenation through the Holocene and resultant impacts on benthic ecosystems. Silled basins provide a unique opportunity to examine both local changes within each basin and to compare oxygenation history across depths when records overlap temporally. Combining a series of silled basins allows for the examination of change through time at multiple water depths and sill depths to investigate the relative impact of surface processes and intermediate water changes in determining oxygenation at depth (Balestra et al., 2018; Moffitt et al., 2014; Wang et al., 2020). 1.3 Benthic foraminiferal assemblages as a metric of past oxygenation and organic matter export Benthic foraminiferal assemblages are an effective and established metric to quantify past changes in marine oxygenation (Balestra et al., 2018; Belanger et al., 2020; Bernhard et al., 1997, 2003; Bernhard & Gupta, 1999; Cardich et al., 2015, 2019; Caulle et al., 2014; De & Gupta, 2010; Kaiho, 1994; Moffitt et al., 2014; Murgese & De Deckker, 2005; Ohkushi et al., 2013; Praetorius et al., 2015). Benthic foraminiferal assemblages are sensitive to small changes in oxygenation in the North Pacific, even in suboxic environments, not exclusively across large biological thresholds of anoxic or sulfidic conditions (Sharon et al., 2021). Multiple methodologies are used to interpret past environmental change from benthic foraminiferal assemblages, which we introduce and review below. Quantifying absolute and relative abundance of benthic foraminiferal species is an established and foundational method; typically, studies quantify species downcore and interpret trends through time using observational or statistical approaches (e.g. Moffitt et al., 2014; McGann, 2011; Gardner et al., 1988). Studies of modern benthic foraminifera from multiple depositional environments and oxygenation regimes have identified oxygenation affinity of benthic foraminiferal species that can be used as indicator species of change through time; this method is most useful to reconstruct relative oxygenation or to

identify past oxygenation thresholds (e.g. Cannariato and Kennett, 1999; Palmer et al., 2020; Balestra et 162 163 al., 2018; Bernhard and Gupta, 1999). 164 165 While these approaches provide relative oxygenation history, transfer functions are used to translate 166 whole or partial benthic species assemblages into absolute oxygenation values (Behl & Kennett, 1996; 167 Kaiho, 1994, 1999; McGann, 2011; Moffitt et al., 2014; Ohkushi et al., 2013; Sharon et al., 2021). Multi-168 species transfer functions including the Kaiho Benthic Foraminiferal Oxygenation Index, Schmiedl 169 Dissolved Oxygen Index, and Behl Dissolved Oxygen Index are used to generate absolute values of past 170 oxygenation by transforming the relative abundance of species into ml L⁻¹ [O₂] (Kaiho, 1994; Ohkushi et 171 al., 2013; Sharon et al., 2021). Transfer functions have typically been constructed using case studies from 172 very low oxygen environments (such as Santa Barbara Basin) or by using the lowest known oxygen 173 tolerance for a species; as such, this approach generates inherently conservative (lower) predictions of 174 oxygenation, with the exception of the analysis by Sharon et al., 2021 which utilized multivariate 175 statistical analysis to group species by oxygenation (Kaiho, 1994; Ohkushi et al., 2013; Sharon et al., 176 2021). Categorization of species into oxic, intermediate hypoxic, and anoxic environments used in 177 multiple transfer functions and as indicator species varies by author (Bernhard et al., 1997; Bernhard & Gupta, 1999; Cannariato & Kennett, 1999; Kaiho, 1994; Moffitt et al., 2014; Palmer et al., 2020; 178 179 Praetorius et al., 2015). Here we follow the convention: weakly suboxic/oxic ([O₂] >1.5 ml L⁻¹), 180 intermediate hypoxic/suboxic ($[O_2]$ 1.5-0.5 ml L⁻¹), and severe hypoxic/dysoxic ($[O_2]$ <0.5 ml L⁻¹) 181 (Moffitt et al., 2014; Palmer et al., 2020; Sharon et al., 2021; Tetard et al., 2021). 182 183 Recent work on paleoecological assemblages, including benthic foraminifera, utilizing analysis of 184 diversity, richness, and multidimensional statistical analysis has expanded our breadth of understanding 185 of how oxygenation impacts seafloor ecosystems (Belanger et al., 2020; Myhre et al., 2017; Sharon et al., 2021). Ecological analysis provides community-scale assessments of environmental change over time and 186 187 often complements analysis of indicator taxa or transfer function calculations (Belanger et al., 2020; 188 Myhre et al., 2017; Sharon et al., 2021). In addition to taxonomic evaluation of benthic foraminifera, 189 studies are increasingly relying on morphometrics to assess past environments. These studies are 190 predicated on morphological response or adaptation to oxygenation such as pore density, size, and shape 191 (i.e., roundedness) and are used in both taxon-specific and taxon-independent analyses (Keating-Bitonti & 192 Payne, 2016, 2018; Keating-Bitonti & Payne, 2017; Rathburn et al., 2018; Tetard et al., 2021). Typically, 193 smaller, thin-walled, elongate species are indicative of low oxygen environments in which high-surface 194 area to volume ratio is advantageous, to maximize oxygen absorption, while in well oxygenated

195 environments, larger, thick-walled and porcelaneous taxa with rounded shapes are dominant (Tetard et al., 196 2021). 197 198 Importantly though, oxygenation is not the only driver of benthic foraminiferal assemblages. Food 199 availability, driven by the timing (pulsed vs. constant) and amount of export of organic matter, water 200 depth, sediment grain size, and water temperature play an important role in structuring seafloor 201 ecosystems, including benthic foraminiferal assemblages (Belanger et al., 2012, 2020; Kaiho, 1999; 202 Venturelli et al., 2018). Here we employ stable isotope and benthic foraminiferal assemblage analyses to 203 investigate cores from three silled basins in the Southern California Borderlands. By combining multiple 204 approaches to assemblage analysis as listed above, we quantify changes in oxygenation below 1000 m, 205 deconstruct surface vs. advective processes as drivers of change, and investigate changes in source waters. This multi-site, multi-faceted approach allows us to reconstruct change over time in each basin 206 207 and to assess regional scale oxygenation history and environmental change in the Southern California 208 Borderlands through the Holocene. 209 210 2 Methods 211 2.1 Study site 212 Cores were collected from three silled basins within the Southern California Borderlands (Figure 1). 213 Tanner Basin is located west of the Channel Islands and is the farthest offshore. The basin has a sill depth 214 of 1160 m and a bottom depth of approximately 1500 m (Figure 1). San Nicolas Basin is located south of 215 San Nicolas Island and west of San Clemente Island. The basin has a sill depth of 1100 m and a bottom 216 depth of approximately 1800 m (Figure 1). San Clemente Basin is located south of San Clemente Island 217 and is the deepest site explored here. The basin has a sill depth of 1815 m and a bottom depth of 218 approximately 1950 m (Figure 1). The water depth of each core was assessed at time of core collection 219 and bottom depths of each basin were measured using GeoMapApp.

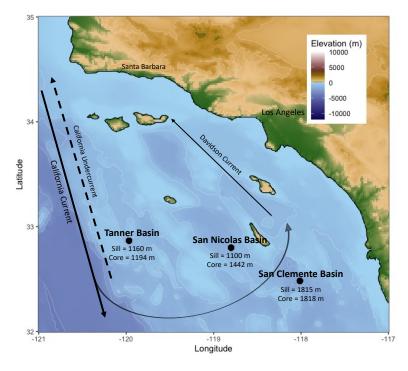


Figure 1: Map of the locations of the three cores used in this study. Schematic of current locations and directions included, solid lines indicate surface currents (California Current and Davidson Current), dashed line indicates subsurface current (California Undercurrent). The sill depth and core collection depth for each core is reported in meters.

2.2 Sediment cores

We investigate three piston cores collected from the Southern California Borderlands on the Maurice Ewing Cruise 9504 in May - June 1995: EW 9504-09PC from Tanner Basin at 1194 m water depth, EW 9504-08PC from San Nicolas Basin at 1442 m water depth, and EW 9504-05 from San Clemente Basin at 1818 m water depth. All cores have 10.16 cm inner diameter and were split and sampled at 2 cm intervals aboard the ship (initial volume of all samples was 162.15 cm³). Sediments from the working half of the core were disaggregated in sodium hexametaphosphate washed over a 63 µm sieve (see Stott et al. 2000). Sediments were dried and stored in glass vials at the University of Southern California until they were processed for this study. The core from Tanner Basin (EW 9504-09PC) was examined from 0-40 cm at 2 cm intervals; intervals below 40 cm were not available for analysis (total of 19 intervals examined). The core from San Nicolas Basin (EW 9504-08PC) was examined at 2 cm intervals from 0-22 cm and every 6 cm from 24-64 cm (due to sample availability, total of 19 intervals examined). The core from San Clemente Basin (EW 9504-05) was examined at 2 cm intervals from 30-52 cm; intervals from 0-30 cm in EW9504-05 did not have sufficient foraminifera for robust analysis (see below) and were thus excluded from analysis (total of 11 intervals examined). Analysis was conducted at the resolution available to the

241 authors and all intervals with sufficient sediment volume and foraminifera (see below) were used in the 242 study. 243 244 2.3 Radiocarbon dating and age model development 245 Radiocarbon based age models were developed for each core using a combination of previously published 246 and newly generated planktic radiocarbon ages (Table S1, Figure S1). Five age dates within the Holocene 247 (1 San Clemente, 1 Tanner, 2 San Nicolas) were previously measured (Stott et al., 2000). These AMS ¹⁴C 248 ages were completed using bulk planktonic foraminifera (weight ~3-5 mg) analyzed at the Lawrence 249 Livermore National Laboratory (Stott et al., 2000). Three additional radiocarbon dates (1 San Clemente, 2 250 Tanner, 1 San Nicolas) from bulk planktic foraminifera were analyzed in this study. All samples for 251 radiocarbon analysis were prepared by picking shell material from the >150 µm fraction, rinsing shells in 252 methanol, sonicating in methanol for 5-10 seconds, and rinsing twice with deionized water. Shells were then dried in a 60°C drying oven. Radiocarbon analysis was completed at the Lawrence Livermore 253 254 National Laboratory using δ^{13} C assumed values following the convention of Stuiver and Polach (1977). 255 The reported age is given in radiocarbon years using the Libby half-life of 5568 years. Accelerator mass 256 spectrometry ages were converted to calendar ages before present (BP) by calibration against the 257 Marine20 curve using the open-source software "R" package Bchron (Haslett & Parnell, 2008; Heaton et 258 al., 2020). Calibration included correction for reservoir ages for the Southern California Coast of 220.0 \pm 259 40.0 years (Ingram & Southon, 1996; Stuiver & Polach, 1977). Age/depth models for each core were 260 generated using the Bayesian age-depth modeling functionality of Bchron (Table S1, Figure S1). 261 262 2.4 Stable Isotope Analysis 263 Stable isotope analyses from planktic and benthic foraminifera from EW9504-09 (Tanner Basin) were 264 conducted on Globigerina bulloides planktic foraminifera and Cibicides mckannai benthic foraminifera from 0-40 cm at 2 cm intervals. Samples were prepared by picking from the >150 µm size fraction (2-5 265 individual C. mckannai per interval, 15-25 G. bulloides per interval) and cleaned using the same 266 267 methodology as above. Planktic foraminifera were analyzed at the UC Davis Stable Isotope Laboratory 268 and benthic foraminifera were analyzed at the UC Santa Cruz Stable Isotope Laboratory. 269 270 Planktic carbon and oxygen isotope samples were analyzed using a GasBench II system interfaced with a 271 Delta V Plus Isotope Ratio Mass Spectrometer at the UC Davis Stable Isotope Laboratory using standard UCD-SM92 (-1.94 for δ^{18} O and 2.08% δ^{13} C) (Ostermann 2000). Values for δ^{13} C and δ^{18} O are expressed 272 in per mil (%) relative to Vienna Pee Dee Belemnite, and values are corrected for changes in linearity and 273 274 instrumental drift. Benthic carbon and oxygen isotope samples were analyzed at the UC Santa Cruz Stable

Isotope Laboratory by acid digestion using an individual vial acid drop Themo Scientific Kiel IV
carbonate device interfaced to Thermo Scientific MAT 253 dual-inlet isotope ratio mass spectrometer. All
samples were measured with several replicates of the externally calibrated Carrera Marble in-house
standard reference material 'CM12' and the NBS-18 limestone international standard reference material.
Values for δ^{13} C and δ^{18} O are expressed in per mil (‰) relative to Vienna Pee Dee Belemnite, and values
are corrected for changes in linearity and instrumental drift. Data were combined with previously
published records of planktic oxygen isotopes and benthic oxygen and carbon isotopes from the same
core examined here (EW9504-09) to increase replicates in the late Holocene and to extend the record
through the entire Holocene (Stott et al., 2000). As such, isotope records reflect analyses from three
different laboratories. Instrument precision from all labs for $\delta^{13}C$ calcite was 0.05-0.06‰ and for $\delta^{18}O$
calcite was 0.06-0.10‰. Thus, we report the maximum uncertainty of 0.06‰ for $\delta^{13}C$ calcite and 0.10‰
for δ^{18} O calcite (Figure 3).
2.5 Benthic Foraminiferal Assemblages
Samples were dry sieved over a 150 µm sieve and picked for benthic foraminifera. Foraminifera greater
than 150 µm have been documented to capture the range of environmental variability in
paleoceanographic reconstructions as well as assemblages containing smaller foraminifera and the use of
larger specimens results in reduced error in identification (Cannariato & Kennett, 1999; Caulle et al.,
2014; Fenton et al., 2018; Palmer et al., 2020). Foraminifera in the size fraction below 150 μm were
excluded from this analysis which may have excluded some smaller taxa or smaller individuals of the taxa
identified here. Individual foraminifera were picked and identified from each interval to obtain >90
individuals per sample (Kemp et al., 2020), as sample sizes as low as 58 have been shown to have stable
assemblages (Belanger et al., 2020; Forcino et al., 2015). The average number of individual foraminifera
identified in each interval was 230, with a range of 90-665. Samples were mounted on micropaleontology
slides using gum tragacanth at time of identification and are archived in the Ocean Climate Laboratory at
the UC Davis Bodega Marine Laboratory. Cores were not laminated; thus, bioturbation is expected and
may have had an averaging effect on assemblages.
2.6 Foraminiferal morphometrics
Morphometrics of benthic foraminifera, including length, width, and surface area were measured using
ImageJ software. Length and width of each individual was measured to the longest and widest margins of
the shell. Three species were selected for analysis to represent a gradient of oxygenation affinity:
Quinqueloculina sp. is weakly suboxic/oxic and B. spissa and U. peregrina are categorized as suboxic
(Table S2). Measurements of <i>Quinqueloculina</i> sp. and <i>B. spissa</i> were quantified in Tanner Basin at 2 cm
(1 auto 52). We asurements of <i>Quinqueiocutina</i> sp. and D. spissa were quantified in 1 aimer Basin at 2 cm

309	intervals from 0-40 cm, measurements of Quinqueloculina sp. and U. peregrina were quantified in San
310	Clemente Basin at 2 cm intervals from 30-44 cm. From each interval, 3-30 individual shells of each
311	species were measured (based on availability of taxa). Results of morphometric analysis were compared
312	to morphometric data (collected using the same methodology) from five open coastal margin sites (300-
313	1175m) near San Diego (see Palmer et al., 2020 for further explanation of the open margin study site, all
314	morphometric data is available on Dryad, see Palmer et al., 2022).
315	
316	2.7 Metazoan microfossil assemblages
317	In addition to picking and identifying benthic foraminifera, ostracods (Arthropoda - Ostracoda) and
318	urchin spines (Echinodermata) were picked from the 150 µm sediment fraction at all intervals examined.
319	Samples were mounted on micropaleontology slides using gum tragacanth at time of identification and
320	are archived in the Ocean Climate Laboratory at the UC Davis Bodega Marine Laboratory. Due to low
321	abundances, presence/absence is recorded, rather than relative abundance.
322	
323	2.8 Benthic foraminiferal oxygenation index
324	Oxygenation was reconstructed using two modified benthic foraminiferal oxygenation indices: Behl and
325	Schmiedl (Schmiedl et al., 2003; Sharon et al., 2021). Data were analyzed using both indices in order to
326	compare the results; the Schmiedl Index uses diversity as a input, but does not include suboxic taxa, while
327	the Behl Index does not include diversity, but includes oxic, suboxic, and weakly suboxic/dysoxic taxa
328	(Ohkushi et al., 2013; Schmiedl et al., 2004). Using a combination of paleo and modern samples, Sharon
329	et al., 2021 used detrended correspondence analysis (DCA) to expand the list of species that can be used
330	as inputs to the Behl Index, thus allowing for its application in a broader range of seafloor environments
331	(Sharon et al., 2021). Here we utilize the Behl dissolved oxygen index following Ohkushi et al., 2013, but
332	expand the list of species included in the assessment following the work of Sharon et al., 2021. We
333	further modified the list of species used in the Behl and Schmiedl indices by adding ten additional species
334	using previously published oxygenation affiliations and morphometric or taxonomic similarities (Table
335	S2). We included all species that made up at least 2% of the total foraminifera identified across all time
336	intervals and cores examined. Oxygenation reconstructions were calculated using two equations:
337	Behl DO index = $((dysoxic \% * 0.1) + (suboxic \% * 0.5) + (weakly suboxic/oxic \% * 1.5))/100$ and
338	$Schmiedl\ index = ((weakly\ suboxic/oxic\ \%)/(weakly\ suboxic/oxic\ \% + hypoxic\ \%) + diversity\ (H`))*0.5$
339	(Ohkushi et al., 2013).
340	

2.9 Statistical analysis

342	Diversity of each interval was calculated using Shannon Index (H). Richness was calculated by tabulating
343	the number of distinct species present in an interval. Non-metric multidimensional scaling (NMDS)
344	ordination, using square root transformation of assemblage species counts and Bray-Curtis similarities
345	and detrended correspondence analysis were conducted and compared (Figure S2, Table S3, S4). We used
346	a single factor ANOVA to determine if there were significant differences among mean morphometrics
347	(length, width, surface area), diversity between basins, diversity through time, and values of reconstructed
348	vs. modern oxygenation. If the results of ANOVA were significant, we performed Tukey's Test to
349	determine where differences in the means occurred. All statistical analyses were completed using the
350	Vegan R package (nmds function, decorana function) or base R functions (tukeyHSD function and
351	res.aov function) (Oksanen et al., 2013; R Core Team, 2021).
352	
353	2.10 Modern oxygenation data
354	Modern oxygen data were sourced from the California Cooperative Oceanic Fisheries Investigations
355	(CalCOFI) for the years 1949-2019. Data were included from all sites in the CalCOFI sampling grid
356	bounded by -116° $-$ -121° W longitude and 32° $-$ 34.5° N latitude (Point Conception is northern
357	boundary)) and depths 1000-2000 m to maximize data availability. Oxygenation was calculated using
358	Winkler Titration of bottle samples at the Scripps Institution of Oceanography (Bograd et al., 2003;
359	Bograd & Lynn, 2003). Flagged data from CalCOFI and property-property and time series analysis
360	excluded 19 data points. The CalCOFI data set used here includes a total of 272 modern oxygen
361	measurements.
362	
363	3 Results
364	3.1 Age Model Development
365	Age models produced a median age and probability distribution (at a range of 95.0% uncertainty, 2σ) at
366	each depth (Figure S1). Age model uncertainty is reported in Table S2. Coherence of age model within
367	each core and between cores supports use of this age model (Figure S1). Mean sedimentation rates for the
368	entire Holocene are similar for all three cores: Tanner has a mean sedimentation rate of 10.8 (\pm 1 σ =5.3
369	cm/ka), San Nicolas has a mean sedimentation rate of 9.9 (\pm 1 σ =1.8 cm/ka), and San Clemente has a
370	mean sedimentation rate of 12.0 (\pm 1 σ =4.8 cm/ka). Sedimentation rates reported here are similar to those
371	observed offshore Central California through the Holocene (McGann, 2011). All following results are
372	discussed as age in thousands of years before present (ka).
373	
374	3.2 Benthic foraminiferal assemblages

375 Benthic foraminiferal assemblages were quantified down core for each core in which sufficient benthic 376 foraminifera were present (Tanner Basin 0-40 cm at 2 cm intervals, San Nicolas Basin 0-22 cm at 2 cm 377 intervals, 24-66 at 6 cm intervals, San Clemente Basin 30-54 cm at 2 cm intervals). The total number of 378 species (richness) in each sample ranged from 15 to 24 (Figure S4) and species diversity (Shannon Index, 379 H) ranged from 2.12 to 2.68, with a mean of 2.40 ($\pm 1\sigma = 0.15$) across all three basins (Figure S4). 380 Species diversity (H) is significantly higher in Tanner Basin (mean=2.51, $\pm 1\sigma = 0.11$), relative to San 381 Nicolas Basin (mean=2.30, $\pm 1\sigma = 0.11$) and San Clemente Basin (mean=2.37, $\pm 1\sigma = 0.10$) (ANOVA, 382 Tukey Test p<0.05) (Figure 3). Comparison of diversity through time in individual basins using 0.5 or 1 383 ka time bins shows that diversity (H) is not statistically different through time in Tanner, San Nicolas, and 384 San Clemente basins (ANOVA, Tukey Test p<0.05 for all sites). Multivariate statistical analysis using 385 nonmetric multidimensional scaling (NMDS) and detrended correspondence analysis (DCA) of down-386 core assemblages shows that, through time, assemblage similarity within sites exceeds similarity to 387 assemblages at any other site (Figure S2). Comparison of results of DCA and NMDS with previously published benthic foraminiferal species categorization by oxygenation do not clearly indicate that 388 389 oxygenation is the dominant factor in controlling species assemblage at the community level. 390 391 In Tanner Basin, the most common species (in descending order) are B. spissa, G. subglobosa, H. 392 soldanii, Quinqueloculina sp., and P. murrhina (Figure 2). Species diversity (H) ranged from 2.32-2.68 393 with a mean of 2.51 ($\pm 1\sigma = 0.11$) (Figure 3). Benthic foraminiferal assemblage gradually changed from 394 5.5 - 1.9 ka; this shift is largely driven by a gradual increase in *Hansenisca soldanii* (previously 395 Gyroidina soldanii) and decrease in B. spissa and U. peregrina (Figure 2). G. subglobosa and 396 Ouinqueloculina sp consistently make up 10-20% of the assemblage through time (Figure 2). In San 397 Nicolas Basin, the most common species (in descending order) are *U. peregrina*, *E. pacifica*, *C.* 398 mckannai, B. spissa, and Uvigerina sp (Figure 2). Species diversity (H) ranges from 2.12-2.56 with a 399 mean of 2.30 (± 0.11) (Figure 3). The assemblage in San Nicolas Basin exhibits little change through time 400 from 10.1-4.7 ka when *U. peregrina* makes up 15-45% of the assemblage. From 4.7-4.2 ka, there is a 401 sharp decline in *U. peregrina* and a sharp increase in *C. mckannai* (Figure 2). From 4.0-1.9 ka, the trend 402 reverses, and the assemblage is very similar to the assemblage from 10.1-4.7 ka. In San Clemente Basin, 403 the most common species (in descending order) are *U. peregrina*, *C. mckannai*, *E. pacifica*, 404 Ouinqueloculina sp., and Uvigerina sp (Figure 2). Species diversity (H) ranges from 2.17-2.51 with a mean of 2.37 (± 0.10) (Figure 3). In San Clemente Basin, relative abundance of benthic foraminifera 405 406 exhibits little change over time from 11.2-7.1 ka as shown by both individual species analysis and DCA 407 (Figure S2).

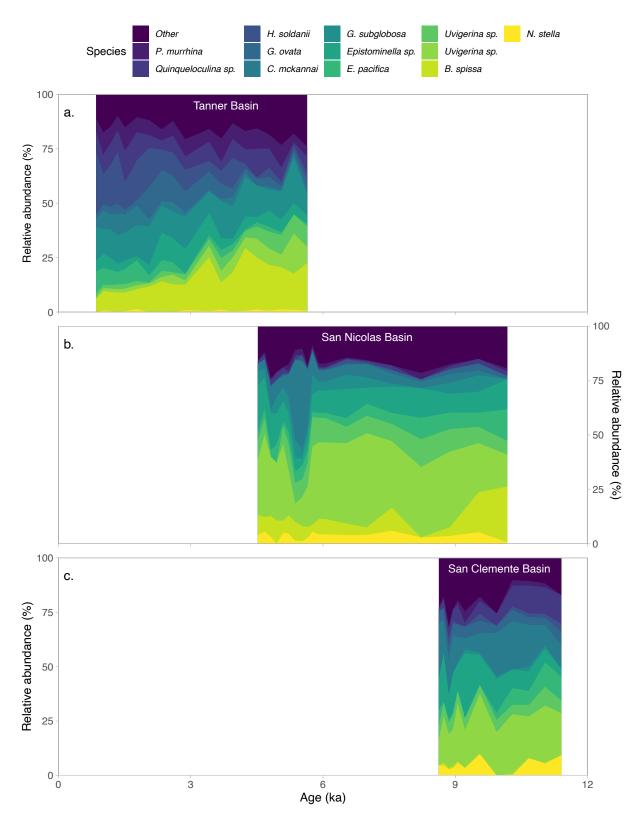


Figure 2: Relative abundance of benthic foraminifera vs. age in thousands of years before present for three cores examined here: Tanner (a.), San Nicolas (b.), and San Clemente (c.). Ten most abundant

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411	species are shown; all other species grouped as "other." Each color represents a taxonomic group
412	(species, genus, or other). See legend at top of plot for color of each. Taxa colors are ordered from
413	dysoxic indicator taxa (yellow/bottom of plot) to oxic indicator taxa (blue/top of plot).
414	
415	3.3 Shell morphometrics
416	Shell size was quantified to examine sub-lethal impacts of environmental change on benthic fauna. Shell
417	size, measured as length, width, and surface area of three taxa (<i>U. peregrina, Quinqueloculina</i> sp., and <i>B.</i>
418	spissa) was compared across the three cores examined here and to size data from a suite of cores from the
419	San Diego Margin (Palmer et al., 2020). Shell size of <i>U. peregrina</i> is statistically significantly larger in
420	coastal sites (p<0.05) relative to San Clemente basin (Figure S3a). Shell size of B. spissa is statistically
421	significantly larger in coastal sites (p<0.05) relative to Tanner Basin (Figure S3a). Shell size of B. spissa
422	and Quinqueloculina from Tanner Basin show relatively little variability through time (5.5-1.9 ka)
423	(Figure S3b, c). As all metrics of shell size (length, width, and surface area) show the same trends, further
424	discussion uses only shell surface area as a metric of whole shell size.
425	
426	3.4 Reconstructed oxygenation using transfer functions
427	Reconstructed dissolved oxygen using the Behl Index ranges from 0.56-1.12 ml L ⁻¹ with a mean of 0.79
428	ml L ⁻¹ (Figure S4) and varies across basins: Tanner Basin mean is 0.86 ml L ⁻¹ and ranges from 0.59-1.12
429	ml L ⁻¹ , San Nicolas Basin mean is 0.67 ml L ⁻¹ and ranges from 0.56-1.02 ml L ⁻¹ , and San Clemente Basin
430	mean is 0.84 ml L ⁻¹ and ranges from 0.68-0.96 ml L ⁻¹ . Reconstructed dissolved oxygen using the
431	Schmiedl Index ranges from 1.36-1.81 ml L ⁻¹ for all three cores, with a mean of 1.62 ml L ⁻¹ (Figure S4).
432	Schmiedl Index reconstructed oxygenation varies across basins: Tanner Basin mean is 1.73 ml L ⁻¹ and
433	ranges from 1.59-1.81 ml L ⁻¹ , San Nicolas Basin mean is 1.50 ml L ⁻¹ and ranges from 1.36-1.66 ml L ⁻¹ ,
434	and San Clemente Basin mean is $1.60 \text{ ml } L^{\text{-1}}$ and ranges from $1.48\text{-}1.71 \text{ ml } L^{\text{-1}}$. Importantly, the number
435	of individual foraminifera identified did not impact the outcomes of the Behl or Schmiedl indices (Figure
436	S4). Species richness is not correlated with outputs of the Behl Index, but is positively correlated with the
437	Schmiedl Index (Figure S4).
438	
439	3.5 Stable isotope record
440	Stable isotope records from Tanner Basin varied slightly through time (Figure 3). Analysis of planktic
441	oxygen and carbon isotopes from G. bulloides yielded the following: δ^{13} C mean is -0.59 (± 1σ = 0.25),
442	range is -0.92 to -0.05, $\delta^{18}O$ mean is 0.22 (± 1σ = 0.31), range is -0.15 to 1.08 (Figure 3). Analysis of
443	benthic oxygen and carbon isotopes from C. mckannai yielded the following: δ^{13} C mean is -0.03 (± 1 σ
444	=0.07), range is -0.13 to 0.10, δ^{18} O mean is 2.53 (± 1 σ = 0.17), range is 2.29 to 2.73 (Figure 3).

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3.6 Metazoan microfossil analysis

Urchin spines and ostracods were present in the Tanner Basin core, but do not present a coeval trend; urchins are present in 13 of 19 intervals, ostracods are present in 5 of 19 samples. Urchin spines and ostracods were scarce in sediments from San Nicolas Basin: urchin spines are present in 6 of 17 and ostracods are present in 2 of 17 intervals. Ostracods and urchin spines were nearly ubiquitous (present in every interval except one) in San Clemente Basin (Figure S5).

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3.7 Modern oxygenation in the Southern California Borderlands

Modern oxygenation decreases linearly with depth from 1000 to 2000 m in this region, and ranges from 0.15-1.67 ml L⁻¹ and has a mean of 0.78 ml L⁻¹ across all depths (1000-2000m). Modern oxygenation in the 50 m above and below the Tanner Basin site (1144-1244 m) has a mean of 0.77 ± 0.18 ml L⁻¹. Modern oxygenation in the 50 m above and below the San Nicolas Basin site (1392-1492 m) has a mean of $0.88 \pm$ 0.37 ml L⁻¹. Modern oxygenation in the 50 m above and below the San Nicolas Basin site (1392-1492 m)

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	Tanner Ba	sin	San Nicola	s Basin	San Cleme	ente Basin
	Mean	Range	Mean	Range	Mean	Range
Behl Index reconstructed oxygen (ml L ⁻¹)	0.86	0.59-1.12	0.67	0.56-1.02	0.84	0.68-0.96
Schmiedl Index reconstructed oxygen (ml L ⁻¹)	1.73	1.59-1.81	1.50	1.36-1.66	1.60	1.48-1.71
Modern dissolved oxygen (ml L ⁻¹)	0.77	0.31-1.45	0.88	0.28-1.39	1.51	1.39-1.61

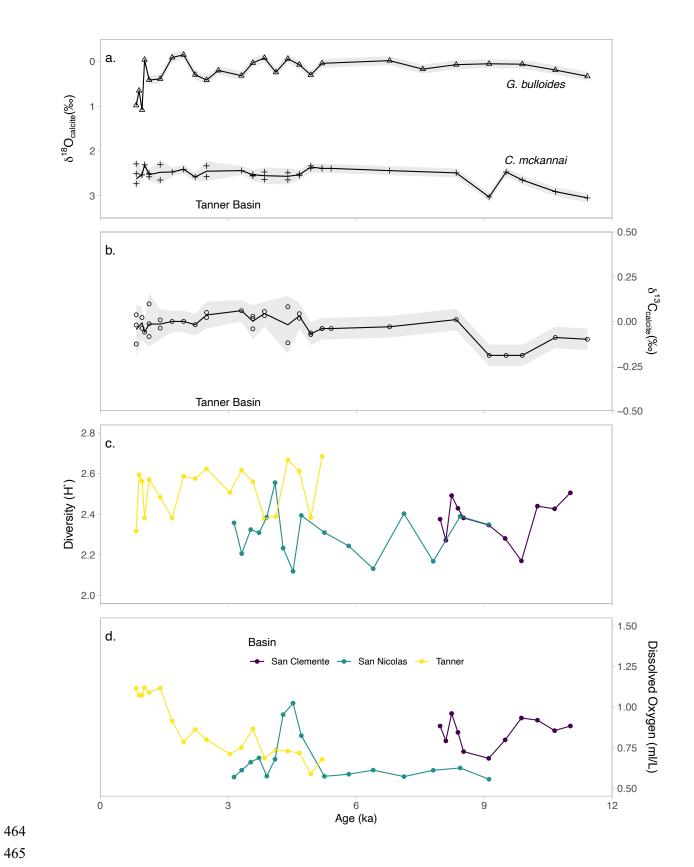
Table 1: Reconstructed and modern dissolved oxygen values for Tanner, San Nicolas, and San Clemente

Basins. Modern oxygenation is from CalCOFI data from 50 m above and below the water depth of core

collection for each basin. 462

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466	Figure 3: Stable isotope record from Tanner Basin (a., b.), benthic foraminiferal diversity (c.), and
467	reconstructed oxygenation (d.) for all three sites using Behl Dissolved Oxygen Index. Oxygen isotopes of
468	planktic foraminifera G. bulloides from this study and Stott et al., 2000 indicated as triangles (a.). Oxygen
469	and carbon isotopes of benthic foraminifera C. mckannai from this study and Stott et al., 2000 indicated
470	as + (b.). Analytical uncertainty for stable isotopes is shown as gray bar ($\delta^{13}C \pm 0.06\%$ and $\delta^{18}O \pm 0.10\%$).
471	In the isotope figures, data from 3.8-0.8 ka includes both data from Stott et al., 2000 and this study, data
472	from 11.4-3.8 ka is from Stott et al., 2000. For diversity and reconstructed dissolved oxygen yellow
473	indicates Tanner, green indicates San Nicolas, and purple indicated San Clemente). Oxygen is reported in
474	ml L ⁻¹ . Discussion
475	4.1 Paleo-oxygenation reconstruction
476	By analyzing the complete assemblage from each basin, we interpret a gradual shift in relative abundance
477	of taxa in Tanner Basin, multi-centennial shifts in the assemblage at San Nicolas Basin, and stability in
478	the assemblage in San Clemente Basin. Examining indicator taxa, we identify a decrease in suboxic taxon
479	B. spissa and a decrease in U. peregrina coeval with an increase in oxic indicator Hansenisca sp. in
480	Tanner Basin from 1.7 to 0.8 ka. In San Nicolas Basin, we observe a short period of divergence from the
481	mean assemblage driven by an increase in the oxic-associated taxon C. mckannai and a sharp decline in
482	U. peregrina from 4.7-4.3 ka. We observe little change over time in the relative abundance of these
483	indicator taxa in San Clemente Basin. We do not identify a relationship between NMDS and DCA species
484	scores and previously published categories of oxygenation indicating that oxygenation may not be the
485	dominant factor in determining the full assemblage. Other parameters such as proximity to the margin of
486	the OMZ, sediment grain size, and organic matter availability also play important roles in structuring
487	foraminiferal diversity and assemblage (e.g., Bernhard et al., 1997; Sharon et al., 2021; Venturelli et al.,
488	2018).
489	
490	Reconstructed oxygen concentration differs between the two transfer functions compared here.
491	Reconstructed dissolved oxygen using Behl Index for all three cores yields values (0.56-1.12 ml L ⁻¹)
492	within "intermediate hypoxia" as defined by Moffitt et al., and "suboxic" as defined by Cannariato and
493	Kennett and Kaiho (Cannariato & Kennett, 1999; Kaiho, 1994; Moffitt et al., 2014). Reconstructed values
494	using the Behl Index are similar to the range of modern values for bottom water oxygenation with notable
495	exceptions discussed below (Figure 3d, 4). In comparison, the Schmiedl Index output is higher (1.36-
496	1.81 ml L ⁻¹) than the Behl Index by approximately 0.5 ml L ⁻¹ (Figure S4). We posit that this is due to the
497	fact that diversity and oxygenation are not necessarily inversely correlated, particularly in intermediate
498	hypoxic environments, yet the Schmiedl Index is based on diversity. Oxygenation plays a more dominant
499	role in structuring diversity across biological thresholds (such as below 0.5 ml L-1 [O ₂]) than across

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oxygen gradients that do not cross biological thresholds (McGann, 2011; Palmer et al., 2020; Sharon et al., 2021; Venturelli et al., 2018). Additionally, the Schmiedl index only considers the end member groups: hypoxic and weakly suboxic/oxic, and does not include the relative abundance of suboxic taxa, taxa which may be important community members and oxygenation indicators at this water depth. Thus, we utilize the Behl Index to reconstruct absolute values of paleo-oxygenation (Figure 3). 4.2 Ecological and environmental change through the Holocene in the Southern California **Borderlands** Geochemical and faunal records from Tanner, San Nicolas, and San Clemente Basins indicate environmental and ecological stability in water below 1000m through the Holocene in the Southern California Borderlands. Planktic δ^{18} O records from Tanner Basin indicate constant sea surface conditions (salinity/temperature) through most of the Holocene (11.75 – 2.3 ka) and a decrease in sea surface temperature or increase in salinity from 1.0-0.8 ka relative to the rest of the Holocene (Figure 3). Benthic δ^{18} O and δ^{13} C of the epibenthic foraminifera C. makannai exhibit little change through the Holocene (Figure 4). Reconstructed seafloor oxygenation using the Behl Index is within the range of intermediate hypoxia $(0.5 - 1.5 \text{ ml L}^{-1})$ and are similar to the range of modern values for bottom water oxygenation with notable exceptions discussed below (Figure 3d, 4). Ecological and environmental change over time as measured by diversity and multi-dimensional community metrics also indicate stability through the Holocene. Diversity varied little between basins and through time within each basin. Shannon Index of diversity (H) ranged from 2.12 to 2.68 across all sites and points in time, and diversity did not significantly change through time at any site across this interval (ANOVA, p>0.05) (Figure 3, S4). Results of NMDS and DCA show species assemblages are distinct in each basin; at any time point, assemblages are more similar to other time points from the same basin than to any other time point from an adjacent basin (Figure S2). This demonstrates that each basinal environment is unique and the analysis of benthic foraminifera as environmental indicators must consider local factors. We hypothesize that the stability in diversity and community-scale ecology indicates that environmental conditions were also relatively stable through this interval. This evidence aligns with results of oxygenation reconstruction from the Behl Index and stable isotope analysis (Figure 3). Changes in benthic foraminifera morphology can reflect environmental variations that are missed by community-based analyses. Comparison of two species (U. peregrina, B. spissa) between basins (studied here) and open margin sediments (San Diego Margin, 300-1200 m water depth see Palmer et al., 2020) shows that both species are larger in the nearshore environment relative to offshore basins (Tanner, San

534 Clemente Basin, see Figure S3). At this scale, we interpret the size difference to be representative of 535 higher relative organic matter input at coastal margin sites or warmer bottom water temperatures at 536 shallower coastal margin sites in comparison to offshore basins (Keating-Bitonti & Payne, 2016, 2018). 537 Importantly though, in comparison to changes between coastal margins and offshore basins, there is 538 relatively little change in shell size of *Ouinqueloculina* sp. (oxic indicator species) or *B. spissa* (suboxic 539 indicator species) over time (5.4-0.8 ka) in Tanner Basin (Figure S3). Thus, benthic foraminiferal δ^{18} O 540 and δ^{13} C, reconstructed dissolved oxygen (Behl Index), community scale ecology, and morphology all indicate relative stability in environmental conditions at water depths 1194-1818 m through the studied 541 542 time interval with few notable exceptions discussed below. 543 544 4.2.1 Ecological and environmental change through time in Tanner Basin 545 In Tanner Basin, the benthic foraminiferal assemblage shifts from more hypoxic taxa at 5.4-1.7 ka to 546 more oxic associated taxa from 1.7-0.8 ka (Figures 2, 3). This shift is largely driven by the increase in 547 Hansenisca soldanii (previously Gyroidina soldanii) and decrease in B. spissa and U. peregrina (Figure 548 2). Hansenisca soldanii has been documented in well oxygenated, oligotrophic, cold deep water with 549 pulsed food supply (De & Gupta, 2010), and both B. spissa and U. peregrina are well documented indicators of low oxygen. Using the modified Behl Index, reconstructed oxygenation increases from 0.58 550 551 to 1.12 ml L⁻¹ across this interval (Figure 3). Thus, we interpret this assemblage shift as an indication of a increase of approximately 0.4 ml L⁻¹ in Tanner Basin occurring around 1.7 ka. 552 553 554 Downcore morphometric analysis shows that neither relative abundance nor surface area of 555 Ouinqueloculina sp. (an oxic indicator species) changes through time. Although B. spissa decreases in 556 relative abundance from 5.4-0.8 ka, shell size (surface area) does not change through time (Figure S3). 557 Stability in shell size concurrent with changes in relative abundance may indicate an interaction of factors including oxygen and organic matter influx in determining lethal vs. sublethal effects on B. spissa. 558 559 Further, the lack of changes in shell size of B. spissa and Quinqueloculina across this interval may 560 indicate a consistent supply of surface-exported organic matter, the main food resource for these benthic 561 foraminifera. Ostracod and echinoderm presence are not correlated with increase in dissolved oxygen at 562 this site (Figure S5). We hypothesize that the shift identified in the benthic foraminiferal assemblage is in response to a relatively small change in oxygenation (less than 1 ml L⁻¹) and does not represent 563 fluctuations across biologically important thresholds for metazoan taxa, as dissolved oxygen was never 564 below 0.4 ml L⁻¹ (Figure 3). 565 566

567	When compared to modern CalCOFI data, reconstructed oxygenation (using the Behl Index) is within 1σ
568	of modern mean oxygenation (Figure 4), except for the interval from $1.7-0.8$ ka in which reconstructed
569	oxygen is higher than modern by 0.3 ml L ⁻¹ (Figure 4). There is no significant difference (ANOVA,
570	Tukey, p=0.51) between reconstructed oxygenation values (mean=0.86 ml L ⁻¹) and all modern
571	oxygenation values (mean=0.77 ml L ⁻ 1) within waters 50 m above and below the Tanner Basin site
572	(Figure 4). Combining assemblage, oxygenation index, and isotopic records of Tanner Basin, we identify
573	increased dissolved oxygen in the basin from 1.7-0.8 ka relative to 5.4-1.7 ka that is not correlated with
574	changes in either the oxygen or carbon isotope records and remains within the modern range of dissolved
575	oxygen variability. The changes in Tanner Basin dissolved oxygen are minor when compared to changes
576	in oxygenation in nearby more shallow basins within the OMZ (Balestra et al., 2018; Moffitt et al., 2014;
577	Palmer et al., 2020; Wang et al., 2020).

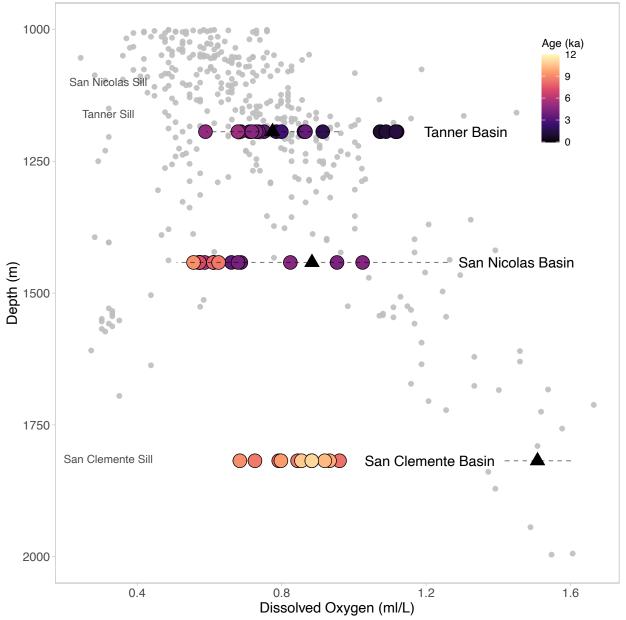


Figure 4: Reconstructed dissolved oxygen concentration (ml L⁻¹) from Tanner, San Nicolas, and San Clemente Basin (large dots colored by age) compared with modern dissolved oxygen concentration (ml L⁻¹) from CalCOFI bottle sampling (1986-2019) (shown in small gray dots). Depth of sill of each basin is labeled. Black triangles show mean modern dissolved oxygen concentration from 50 m above and below each site and dashed gray line shows 1σ range in modern dissolved oxygen concentration from 50 m above and below each site.

4.2.2 Ecological and environmental change through time in San Nicolas Basin

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In San Nicolas Basin, low oxygen-associated benthic foraminiferal taxa dominate from 9.8 to 4.7 ka. A short period of anomalous assemblage driven by an increase in C. mckannai, an oxic-associated taxon, and a sharp decline in *U. peregrina*, an intermediate hypoxia-associated taxon, occurs at 4.7-4.3 ka and is interpreted as an interval of increased oxygenation. Using the modified Behl Index, reconstructed oxygenation increases from 0.59 ml L⁻¹ before the interval to a maximum of 1.03 ml L⁻¹ during the oxygenation interval (Figure 3). Oxygenation returned to pre-event levels of dissolved oxygen after 4.3 ka and pre and post event assemblages are similar, indicating an absence of a legacy effect of the event on the community (Figure 2). Metazoan invertebrate microfossils (ostracods and urchin spines) are scarce in this core and changes in presence/absence of metazoans do not correlate with shifts in benthic foraminiferal assemblages (Figure S5). Modern studies of circulation in San Nicolas Basin indicate that residence time for bottom waters in the basin are 9 ± 2 months (Berelson, 1991). As such, we hypothesize that the assemblage response recorded here represents persistent oxygenation or repeated ventilation on a decadal-to centennial scale, rather than a single episode of ventilation. Yet, the absolute change in oxygenation is relatively minor and still within a threshold of intermediate hypoxia and within the range of modern values (Figure 3, 4). When compared to modern CalCOFI data, reconstructed oxygenation in San Nicolas Basin is slightly lower than modern mean dissolved oxygen yet remains within 1 \u03c4 of the modern mean at all points throughout the Holocene (Figure 4). Reconstructed oxygenation (mean=0.67 ml L⁻¹) is statistically lower than modern oxygenation from waters 50 m above and below the depth of the San Nicolas site (ANOVA, Tukey p<0.05) (Figure 4). Further, the variance in reconstructed dissolved oxygen is similar to variance in modern oxygenation at the sill depth of San Nicolas Basin (1100m, see Figure 4). The scale of change in reconstructed dissolved oxygen, relative abundance of species, and species diversity in San Nicolas Basin across this interval of change is minor when compared to intervals of change in nearby basins across the deglacial and in comparison to Holocene changes within basins that experienced shifts between anoxic and oxic conditions, such as the Macoma Event in the Santa Barbara Basin (Balestra et al., 2018; Cannariato & Kennett, 1999; Moffitt et al., 2015; Praetorius et al., 2015; Schimmelmann et al., 2013). 4.2.3 Ecological and environmental change through time in San Clemente Basin Environmental conditions remained relatively constant in San Clemente Basin from 11.0-8.0 ka as demonstrated by the lack of variability in the benthic foraminiferal assemblages. Species diversity does not change significantly through time (Figure 3, S4). Reconstructed oxygenation from the modified Behl Index ranges from 0.67 to 0.96 ml L⁻¹ across this interval; this variability is reduced relative to Tanner and

San Nicolas Basins (Figure 3). The assemblage at this depth is distinct from coastal margin environments

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and other, shallower basins (Balestra et al., 2018; Moffitt et al., 2014; Palmer et al., 2020). Ostracods and urchin spines are present in every interval (except one) examined here, indicating a well-oxygenated seafloor environment that supported metazoan life and motile organisms (Figure S5) (Moffitt et al., 2015). Ecological differences between the shallow sill basins (San Nicolas and Tanner) vs. the deep sill basin (San Clemente) documented here are mirrored in previously documented seafloor metazoan assemblages (France, 1994). We hypothesize that at 1815 m water depth, this site was not impacted by changes in the oxygen minimum zone during the Holocene and that intermediate and deep water reaching this site had a consistent source from 11.0-8.0 ka. In comparison to modern CalCOFI data, reconstructed dissolved oxygen (mean=0.84 ml L⁻¹) in the early and mid-Holocene is statistically lower than modern dissolved oxygen (mean=1.50 ml L⁻¹) within 50 m above and below the depth of the San Clemente Site (p<0.05) (Figure 4). We propose three potentially overlapping hypotheses for this difference. First, this may be an artifact of categorizations of some species used in the Behl Index that use lowest known thresholds of oxygenation and thus may predict the minima of dissolved oxygen concentrations. Second, in the early Holocene, sea level was lower than modern (by approximately 70 m), thus, the sites sampled here may have been less well oxygenated simply due to being at a shallower depth, but this would not entirely explain the difference from modern (Figure 4) (Fleming et al., 1998; Moffitt et al., 2014). Finally, the preformed oxygenation of the incoming water masses may have been lower than modern, a pattern that is reflected in Santa Monica and Santa Barbara Basins at this time (Balestra et al., 2018; Wang et al., 2020). 4.3 Environmental stability in Southern California Borderlands through the Holocene and comparison to modern ocean conditions Environmental conditions in Tanner, San Nicolas, and San Clemente Basins remained relatively stable across the time intervals examined. In the early to mid-Holocene (11.0-4.7 ka), seafloor oxygenation and environmental conditions more broadly below 1400 m were stable in the Southern CA Borderlands, evidenced by stability in assemblage records from San Clemente Basin (1815 m) and San Nicolas Basin (1442 m), and was likely driven by consistent source and composition of intermediate waters. Benthic foraminiferal assemblage records from the Santa Lucia Slope (ODP 1017, 955 m) offshore Point Conception indicate a stable suboxic environment (reconstructed [O₂] 1.5-0.5 ml L⁻¹) through the early and mid-Holocene and are similar in taxonomic composition to the records presented here from San Nicolas and San Clemente Basins (Cannariato & Kennett, 1999; Sharon & Belanger, 2022). Synchrony in stability and assemblages across San Nicolas Basin (1442 m), San Clemente Basin (1818 m), and the Santa Lucia Slope (955 m) contrasts with evidence of fluctuations in age of North Pacific Intermediate

656 Water entering SBB (as recorded by benthic-planktic radiocarbon age differences) at 9 ka (Roark et al., 657 2003), expansion of the OMZ in SBB in the early Holocene (Wang et al., 2020), and a turning point from 658 hypoxic to suboxic conditions at 9 ka in Santa Monica Basin (indicated by benthic foraminiferal 659 assemblages and geochemical records) (Balestra et al., 2018). Differences in oxygenation between sites 660 may be explained by differences across water depth; we propose that changes in the source and strength 661 of NPIW impacted sites at 400-1000m water depth (SBB, SMB) (Balestra et al., 2018; Roark et al., 2003; 662 Wang et al., 2020), but had less of an effect on the water column below 1000 m. Similar patterns of stable 663 suboxic conditions below the depth of the OMZ and variability in dissolved oxygen at shallower, OMZ-664 impacted depths through the Holocene were observed further north, offshore British Columbia, and have been attributed to changes in the strength of NPIW (Sharon & Belanger, 2022). In the mid to late 665 Holocene, San Nicolas Basin experienced a multi-centennial oxygenation interval from 4.7-4.3 ka and 666 oxygenation increased in Tanner Basin beginning at 1.0 ka, yet all changes were within 1 ml L⁻¹ [O₂]. 667 668 Drivers of oxygenation in each basin may have been due to "indirect" ventilation through diffusion of dissolved oxygen from overlying waters, rather than "direct" ventilation due to advection (Talley, 1993), 669 670 a change in oxygenation of source waters, or basinal processes related to the shape, depth, and overlying 671 surface water productivity in each basin. 672 673 Although oxygenation does vary across the Holocene in Tanner, San Nicolas, and San Clemente basins, all variability was within 1 ml L⁻¹ [O₂] and does not cross critical biological thresholds below 0.5 ml L⁻¹ 674 or above 1.5 ml L⁻¹ in any basin. As such, variability in oxygenation and ventilation of San Nicolas, 675 676 Tanner, and San Clemente basins is reduced relative to shallower sites, including Santa Barbara Basin and 677 Santa Monica Basin, across the entire Holocene (Balestra et al., 2018; Moffitt et al., 2014; Ohkushi et al., 678 2013). Additionally, Holocene-scale oxygenation changes (presented here) are reduced relative to 679 glacial/interglacial changes in oxygenation in the basins examined here and in the nearby Santa Lucia Slope (Cannariato & Kennett, 1999; Stott et al., 2000). This indicates that Holocene-scale climate changes 680 681 driving oxygenation change shallower than 1000 m are not impacting waters below 1000 m and that 682 climate changes within the Holocene do not significantly impact oxygenation below 1000 m, despite 683 changes in the intensity and extent of the OMZ shallower than 1000 m. As such, we hypothesize that 684 deoxygenation due to anthropogenic climate change will also have a greater impact on the water column 685 above 1000 m, relative to below 1000m. 686 687 Direct comparison of reconstructed dissolved oxygen with modern measured dissolved oxygen indicates 688 that the variance across millennia is similar to decadal-scale variance in the modern ocean (Figure 4). 689 Variance in dissolved oxygen occurring on centennial to millennial timescales does not exceed the

variance observed in this modern ocean, indicating stability in intermediate waters and oxygenation below 1100 m in the Southern California Borderlands. We hypothesize that if there are significant changes to oxygenation below 1000m due to anthropogenic climate change, these changes may have large impacts on benthic ecosystems as they have not experienced significant changes in dissolved oxygen over the last 11 ka.

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5 Conclusion

Reconstruction of past oxygenation using analysis of benthic microfaunal communities (foraminiferal and metazoan) is optimized by combining multiple approaches including analysis of indicator taxa, reconstruction of oxygenation using multi-taxa indices, and community scale-analysis such as multidimensional analysis and diversity. Here we demonstrate the utility of combined approaches, and we expand the use of the Behl Index for paleo-oxygenation reconstruction. Analysis of benthic foraminiferal assemblages from three silled basins (Tanner, San Nicolas, San Clemente) in the Southern California Borderlands combined with benthic and planktic stable isotope analysis from Tanner Basin show largely stable oxygenation except for a gradual increase (approximately 0.4 ml L⁻¹ [O₂]) in oxygenation in Tanner Basin occurring at 1.7 ka and multi-centennial variability in oxygenation (approximately 0.5 ml L^{-1} [O₂]) in San Nicolas Basin. The seafloor environment is stable in San Clemente Basin from 11.0-8.0 ka, yet reconstructed oxygenation is lower than modern at this site. Holocene scale climate changes did not drive significant changes (> 1 ml L⁻¹) in marine oxygenation below 1000 m in the Southern California Borderlands. In the context of modern oxygenation changes, findings from this analysis show that seafloor oxygenation of the Southern California Borderlands through the Holocene below 1000 m remained relatively stable and variance in oxygenation across millennia is similar to decadal-scale variance in the modern ocean. As such, we expect that future changes to marine oxygenation will be greater at depths above 1000 m relative to deeper waters and note that if anthropogenic climate change induced changes in oxygenation do cause shifts in dissolved oxygen greater than > 1 ml L⁻¹ below 1000 m, it will represent a divergence from scales of variability over the last 11 ka.

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717	Acknowledgements
718	The authors declare no conflicts of interest. We acknowledge funding for this project provided through
719	National Science Foundation Grant OCE 1832812 to TMH and PDR and the University of California,
720	Davis Dissertation Year Grant and the University of California, Davis Earth and Planetary Sciences
721	Durrell Fund Research Award. We thank Sarah Merolla and Kimberly Bowman for their support with
722	sample preparation and processing. We thank Robin Trayler for his support and advice in developing the
723	age model. We also thank Sharon for their assistance in multivariate statistics. We acknowledge three
724	helpful reviewers whose input significantly improved this paper.
725	
726	Open Research
727	The benthic foraminiferal assemblage data, radiocarbon age model data, carbon and oxygen stable isotope
728	data, and morphometric data used for environmental reconstruction in the study are available at Dryad via
729	Palmer et al. (2022).
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Paleoceanography and Paleoclimatology

Supporting Information for

Ecological and environmental stability in offshore Southern California Marine Basins through the Holocene

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Figures S1 to S5 Tables S1 to S2

Introduction

Supporting information contains five supporting figures and two supporting tables. All methodology is discussed in main text.

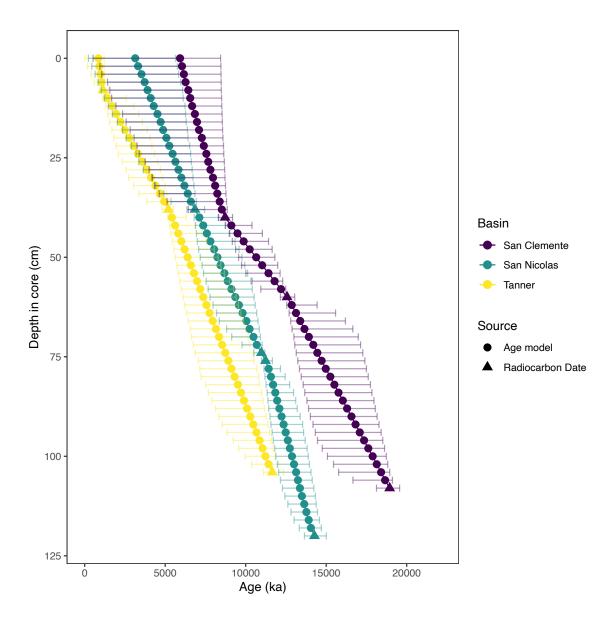


Figure S1. Radiocarbon-based age model as generated through Bchron for three cores included in study. Age shown in thousands of years before present. Triangles are calibrated radiocarbon ages, circles are median ages generated through Bchron age model: Tanner= yellow, San Nicolas = green, San Clemente = purple). Error bars indicate 95% credible interval at each depth.

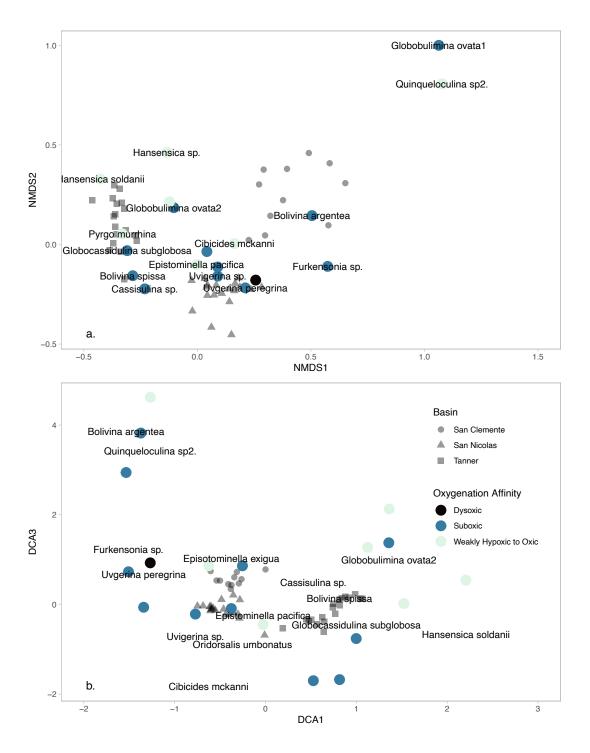


Figure S2. Non-metric multidimensional scaling plot (a.) and detrended correspondence plot (b.) of all benthic foraminiferal assemblages through time from three cores. Axes that represent the most variance are shown here. Species are large circles color coded by published oxygenation affinity (dark blue is dysoxic, blue is suboxic, and teal gray is weakly hypoxic to oxic). All other points represent assemblages at each interval through time at each of the three cores; gray dots are San Clemente assemblage, gray triangles are San Nicolas assemblage, and gray squares are Tann assemblage.

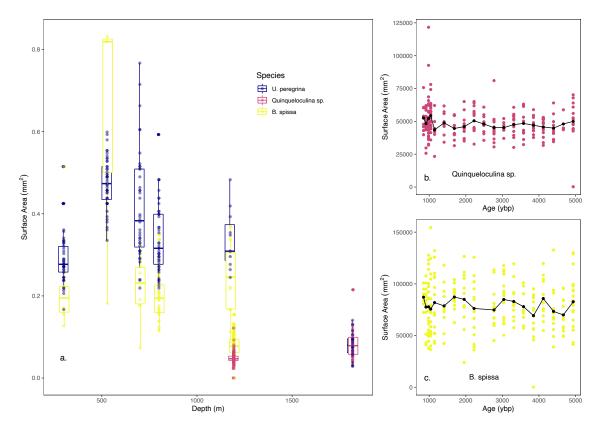


Figure S3. Shell surface area (mm²) vs. water depth (m) at all points in time (a.). Data shown here indicate all points in time from each water depth. Colors represent species: U. peregrina = purple, *Quinqueloculina* sp. = magenta, *B. spissa* = yellow. All points are shown in addition to box plot. Box plot is slightly offset to better display two species at each water depth. Surface area of *Quinqueloculina* sp. (b.) and *B. spissa* (c.) vs. time in years before present from Tanner Basin.

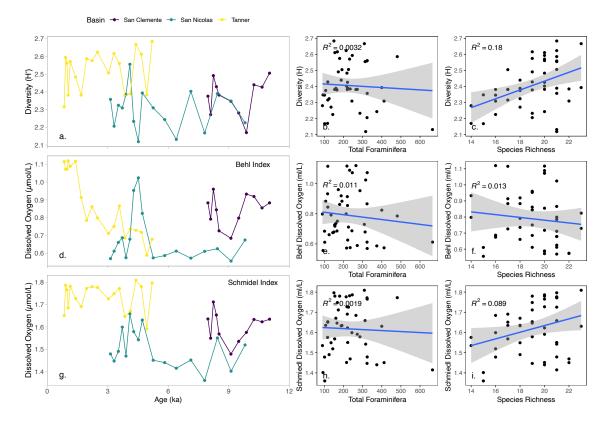


Figure S4. Diversity (Shannon Index), vs. age in thousands of years before present (a.), total foraminifera (b.), and species richness (c.). Reconstructed dissolved oxygen (ml/L) using Behl Index vs. age in thousands of years before present (a.), total foraminifera (b.), and species richness (c.). Reconstructed dissolved oxygen (ml/L) using Schmiedl Index vs. age in thousands of years before present (a.), total foraminifera (b.), and species richness (c.). Trendlines in panels b., c., e., f., h., and i., use linear regression, R² is shown on plot.

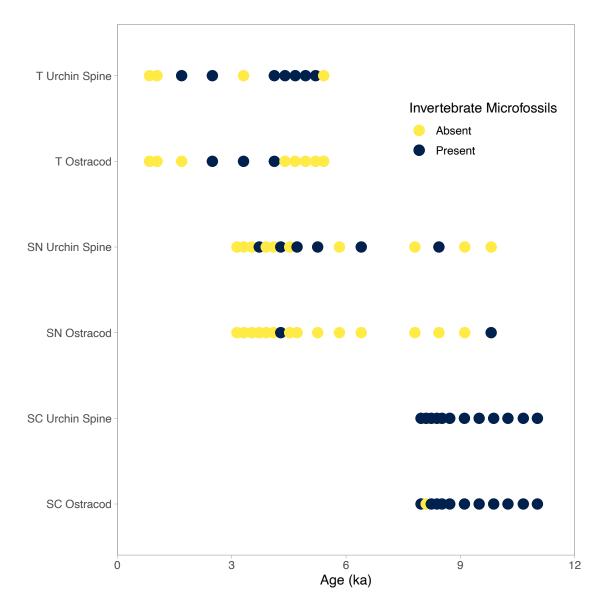


Figure S5: Presence and absence of metazoan microfossils (urchin spines and ostracods) through time (age in thousands of years before present) for each basin (Tanner (T), San Nicolas (SN), and San Clemente (SC)). Presence is blue, absence is yellow.

Basin	Core	Sample Interval	Source	Radiocarbon date (14C years)	±	Calendar age (years before present)	95% Credible Interval (years before present)
Tanner	EW95 04-09	8-10 cm	Stott et al., 2000	1950	50	1146	905-1468
Tanner	EW95 04-09	38- 40 cm	This paper	5290	30	5201	4809-5511
Tanner	EW95 05-09	104-106 cm	Stott et al., 2000	10790	60	11647	11102-12350
San Nicolas	EW95 04-08	38-40 cm	Stott et al., 2000	6780	50	6849	6351-7448
San Nicolas	EW95 04-08	74-76 cm	This paper	10330	35	10976	10511-11190
San Nicolas	EW95 04-08	76-78 cm	Stott et al., 2000	10460	70	11233	10987-11657
San Nicolas	EW95 04-08	120-122 cm	Stott et al., 2000	12870	160	14269	13647-15006
San Clemente	EW95 04-08	40-42 cm	Stott et al., 2000	8550	60	8721	8308-9185
San Clemente	EW95 04-08	60-62 cm	This paper	11430	40	12570	12155-13043
San Clemente	EW95 04-08	108-110 cm	Stott et al., 2000	16650	50	18942	181123-19560

Table S1. Radiocarbon ages and age model for all three cores examined here. Radiocarbon age calibrated using reservoir age of 220 + -40 and Behron Bayesian approach.

	Oxygen	Modified from	
Species	Classification	Sharon-Behl	Citation for oxygenation affinity
Bolivina argentea	Suboxic		Sharon et al., 2021
Bolivina pseudobeyrichi	Dysoxic		Sharon et al., 2021
Bolivina spissa	Suboxic		Sharon et al., 2021
Bulimina tenuata	Dysoxic		Sharon et al., 2021
Cassidulina sp.	Suboxic		Sharon et al., 2021
Chlistomella ovoidea	Suboxic		Sharon et al., 2021
	Weakly hypoxic to		
Cibicides mckannai	oxic	Modified	Kaiho 1994
	Weakly hypoxic to		
Cibicides sp.	oxic	Modified	Kaiho 1994
	Weakly hypoxic to		
Elphidium sp.	oxic		Sharon et al., 2021
Epistominella exigua	Suboxic		Sharon et al., 2021
	Weakly hypoxic to		
Epistominella pacifica	oxic	Modified	Cannariato and Kennett 1999
Fursenkoina sp.	Suboxic	Modified	Kaiho 1994
Globobulimina barbata	Suboxic	Modified	Ohkushi et al., 2013
Globobulimina ovata1	Suboxic	Modified	Ohkushi et al., 2013
Globobulimina ovata2	Suboxic	Modified	Ohkushi et al., 2013
Globobulimina pacifica	Suboxic	Modified	Ohkushi et al., 2013
Globocassidulina sp.	Suboxic		Sharon et al., 2021
Globocassidulina			
subglobosa	Suboxic		Sharon et al., 2021
	Weakly hypoxic to		
Hansenisca soldanii	oxic	Modified	De and Gupta 2010
	Weakly hypoxic to		
Hansenisca sp.	oxic	Modified	De and Gupta 2010
Melonis affinis	Suboxic	Modified	Kaiho 1994
Nonionella stella	Dysoxic		Sharon et al., 2021
Oridorsalis umbonatus	Suboxic		Sharon et al., 2021
Other			
	Weakly hypoxic to		
Pyrgo murrhina	oxic		Sharon et al., 2021
	Weakly hypoxic to		
Quinqueloculina sp.	oxic		Sharon et al., 2021
	Weakly hypoxic to		
Quinqueloculina sp2.	oxic		Sharon et al., 2021
Uvigerina peregrina	Suboxic		Sharon et al., 2021
Uvigerina proboscoidea	Suboxic		Sharon et al., 2021
Uvigerina sp.	Suboxic		Sharon et al., 2021

Table S2. Oxygen affinity for all species examined here and utilized in oxygen transfer functions.

	NMDS1	NMDS2	NMDS3	NMDS4
	(43.0%)	(15.8%)	(3.1%)	(0.8%)
Bolivina spissa	-0.284	-0.157	0.167	-0.042
Cassisulina sp.	-0.232	-0.223	0.080	-0.085
Cibicides mckanni	0.163	0.005	-0.153	0.166
Epistominella pacifica	-0.004	-0.100	-0.046	0.122
Episotominella exigua	0.089	-0.113	-0.040	-0.059
Globobulimina ovata2	-0.104	0.185	0.005	-0.089
Globocassidulina subglobosa	-0.309	-0.031	0.045	0.033
Hansensica soldanii	-0.427	0.329	-0.075	0.076
Oridorsalis umbonatus	0.042	-0.035	-0.123	0.103
Pyrgo murrhina	-0.330	0.053	-0.007	0.049
Quinqueloculina sp.	-0.123	0.214	0.101	-0.041
Uvigerina sp.	0.090	-0.159	-0.068	0.038
Uvgerina peregrina	0.211	-0.218	-0.028	0.004
Hansensica sp.	-0.135	0.463	-0.357	-0.239
Nonionella stella	0.257	-0.179	-0.072	-0.141
Bolivina argentea	0.505	0.145	-0.813	-0.953
Furkensonia sp.	0.574	-0.110	0.239	0.125
Quinqueloculina sp2.	1.077	0.806	0.276	-0.216
Globobulimina ovata1	1.063	1.000	0.611	0.317

Table S3. NMDS species scores. Proportion of variance for each axis is reported in parenthetical. Proportion of variance of each axis was calculated summarized by each axis using the Pearson correlation between the Euclidean dissimilarity of the sites scores and the Bray-Curtis dissimilarity of the faunal abundances for each site with the mantel function in the "ecodist" package (Sharon et al., 2021).

	DCA1	DCA2	DCA3	DCA4
	(45.6%)	(6.2%)	(11.3%)	(2.1%)
Bolivina spissa	0.815	0.132	-1.676	-0.235
Cassisulina sp.	0.527	0.490	-1.701	-0.720
Cibicides mckanni	-0.626	-1.833	0.864	0.333
Epistominella pacifica	-0.026	-0.248	-0.447	1.073
Episotominella exigua	-0.376	1.024	-0.091	-1.480
Globobulimina ovata2	1.358	0.997	1.375	-1.259
Globocassidulina subglobosa	0.999	-0.456	-0.761	-0.038
Hansensica soldanii	2.207	-0.665	0.544	0.706
Oridorsalis umbonatus	-0.253	-0.897	0.861	-0.859
Pyrgo murrhina	1.523	-0.763	0.020	0.498
Quinqueloculina sp.	1.124	0.525	1.273	1.365
Uvigerina sp.	-0.772	-0.713	-0.215	0.231
Uvgerina peregrina	-1.341	0.669	-0.063	0.199
Hansensica sp.	1.362	-0.780	2.129	-2.557
Nonionella stella	-1.269	1.092	0.928	0.723
Bolivina argentea	-1.535	3.864	2.943	1.728
Furkensonia sp.	-1.508	1.231	0.729	1.057
Quinqueloculina sp2.	-1.267	3.421	4.619	2.955
Globobulimina ovata1	-1.372	3.262	3.822	4.412

Table S4. DCA species scores. Proportion of variance for each DCA axis is reported in parenthetical. Proportion of variance of each axis was calculated summarized by each axis using the Pearson correlation between the Euclidean dissimilarity of the DCA sites scores and the Bray-Curtis dissimilarity of the faunal abundances for each site with the mantel function in the "ecodist" package (Sharon et al., 2021).