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- Ocean biogeochemistry in the Canadian Earth System Model version 5.0.3: CanESM5 and
 CanESM5-CanOE
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Abstract. The ocean biogeochemistry components of two new versions of the Canadian Earth 19 System Model are presented and compared to observations and other models. CanESM5 20 employs the same ocean biology model as CanESM2 whereas CanESM5-CanOE ("Canadian 21 Ocean Ecosystem model") is a new, more complex model developed for CMIP6, with multiple 22 food chains, flexible phytoplankton elemental ratios, and a prognostic iron cycle. This new 23 24 model is described in detail and the outputs (distributions of major tracers such as oxygen, dissolved inorganic carbon, and alkalinity, the iron and nitrogen cycles, plankton biomass, and 25 historical trends in CO₂ uptake and export production) compared to CanESM5 and CanESM2, as 26 27 well as to observations and other CMIP6 models. Both CanESM5 models show gains in skill relative to CanESM2, which are attributed primarily to improvements in ocean circulation. 28 CanESM5-CanOE shows improved skill relative to CanESM5 for most major tracers at most 29 depths. CanESM5-CanOE includes a prognostic iron cycle, and maintains high nutrient / low 30 chlorophyll conditions in the expected regions (in CanESM2 and CanESM5, iron limitation is 31 specified as a temporally static 'mask'). Surface nitrate concentrations are biased low in the 32 subarctic Pacific and equatorial Pacific, and high in the Southern Ocean, in both CanESM5 and 33 CanESM5-CanOE. Export production in CanESM5-CanOE is among the lowest for CMIP6 34 35 models; in CanESM5 it is among the highest, but shows the most rapid decline after about 1980. CanESM5-CanOE shows some ability to simulate aspects of plankton community structure that a 36 single-species model can not (e.g., seasonal dominance of large cells), but is biased towards low 37 38 concentrations of zooplankton and detritus relative to phytoplankton. Cumulative ocean uptake of anthropogenic carbon dioxide through 2014 is lower in both CanESM5 models than in 39 observation-based estimates (145 PgC) or the model ensemble mean (144 PgC), and is lower in 40 41 CanESM5-CanOE (122 PgC) than in CanESM5 (132 PgC).

42

43 1. Introduction

45	The Canadian Centre for Climate Modelling and Analysis has been developing coupled models
46	with an interactive carbon cycle for more than a decade (Arora et al., 2009; 2011; Christian et al.,
47	2010). The Canadian Earth System Model version 5 (CanESM5, Swart et al., 2019a) is an
48	updated version of CanESM2 (Arora et al., 2011), with a new ocean model based on the Nucleus
49	for European Modelling of the Ocean (NEMO) system version 3.4. The ocean biogeochemistry
50	modules were developed in-house. CanESM5 uses the same ocean biology model as CanESM1
51	(Christian et al., 2010) and CanESM2 (Arora et al., 2011), the Canadian Model of Ocean Carbon
52	(CMOC; Zahariev et al., 2008). An additional model was developed for CMIP6, called the
53	Canadian Ocean Ecosystem model (CanOE). The biological components of CanOE are of
54	substantially greater complexity than CMOC, including multiple food chains, flexible
55	phytoplankton elemental ratios, and a prognostic iron (Fe) cycle. The two coupled models are
56	known as CanESM5 and CanESM5-CanOE, respectively.
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57 58 59 60 61 62	The reasons for developing both models are, firstly, to evaluate the effect of changes in ocean circulation between CanESM2 and CanESM5 on ocean biogeochemistry by running the new climate model with the same ocean biogeochemistry, and secondly because CanOE is substantially more expensive computationally (as it has 19 tracers vs 7 the total computation cost is 2-3 times greater). Most CMIP6 experiments were run with CanESM5 only, as ocean

The CMIP6 experiments published for CanESM5-CanOE are listed in Supplementary Table S1. 66 67

CMOC is a nutrient-phytoplankton-zooplankton-detritus (NPZD) model with highly 68 parameterized representations of phytoplankton Fe limitation, dinitrogen (N_2) fixation and 69 denitrification, and calcification and calcite dissolution (Zahariev et al., 2008; Supplementary 70 Figure S1). CanESM1 and CanESM2 did not include oxygen;CanESM5 includes oxygen as a 71 72 purely 'downstream' tracer that does not affect other biogeochemical processes. In CanESM5-CanOE, denitrification is prognostic and dependent on the concentration of oxygen. Among the 73 less satisfactory aspects of CMOC biogeochemistry are, firstly, that Fe limitation is specified as a 74 75 static 'mask' that does not change with climate (it is calculated from the present-day climatological distribution of nitrate, based on the assumption that regions without iron 76 limitation will have complete drawdown of surface nitrate at some point in the year), and 77 78 secondly, that denitrification is parameterized so that nitrogen (N) is conserved within each vertical column, i.e., collocated with N₂ fixation in tropical and subtropical open-ocean regions 79 (Zahariev et al., 2008; Riche and Christian, 2018). This latter simplification produced excessive 80 accumulations of nitrate in Eastern Boundary Current (EBC) regions where most denitrification 81 82 occurs. CMOC also has a tendency to produce rather stark extremes of high and low primary and 83 export production (Zahariev et al., 2008), a well-known problem of NPZD models (Armstrong, 1994; Friedrichs et al., 2007). Our intent in developing CanOE was to alleviate, or at least 84 reduce, these biases, by including multiple food chains, a prognostic Fe cycle, and prognostic 85 86 denitrification. Dinitrogen fixation is still parameterized, but the CanOE parameterization 87 includes Fe (but not P) limitation, whereas in CMOC N₂ fixation tends to grow without bound in 88 a warming ocean as there is no P or Fe limitation (Riche and Christian, 2018).

90	In this paper we present a detailed model description for CanOE and an evaluation of both
91	CanESM5 and CanESM5-CanOE relative to observational data products and other available
92	models. CMOC has been well described previously (Zahariev et al., 2008) and the details are not
93	reiterated here. In some cases, CanESM2 results are also shown to illustrate which differences in
94	the model solutions arise largely from the evolution of the physical ocean model, and which are
95	specifically associated with different representations of biogeochemistry. An overall evaluation
96	of the CanESM5 physical ocean model is given in Swart et al. (2019a). Here, we focus on
97	biogeochemical variables, and have evaluated model performance in three main areas: (1) the
98	distribution of major tracers like oxygen, DIC and alkalinity, and the resulting saturation state for
99	CaCO ₃ minerals, (2) the iron cycle and its interaction with the nitrogen cycle, and (3) plankton
100	community structure and the concentration and export of particulates. We first address the major
101	chemical species that are common to both models (and almost all other Earth System Models) to
102	determine whether a more complex biology model measurably improves skill, and whether the
103	updated circulation model improves skill relative to CanESM2. Then we examine the areas
104	where our two models differ: the presence of a prognostic iron cycle and multiple food chains in
105	CanOE. More specifically, does CanESM5-CanOE reproduce the geographic distribution of
106	High-Nutrient, Low-Chlorophyll (HNLC) regions? Does the large phytoplankton / large
107	zooplankton food chain become dominant under nutrient-rich conditions, and how does having
108	multiple detrital size classes affect particle flux and remineralization length scale? Following this
109	model evaluation, we present historical trends in ocean anthropogenic CO ₂ uptake, export
110	production, and total volume of low-oxygen waters over the historical (1850-2014) experiment.
111	Possible future changes under Shared Socioeconomic Pathway experiments will be addressed in

subsequent publications.

2. Model Description

116	CanESM5 (Swart et al., 2019a) is an updated version of CanESM2 (Arora et al., 2011), with an
117	entirely new ocean. The atmosphere model has the same T63 horizontal resolution, and contains
118	some important improvements in atmospheric physics (Swart et al., 2019a). The land surface
119	(Canadian Land Surface Scheme) and terrestrial carbon cycle (Canadian Terrestrial Ecosystem
120	Model) models are substantially the same as in CanESM2 with minor modifications as described
121	by Arora et al. (2020). The CanESM5 ocean is based on the NEMO modelling system version
122	3.4, with a horizontal resolution of 1° , telescoping to $1/3^\circ$ in the tropics, and 45 vertical levels
123	ranging in thickness from ~ 6 m near the surface to ~ 250 m in the deep ocean (Swart et al.,
124	2019a). All physical climate model components are the same in CanESM5 and CanESM5-
125	CanOE. There are no feedbacks between biology and the physical ocean model, so the physical
126	climate of CanESM5 and CanESM5-CanOE is identical in experiments with prescribed
127	atmospheric CO ₂ concentration.
128	

The NEMO system is a publicly available archive of codes based on the OPA (Océan
PArallelisé) ocean model (Madec and Imbard, 1996; Guilyardi and Madec, 1997) and the
Tracers in Ocean Paradigm (TOP) module for tracer advection and mixing. Our ocean
biogeochemistry modules are built within TOP, using NEMO v3.4.1, but have also been
implemented in NEMO 3.6 for regional downscaling applications (Holdsworth et al., 2021).

135 Carbon chemistry is based on the Best Practices Guide (Dickson et al., 2007) and the OMIP-BGC data request (Orr et al., 2017) and are identical in CanESM5 and CanESM5-CanOE. All 136 calculations are done on the total scale and the recommended formulae for the equilibrium 137 constants are employed. The carbon chemistry solver was run for a fixed number of iterations 138 (ten in the surface layer, and five in the subsurface layers in CanESM5-CanOE). CanESM5 does 139 not solve the carbon chemistry equations in the subsurface layers. OMIP-BGC formulations for 140 CO₂ and O₂ solubility and gas exchange are employed. It is important to note here that the 141 carbon chemistry and gas exchange formulations used in CanESM2 (and other CMIP5 models) 142 are slightly different than those used in CMIP6. However, this difference is of little functional 143 significance, i.e., it will have a negligible impact on the distribution of [CO3⁻⁻] compared to the 144 differences in DIC and alkalinity distribution. The initialization fields for nitrate, DIC and 145 alkalinity were also different in CanESM2. This will affect the total ocean inventory of DIC but 146 not the spatial distribution if the model is well equilibrated. 147

148

The CanOE biology model is a substantially new model based on the cellular regulation model 149 of Geider et al. (1998). There are two phytoplankton size classes, and each group has four state 150 variables: C, N, Fe and chlorophyll. Photosynthesis is decoupled from cell production and 151 photosynthetic rate is a function of the cell's internal N and Fe quotas. Each functional group has 152 a specified minimum and maximum N quota and Fe quota, and nutrient uptake ceases when the 153 154 maximal cell quota is reached. Chlorophyll synthesis is a function of N uptake and increases at 155 low irradiance. There are also two size classes each of zooplankton and detritus. Small zooplankton graze on small phytoplankton, while large zooplankton graze on both large 156 phytoplankton and small zooplankton. Small detritus sinks at 2 m d⁻¹ and large detritus at 30 157

m d⁻¹ (in CanESM5 there is a single detrital pool with a sinking rate of 8 m d⁻¹). Model
parameters and their values are listed in Table 1. A schematic of the model is shown in Figure 1.
2.1 Photosynthesis and Phytoplankton Growth

For simplicity and clarity, the equations are shown here for a single phytoplankton species, and
do not differ structurally for small and large phytoplankton. Some parameter values differ for the
two phytoplankton groups; all parameter values are listed in Table 1.

166

167 Temperature dependence of photosynthetic activity is expressed by the Arrhenius equation

168
$$T_f = \exp(-\frac{E_{ap}}{R}(\frac{1}{T} - \frac{1}{T_{ref}}))$$
 (1)

169 where E_{ap} is an enzyme activation energy that corresponds approximately to that of RuBisCo (cf.

170 Raven and Geider 1988), *R* is the gas constant (8.314 J mol⁻¹ K⁻¹), and temperature *T* and

171 reference temperature T_{ref} are in Kelvin. Maximal rates of nutrient (either N or Fe, but

172 generically referred to here with the superscript X) uptake are given by

173
$$V_{max}^{X} = V_{ref}^{X} T_f \left(\frac{Q_{max}^{X} - Q^{X}}{Q_{max}^{X} - Q_{min}^{X}} \right)^{0.05}$$
(2)

where V_{max}^{X} is the maximal uptake rate in mg of nutrient X per mg of cell C, X can represent N or Fe, Q is the nutrient cell quota and Q_{min} and Q_{max} its minimum and maximum values, and V_{ref}^{X} is a (specified) basal rate at $T=T_{ref}$ and $Q=Q_{min}$. These maximum rates are then reduced according to the ambient nutrient concentration, i.e.

178
$$V^N = V_{max}^N (L_{NH4} + (1 - L_{NH}) L_{NO})$$
 (3a)

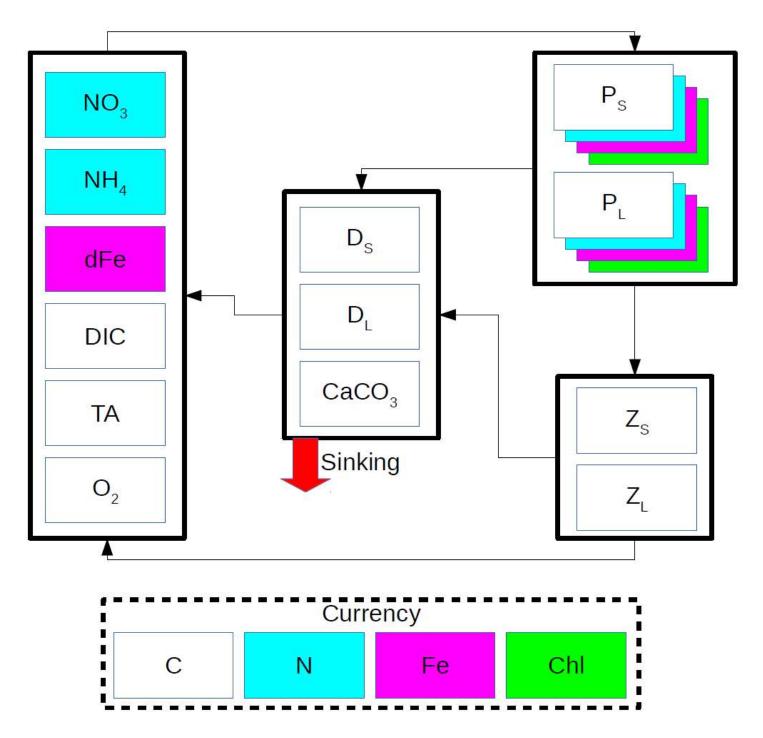


Figure 1 - Schematic of the CanOE biology model. Model currencies including chlorophyll (Chl) are indicated by coloured boxes except oxygen (O₂) and carbonate (CaCO₃). Arrows indicate flows of carbon (C), nitrogen (N) and iron (Fe) between compartments containing small (S) and large (L) phytoplankton (P), zooplankton (Z), and detritus (D) components; counterflows of oxygen are not shown.

Table 1 – Ecosystem model parameters.

Symbol	Description	Unit	
<u> </u>	D. C	IZ IZ	200.15
T _{ref}	Reference temperature	K	298.15
E_{ap}	Activation energy for photosynthesis	kJ mol ⁻¹	37.4
Q ^N mins	Small phytoplankton minimum N quota	g N g C ⁻¹	0.04
Q ^N _{maxs}	Small phytoplankton maximum N quota	g N g C ⁻¹	0.172
Q^{N}_{minl}	Large phytoplankton minimum N quota	g N g C ⁻¹	0.04
Q^{N}_{maxl}	Large phytoplankton maximum N quota	g N g C ⁻¹	0.172
$Q^{Fe}_{\ mins}$	Small phytoplankton minimum Fe quota	μg Fe g C ⁻¹	4.65
Q ^{Fe} maxs	Small phytoplankton maximum Fe quota	μg Fe g C ⁻¹	93.
Q ^{Fe} minl	Large phytoplankton minimum Fe quota	μg Fe g C ⁻¹	6.5
Q ^{Fe} maxl	Large phytoplankton maximum Fe quota	μg Fe g C ⁻¹	70.
V ^N _{ref}	Reference rate of N uptake	g N g C ⁻¹ d ⁻¹	0.6
V ^{Fe} _{ref}	Reference rate of Fe uptake	μg Fe g C ⁻¹ d ⁻¹	79.
P ^C _{ref}	Reference rate of photosynthesis	g C g C ⁻¹ d ⁻¹	3
k _{XU}	Rate coefficient for exhudation	d ⁻¹	1.7
k _{dgr}	Rate coefficient for chlorophyll degradation	d ⁻¹	0.02
ζ	Respiratory cost of biosynthesis	g C g N ⁻¹	2
α_{chl}	Initial slope of P-E curve	$g C g CHL^{-1} h^{-1} (\mu mol m^{-2} s^{-1})^{-1}$	1.08
$\Theta_{max}{}^N$	Maximum chlorophyll-nitrogen ratio	g g ⁻¹	0.18
K _{NiS}	Half-saturation for small phytoplankton nitrate uptake	mmol ⁻¹ m ³	0.1
K _{NaS}	Half-saturation for small phytoplankton ammonium uptake	mmol ⁻¹ m ³	0.05
K _{FeS}	Half-saturation for small phytoplankton iron uptake	nmol ⁻¹ m ³	100
K _{NiL}	Half-saturation for large phytoplankton nitrate uptake	mmol ⁻¹ m ³	1.0
K _{NaL}	Half-saturation for large phytoplankton ammonium uptake	mmol ⁻¹ m ³	0.05
K _{FeL}	Half-saturation for large phytoplankton iron uptake	nmol ⁻¹ m ³	200
m _{1S}	Small phytoplankton/zooplankton mortality rate (linear)	d ⁻¹	0.05
m _{2S}	Small phytoplankton/zooplankton mortality coefficient	(mmol C m ⁻³) ⁻¹ d ⁻¹	0.06
m _{1L}	Large phytoplankton/zooplankton mortality rate (linear)	d ⁻¹	0.1

m_{2L}	Large phytoplankton/zooplankton mortality coefficient	$(\text{mmol C m}^{-3})^{-1} d^{-1}$	0.06
X _{minp}	Minimum phytoplankton concentration for linear mortality	mmol C m ⁻³	0.01
aL	Large zooplankton grazing parameter	(mmol C m ⁻³) ⁻¹	0.25
GL0	Large zooplankton maximum grazing rate	d-1	0.85
a _S	Small zooplankton grazing parameter	(mmol C m ⁻³) ⁻¹	0.25
G _{S0}	Small zooplankton maximum grazing rate	d-1	1.7
λ	Assimilation efficiency	n.d.	0.8
r _{zs}	Microzooplankton specific respiration rate at T _{ref}	d-1	0.3
r _{zl}	Mesozooplankton specific respiration rate at T _{ref}	d-1	0.1
r ₁	Small detritus remineralization rate at T _{ref}	d ⁻¹	0.25
r ₂	Large detritus remineralization rate at T _{ref}	d ⁻¹	0.25
Ear	Activation energy for detritus remineralization	kJ mol ⁻¹	54.0
Ws	Small detritus sinking speed	m d ⁻¹	2.
Wl	Large detritus sinking speed	m d ⁻¹	30.
WCa	CaCO ₃ sinking speed	m d ⁻¹	20.
P _{Ca}	CaCO ₃ production as fraction of mortality	mol CaCO ₃ molC ⁻¹	0.05
k _{Ca}	CaCO ₃ dissolution rate	d ⁻¹	0.0074
S _{Fe1}	Dissolved iron scavenging loss rate (Fe≤L _{Fe})	d ⁻¹	0.001
S _{Fe2}	Dissolved iron scavenging loss rate (Fe>L _{Fe})	d ⁻¹	2.5
L _{Fe}	Ligand concentration	nmol Fe m ⁻³	600.
P _{Fe}	POC-dependence parameter for Fe scavenging	(mmolC m ⁻³) ⁻¹	0.66
k _{NH4ox}	Nitrification rate constant	d ⁻¹	0.05
K _E	Half-saturation for irradiance inhibition of nitrification	W m ⁻²	1.
k _{dnf}	Light and nutrient saturated rate of N ₂ fixation at 30°C	mmol m ⁻³ d ⁻¹	0.0225
a	Initial slope for irradiance-dependence of N ₂ fixation	(W m ⁻²) ⁻¹	0.02
K _{Fe}	Half-saturation for Fe dependence of N ₂ fixation	nmol m ⁻³	100.
K _{NO3}	Half-saturation for DIN inhibition of N ₂ fixation	mmol m ⁻³	0.1
O _{mxd}	O ₂ concentration threshold for denitrification	mmol m ⁻³	6.
A _f	Anammox fraction of N loss to denitrification	n.d.	0.25

182 where $L_{\text{NH4}} = \frac{N_a}{K_{\text{NaX}} + N_a}$ and $L_{\text{NO3}} = \frac{N_i}{K_{\text{NiX}} + N_i}$, with N_i and N_a indicating nitrate and ammonium

183 respectively, and

184
$$V^{Fe} = V_{max}^{Fe} \left(\frac{Fe}{K_{FeX} + Fe}\right)$$
 (3b)

where X indicates large or small phytoplankton (Table 1). The maximal carbon-based growthrate is given by

187
$$P_{max}^{C} = P_{ref}^{C} T_{f} \min\{\frac{Q^{N} - Q_{min}^{N}}{Q_{max}^{N} - Q_{min}^{N}} \cdot \frac{Q^{Fe} - Q_{min}^{Fe}}{Q_{max}^{Fe} - Q_{min}^{Fe}}\}$$
(4)

188 where P^{C}_{ref} is the rate at the reference temperature T_{ref} under nutrient-replete conditions

189 $(Q=Q_{max})$. The light-limited growth rate is then given by

190
$$P_{phot}^{C} = P_{max}^{C} \left(1 - e^{-\alpha_{chl} E \theta_C} / P_{max}^{C} \right)$$
(5)

where E is irradiance and θ_C is the chlorophyll-to-carbon ratio. The rate of chlorophyll synthesis is

193
$$\rho_{\rm chl} = \theta_{max}^{N} \frac{P_{phot}^{C}}{\epsilon \alpha_{\rm chl} \theta}$$
(6)

These rates are then used to define a set of state equations for phytoplankton carbon (C_p),
nitrogen (N_p), iron (Fe_p), and chlorophyll (M).

196
$$\frac{dC_p}{dt} = (P_{phot}^C - \zeta V_N)C_p - (G + C_{XS}) - m_1C_p - m_2C_p^2 - k_{XU}C_{INTR}$$
(7)

197 where
$$\zeta$$
 is the respiratory cost of biosynthesis, *G* is the grazing rate (equation 12), C_{XS} is the
198 excess (above the ratio in grazer biomass) carbon in grazing losses (see below equation 16a), m_1
199 and m_2 are coefficients for linear and quadratic nonspecific mortality terms, C_{INTR} is the
200 concentration of intracellular carbohydrate carbon in excess of biosynthetic requirements, and
201 k_{XU} is a rate coefficient for its exudation to the environment. The nonspecific mortality terms are

set to 0 below 0.01 mmol C m⁻³, to prevent biomass from being driven to excessively low levels
in the high latitudes in winter; linear mortality terms can result in biomass declining to levels
from which recovery would take much longer than the brief Arctic summer (Hayashida, 2018).
The full equation for phytoplankton N, Fe and chlorophyll are

206
$$\frac{dN_p}{dt} = \frac{V^N}{Q_N} - (G + m_1 C_p + m_2 C_p^2) R_{\rm NC} - N_{\rm XS}$$
(8)

207
$$\frac{dFe_p}{dt} = \frac{V^{Fe}}{Q_{Fe}} - (G + m_1 C_P + m_2 C_p^2) R_{FeC} - Fe_{XS}$$
(9)

208
$$\frac{\mathrm{d}M}{\mathrm{dt}} = \frac{\rho_{\mathrm{chl}} V^N}{\theta_C} M - (G + m_1 C_p + m_2 C_p^2) \theta_C - k_{\mathrm{dgr}} M \tag{10}$$

where k_{dgr} is a rate coefficient for nonspecific losses of chlorophyll e.g., by photooxidation, in addition to losses to grazing and other processes that also affect C_p , N_p , and Fe_p . N_{XS} and Fe_{XS} are remineralization of "excess" (relative to grazer or detritus ratios) N or Fe and are defined below (equation 16).

213

214 **2.2 Grazing and Food Web Interactions**

215

216 Grazing rate depends on the phytoplankton carbon concentration, which most closely represents

the food concentration available to the grazer (Elser and Urabe 1999; Loladze et al. 2000).

218 Zooplankton biomass is also in carbon units. State equations for small and large zooplankton are

219
$$\frac{dZ_s}{dt} = \lambda G_s - (R + G_Z + m_{1s}Z_s + m_{2s}Z_s^2)$$
(11a)

220
$$\frac{dZ_L}{dt} = \lambda G_L - (R + m_{1L}Z_L + m_{2L}Z_L^2)$$
(11b)

221 where

222
$$G_s = G_{so}(1 - e^{-a_s C_{ps}})Z_s$$
 (12a)

223
$$G_L = G_{L0}(1 - e^{-a_l(C_{pl} + Z_s)})Z_L$$
(12b)

for small and large zooplankton respectively, G_Z is grazing of small zooplankton by large zooplankton, *R* is respiration, and m_1 and m_2 are nongrazing mortality rates. Large zooplankton grazing is divided into grazing on large phytoplankton and small zooplankton in proportion to the relative abundance of each

$$228 \qquad G_P = G_L \frac{P_l}{P_l + Z_s} \tag{13a}$$

$$G_Z = G_L \frac{Z_s}{P_l + Z_s}$$
(13b)

230 Zooplankton biomass loss to respiration is given by

231
$$R = max\{r_z T_f Z - C_{\rm XS}, 0\}$$
(14)

and uses the same activation energy as photosynthesis. Respiration (R) is assumed to consume 232 only carbon and not result in catabolism of existing biomass when "excess" carbon is available 233 234 in the prey. In addition, conservation of mass must be maintained by recycling to the dissolved pool grazer consumption of elements in excess of biosynthetic requirements when grazer and 235 prey elemental ratios differ. In the case where the nutrient quota (relative to carbon) exceeds the 236 237 grazer fixed ratio, the excess nutrient is remineralized to the dissolved inorganic pool. In the case where the nutrient quota is less than the grazer ratio, the grazer intake is reduced to what can be 238 supported by the least abundant nutrient (relative to the grazer biomass ratio) and excess carbon 239 is remineralized. For the case of two nutrients (in this case N and Fe) it is necessary to define 240

241
$$G' = G\min\left\{\frac{N_P}{C_P}R_{\rm CN}, \frac{Fe_P}{C_P}R_{\rm CFe}, 1\right\}$$
(15)

where G is equal to G_S (equation 12a) for small zooplankton and G_P (equation 13a) for large

243 zooplankton, and R_{XY} indicates the fixed ratio of element X to element Y in grazer biomass. The

244 'excess' carbon available for respiration is

245
$$C_{\rm XS} = G' \left\{ \frac{C_P}{N_P} R_{\rm NC} - 1, \frac{C_P}{Fe_P} R_{\rm FeC} - 1, 0 \right\}$$
 (16a)

and the excess nutrients remineralized to their inorganic pools are

247
$$N_{\rm XS} = G'\max\left\{\frac{N_P}{C_P} - R_{\rm NC}.0\right\}\varepsilon + G'\max\left\{R_{\rm NC}\left(\frac{N_P}{Fe_P}R_{\rm FeN} - 1\right).0\right\}(1-\varepsilon)$$
(16b)

248
$$\operatorname{Fe}_{\mathrm{XS}} = G'\max\left\{\frac{\operatorname{Fe}_{P}}{c_{P}} - R_{\mathrm{FeC}}, 0\right\}\varepsilon + G'\max\left\{R_{\mathrm{FeC}}(\frac{\operatorname{Fe}_{P}}{N_{P}}R_{\mathrm{NFe}} - 1), 0\right\}(1 - \varepsilon)$$
(16c)

249 where

$$250 \qquad \varepsilon = \frac{\max\{C_{xs}, 0\}}{C_{xs} + \Delta}$$

is a switch to prevent double-counting in cases where one of the terms is redundant (the excess relative to the least abundant element is included in the other term), but would otherwise be nonzero (Δ is a constant equal to 10⁻¹⁵, to prevent divide-by-zero). For three elements, there are 3! = 6 possible cases: for N greater or less than C_PR_{NC}, Fe may be either in excess relative to both C and N, deficient relative to both, or in excess relative to one but not the other (Table 2).

Table 2 - Cases where the 'excess' terms are nonzero. These terms are always greater than or equal to zero, and always zero when the phytoplankton elemental ratio is equal to the grazer biomass ratio. A plus (+) sign indicates that a specific term is positive. N_1 and N_2 , Fe_1 and Fe_2 indicate the first and second terms in equations 16b and 16c. R_{NC} is the grazer N/C (Redfield) ratio.

	Fe in excess relative to				Fe in excess relative to C				Fe deficient relative to						
	both C and N				or N but not both					both C and N					
	C	N_1	N ₂	Fe ₁	Fe ₂	C	N_1	N_2	Fe ₁	Fe ₂	С	N_1	N ₂	Fe ₁	Fe ₂
N/C>R _{NC}		+		+			+		+		+		+		
N/C <r<sub>NC</r<sub>	+				+	+				+	+		+		

262 2.3 Organic and Inorganic Pools

263

There are two pools of detritus with different sinking rates but the same fixed elemental ratios.
Detrital C/N/Fe ratios are the same as zooplankton, so zooplankton mortality or grazing of small
zooplankton by large zooplankton produce no 'excess'. Phytoplankton mortality, and defecation

by zooplankton grazing on phytoplankton, produces excess nutrient or excess C that needs to be

recycled into the inorganic pool in a similar fashion as outlined above for the assimilated fraction

269 of grazing on phytoplankton.

270 The conservation equations for detrital C are

271
$$\frac{dD_s}{dt} = m_1(C_{ps} + Z_s) + m_2(C_{ps}^2 + Z_s^2) - r_1 D_s T_g - w_s \frac{dD_s}{dz}$$
(17a)

272
$$\frac{\mathrm{d}D_l}{\mathrm{d}t} = m_1(C_{\mathrm{pl}} + Z_L) + m_2(C_{pl}^2 + Z_L^2) - r_2 D_l T_g - w_l \frac{\mathrm{d}D_l}{\mathrm{d}z}$$
(17b)

where T_g is an Arrhenius function for temperature dependence of remineralization and w is the sinking speed. The conservation equations for inorganic C, N, and Fe are

275
$$\frac{dC_i}{dt} = (\zeta V^N - P_{\text{phot}}^C)C_p + R + C_{\text{XS}} + (r_1 D_s + r_2 D_l)T_g$$
(18a)

276
$$\frac{dN_i}{dt} = -\frac{V^N}{Q^N} N_p(\frac{L_{N03}}{L_{N03} + L_{NH4}}) + N_{ox} - N_{dentr}(1 - A_f)$$
(18b)

277
$$\frac{dN_a}{dt} = -\frac{V^N}{Q^N} N_p \left(\frac{L_{\text{NH4}}}{L_{\text{NO3}} + L_{\text{NH4}}}\right) + \frac{R}{R_{\text{CN}}} + N_{\text{XS}} + (r_1 D_s + r_2 D_l) R_{\text{NC}} T_g - N_{\text{ox}} + N_{\text{dnf}} - N_{\text{dentr}} A_f \quad (18c)$$

278
$$\frac{\mathrm{dFe}}{\mathrm{dt}} = \frac{V^{\mathrm{Fe}}}{Q^{\mathrm{Fe}}} \mathrm{Fe}_p + \frac{R}{R_{\mathrm{CFe}}} + \mathrm{Fe}_{\mathrm{XS}} + (r_1 D_s + r_2 D_l) R_{\mathrm{FeC}} T_g$$
(18d)

- where N_{ox} is microbial oxidation of ammonium to nitrate (nitrification), N_{dnf} and N_{dentr} are
- sources and sinks associated with dinitrogen fixation and denitrification, and A_f is the ammonium

fraction of denitrification losses, associated with anaerobic ammonium oxidation ("anammox").

The oxygen equation is essentially the inverse of equation 18a, with additional terms foroxidation and reduction of N, i.e.,

284
$$\frac{dO_2}{dt} = -\frac{dC_i}{dt} + 2\frac{V^N}{Q^N}N_p(\frac{L_{NO3}}{L_{NO3}+L_{NH4}}) - 2N_{ox}$$
(19)

285 Nitrification is given by

286
$$N_{\rm ox} = k_{\rm NH4ox} N_a \frac{\kappa_E}{\kappa_E + E(z)}$$
(20)

where E(z) is the layer mean irradiance at depth *z*. Dinitrogen fixation is parameterized as an external input of ammonium dependent on light, temperature and Fe availability, and inhibited by high ambient concentrations of inorganic N,

290
$$N_{\rm dnf} = k_{\rm dnf} T_{\rm dnf} (1 - e^{-aE}) (\frac{Fe}{K_{\rm Fe} + Fe}) (\frac{K_{\rm N03}}{K_{\rm N03} + N_i + N_a})$$
 (21)

where T_{dnf} =max(0, 1.962(T_f - 0.773)), i.e., a linear multiple of equation (1) that is 0 at T<20°C and unity at T=30°C.

293

294 Denitrification is parameterized as a fraction of total remineralization that increases as a linear 295 function of oxygen concentration for concentrations less than a threshold concentration O_{mxd}

296
$$N_{\text{frxn}} = 1 - \frac{\min(O_2, O_{\text{mxd}})}{O_{\text{mxd}}}$$
 (22)

Remineralization is then divided among oxygen $(1-N_{frxn})$, nitrate $(0.875N_{frxn})$, and ammonium ($0.125N_{frxn}$) assuming an average anammox contribution of 25% (Babbin et al., 2014). We use this average ratio of anammox to classical denitrification to partition fixed N losses between NO₃⁻ and NH₄⁺; the DIC sink and organic matter source associated with anammox are small and are neglected here.

303 **2.4 Calcification, Calcite Dissolution, and Alkalinity**

304

In CanOE, calcification is represented by a prognostic detrital calcite pool with its own sinking 305 rate (distinct from that of organic detritus), and calcite burial or dissolution in the sediments 306 depends on the saturation state (100% burial when $\Omega_{\rm C} \ge 1$, 100% dissolution when $\Omega_{\rm C} < 1$). 307 Calcification is represented by a detrital calcium carbonate (CaCO₃) state variable, but no 308 309 explicit calcifier groups. Detrital CaCO₃ sinks in the same fashion as detrital particulate organic carbon (POC), with a sinking rate independent of those for large and small organic detritus. 310 Calcite production is represented as a fixed fraction of detritus production from small 311 312 phytoplankton and small zooplankton mortality:

313
$$\frac{dCa}{dt} = m_1 (C_{ps} + Z_s) P_{Ca} + m_2 (C_{ps}^2 + Z_s^2) P_{Ca} - k_{Ca} Ca - w_{Ca} \frac{dCa}{dz}$$
(23)

314 Calcite dissolution occurs throughout the water column as a first order process (i.e., no dependence on temperature or saturation state). Approximately 80% of calcite produced is 315 exported from the euphotic zone. Burial in the sediments is represented as a simple 'on/off' 316 317 switch dependent on the calcite saturation state (zero when $\Omega_C < 1$ and 1 when $\Omega_C \geq 1$). In 318 CanESM5, calcification is parameterized by a temperature dependent "rain ratio" (Zahariev et al., 2008) and 100% burial of calcite that reaches the seafloor is assumed. Calcite burial in both 319 models is balanced by an equivalent source of DIC and alkalinity at the ocean surface (in the 320 321 same vertical column) as a crude parameterization of fluvial sources.

322

323 For each mole of calcite production two moles of alkalinity equivalent are lost from the

324 dissolved phase; the reverse occurs during calcite dissolution. There are additional sources and

sinks for alkalinity associated with phytoplankton nutrient (NH_4^+, NO_3^-) uptake, organic matter

326 remineralization, nitrification, denitrification and dinitrogen fixation (Wolf-Gladrow et al., 2007, see Supplementary Table S2). The anammox reaction does not in itself contribute to alkalinity 327 (Jetten at al., 2001), but there is a sink associated with ammonium oxidation to nitrite (the model 328 does not distinguish between nitrite and nitrate). Autotrophic production of organic matter by 329 anammox bacteria is a net source of alkalinity (Strous et al., 1998) but this source is extremely 330 331 small (~0.03 mol/molN) and is neglected here. Globally, the sources and sinks of alkalinity from the N cycle offset each other such that there is no net gain or loss as long as the global fixed N 332 pool is conserved (see below Sect. 2.5). If dinitrogen fixation and denitrification are allowed to 333 334 vary freely, there will generally be a net gain or loss of fixed N and, therefore, of alkalinity.

335

- 336 **2.5 External Nutrient Sources and Sinks**
- 337

External sources and sinks consist of river inputs, aeolian deposition, biological N₂ fixation, 338 339 denitrification, mobilization of Fe from reducing sediments, loss of Fe to scavenging, and burial of calcium carbonate in the sediments. Aeolian deposition of Fe is calculated from a climatology 340 341 of mineral dust deposition generated from offline (atmosphere-only) simulations with CanAM4 342 (von Salzen et al., 2013), with an Fe mass fraction of 5% and a fractional solubility of 1.4% in 343 the surface layer. Subsurface dissolution is parameterized based on PISCESv2 (Aumont et al., 344 2015); the total dissolution is 6.35%, with 22% of soluble Fe input into the first vertical layer 345 (see Supplementary material). Iron from reducing sediments is also based on PISCES, with a constant areal flux of 1000 nmol m⁻² d⁻¹ in the first model level, declining exponentially with 346 increasing seafloor depth (i.e., assuming that shelf sediments are the strongest source and the 347 348 sediments become progressively more oxygenated with increasing seafloor depth) with an efolding length scale of about 200 m. Scavenging of dissolved iron is first-order with a high rate (2.5 d⁻¹) for concentrations in excess of 0.6 nM (Johnson et al., 1997). For concentrations below this threshold, the rate is much lower (0.001 d⁻¹) and is weighted by the concentration of organic detritus (Christian et al., 2002b), i.e.,

353
$$\frac{dFe}{dt} = -FeS_{Fe1}min\{(D_S + D_L)P_{Fe}, 1\}$$
(24)

where Fe is the dissolved iron concentration, D_S and D_L are the small and large detritus 354 355 concentrations, S_{Fe1} is the first-order scavenging rate in surface waters with abundant particulates, and PFe is an empirical parameter to determine the dependence on particle 356 concentration (Table 1). The basis for this parameterization is that the rate of scavenging must 357 depend not only on the concentration of iron but on the concentration of particles available for it 358 359 to precipitate onto, and assumes that POC is strongly positively correlated with total particulate matter. Scavenging is treated as irreversible, i.e., scavenged Fe is not tracked and does not 360 reenter the dissolved phase. 361

362

N₂ fixation and denitrification vary independently in CanOE, so the global total N pool can 363 change. Conservation is imposed by adjusting the global total N pool according to the difference 364 between the gain from N₂ fixation and the loss to denitrification. A slight adjustment is applied 365 to the nitrate concentration at every grid point, while preserving the overall spatial structure of 366 367 the nitrate field. Adjustments are multiplicative rather than additive to avoid producing negative concentrations. This adjustment does not maintain (to machine precision) a constant global N 368 inventory but is intended to minimize long term drift, keeping it much smaller than the free 369 surface error (see below). This adjustment is applied every 10 days and has a magnitude of 370 approximately $7x10^{-8}$ of the total N. 371

373	When the total fixed N adjustment is applied, one mole of alkalinity is removed per mole of N
374	added or removed, to account for the alkalinity sources associated with N_2 fixation (creation of
375	new NH4 ⁺) and denitrification (removal of NO3 ⁻) (Wolf-Gladrow et al., 2007, see Supplementary
376	Table S2). As there is a 2 mol/molN sink associated with nitrification, this formulation is
377	globally conservative. As noted above, in CanOE CaCO3 can dissolve or be buried in the
378	sediments depending on the calcite saturation state. DIC and alkalinity lost to burial are
379	reintroduced at the ocean surface, at the same grid point as burial occurs, providing a crude
380	parameterization of river inputs so that global conservation is maintained (fresh water runoff
381	contains no DIC or alkalinity). However, the OPA free surface formulation is inherently
382	imperfect with regard to tracer conservation. Drift in total ocean alkalinity and nitrogen over
383	time is on the order of 0.01% and 0.03% per thousand years, respectively (losses due to the free
384	surface are generally larger for tracers with less homogeneous distributions).

2.6 Ancillary data

For first-order model validation we have relied largely on global gridded data products rather
than individual profile data. Global gridded data from World Ocean Atlas 2018 (WOA2018)
(Locarnini et al., 2018; Zweng et al., 2018; Garcia et al., 2018a; 2018b) were used for
temperature, salinity, and oxygen and nitrate concentration. DIC and alkalinity were taken from
the GLODAPv2.2016b gridded data product (Key et al., 2015; Lauvset et al., 2016). Offline
carbon chemistry calculations were done following the Best Practices Guide (Dickson et al.,
2007) and the OMIP-BGC protocols (Orr et al., 2017), and are identical to those used in the

models except that constant reference concentrations were used for phosphate (1 μ M) and silicate (10 μ M).

397

398	There is no global gridded data product for Fe, but we have made use of the GEOTRACES
399	Intermediate Data Product 2017 (Schlitzer et al., 2018), and the data compilations from MBARI
400	(Johnson et al., 1997; 2003) and PICES Working Group 22 (Takeda et al., 2013). The latter two
401	are concentrated in the Pacific, while GEOTRACES is more global. The combined data sets
402	provide more than 10000 bottle samples from more than 1000 different locations (Supplementary
403	Figure S9a) (excluding some surface transect data that involve frequent sampling of closely
404	spaced locations along the ship track). More detail about model comparison to these data
405	compilations and the list of original references are given in the Supplementary information.
406	
407	Satellite ocean colour estimates of surface chlorophyll were taken from the combined
407 408	Satellite ocean colour estimates of surface chlorophyll were taken from the combined SeaWiFS/MODIS climatology described by Tesdal et al. (2016). Climatological satellite POC
408	SeaWiFS/MODIS climatology described by Tesdal et al. (2016). Climatological satellite POC
408 409	SeaWiFS/MODIS climatology described by Tesdal et al. (2016). Climatological satellite POC was downloaded from the NASA ocean colour web site and is based on the algorithm of
408 409 410	SeaWiFS/MODIS climatology described by Tesdal et al. (2016). Climatological satellite POC was downloaded from the NASA ocean colour web site and is based on the algorithm of Stramski et al. (2008) using MODIS-Aqua data. This climatology differs slightly from the
408 409 410 411	SeaWiFS/MODIS climatology described by Tesdal et al. (2016). Climatological satellite POC was downloaded from the NASA ocean colour web site and is based on the algorithm of Stramski et al. (2008) using MODIS-Aqua data. This climatology differs slightly from the chlorophyll one in terms of years included and sensors utilized, but as only climatological
408 409 410 411 412	SeaWiFS/MODIS climatology described by Tesdal et al. (2016). Climatological satellite POC was downloaded from the NASA ocean colour web site and is based on the algorithm of Stramski et al. (2008) using MODIS-Aqua data. This climatology differs slightly from the chlorophyll one in terms of years included and sensors utilized, but as only climatological concentrations are considered and each climatology covers ~15 years, these differences will have

416

417 CMIP6 model data were regridded by distance-weighted averaging using the Climate Data Operators (https://code.mpimet.mpg.de/projects/cdo/) to a common grid (2x2°, 33 levels) to 418 facilitate ensemble averaging. The vertical levels used are those used in GLODAP and in earlier 419 (through 2009) versions of the World Ocean Atlas (e.g., Locarnini et al., 2010). For large scale 420 tracer distributions, using a 1° or 2° grid makes little difference (for example, the spatial pattern 421 correlation between CanESM5 and observed oxygen concentration at specific depths differs by 422 an average of 0.0011). The years 1986-2005 of the Historical experiment were averaged into 423 climatologies or annual means, for meaningful comparison with observation-based data products. 424 425 The CMIP6 Historical experiment runs from 1850-2014 with atmospheric CO₂ concentration (and other atmospheric forcings) based on historical observed values. A single realization was 426 used in each case (see Table S3); as 20 year averages are used, internal variability is assumed to 427 have little effect (e.g., Arguez and Vose, 2011, see Table S4). Where time series are shown, 5-428 year means are used. 429

430

Sampling among CMIP6 models was somewhat opportunistic and the exact suite of models 431 varies among the analyses presented. When we conducted a search for a particular data field, we 432 433 included in the search parameters all models that published that field, and repeated the search at 434 least once for models that were unavailable the first time the search was executed. In some cases, model ensemble means excluded all but one model from a particular 'family' (e.g., there are 435 436 three different MPI-ESM models for which ocean biogeochemistry fields were published), as the 437 solutions were found to be similar and would bias the ensemble mean towards their particular climate. The models used are ACCESS-ESM1-5, CESM2, CESM2-WACCM, CNRM-ESM2-1, 438 439 GFDL-CM4, GFDL-ESM4, IPSL-CM6A-LR, MIROC-ES2L, MPI-ESM-1-2-HAM, MPI-

ESM1-2-LR, MPI-ESM1-2-HR, MRI-ESM2-0, NorESM2-LM, NorESM2-MM, and UKESM10-LL. Details of which variables and realizations are used for which models are given in
Supplementary Table S3.

443

444 **3. Results**

445

We first describe here the large-scale distribution of oxygen, DIC, alkalinity, and the saturation 446 state with respect to CaCO₃ that derives from these large-scale tracer distributions. Tracer 447 distributions result partly from ocean circulation and partly from biogeochemical processes. An 448 449 overall evaluation of the ocean circulation model is given in Swart et al. (2019a). Analyzing CanESM5 and CanESM5-CanOE (with identical circulation) as well as CanESM2 where 450 possible (same biogeochemistry as CanESM5 but different circulation) allows us to separate the 451 452 effects of physical circulation and biogeochemistry on evolving model skill with respect to largescale tracer distributions. In subsequent sections we address the main areas where CanESM5 and 453 CanESM5-CanOE differ, such as the interaction of the iron and nitrogen cycles and plankton 454 community structure. Finally, we present some temporal trends over the course of the historical 455 experiment (1850-2014). 456

457

458 **3.1 Distribution of oxygen**

459

460 The spatial distribution of oxygen concentration ([O₂]) at selected intermediate depths (400, 900,

and 1400 m) is shown in Figure 2 for gridded data from WOA2018 and differences of

462 CanESM5, CanESM5-CanOE, a model ensemble mean (MEM) of CMIP6 models (excluding

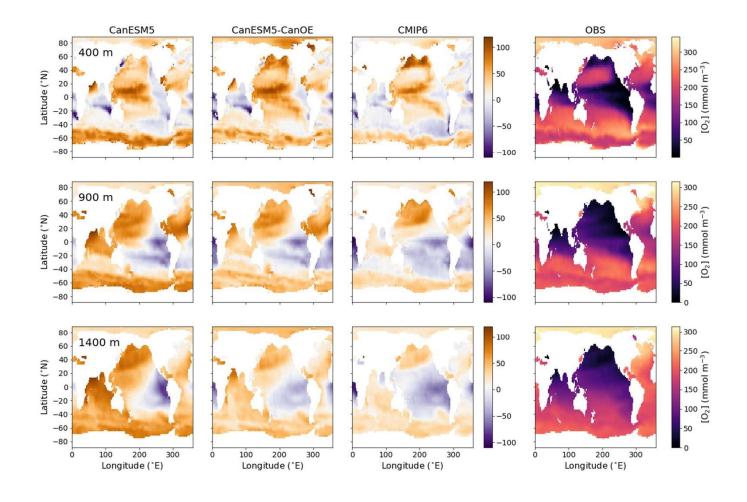


Figure 2 - Global distribution of oxygen (O_2) concentration in mmol m⁻³ at 400, 900, and 1400 m (rows). Observations (WOA2018) are in the right hand column; other columns show the difference from the observations of CanESM5-CanOE, CanESM5, and the mean for other CMIP6 models. Note different colour scales for different rows. Model concentrations are shown in Supplementary Figure S3.

464 CanESM5 and CanESM5-CanOE) from the observational data product. The depths were chosen to span the depth range where low oxygen concentrations exist; these low-oxygen environments 465 are of substantial scientific and societal interest and are sensitive to model formulation. The 466 major features are consistent across the models. Both CanESM models as well as the MEM show 467 elevated oxygen concentrations relative to observations, particularly in the North Pacific, the 468 469 North Atlantic and the Southern Ocean. In the Indian Ocean, both CanESM models show high oxygen concentrations in the Arabian Sea and deeper layers of the Bay of Bengal relative to 470 observations and the MEM; these biases are somewhat smaller in CanESM5-CanOE than in 471 472 CanESM5 (Figure 2).

473

The ocean's oxygen minimum zones (OMZs) are mostly located in the eastern Pacific Ocean, the 474 northern North Pacific, and the northern Indian Ocean; the spatial pattern changes with 475 476 increasing depth (Figure 2), but the OMZs are mostly located between 200 and 2000 m depth. Biases in the EBC regions are depth and model specific. CanESM5 shows particularly strong 477 oxygen depletion at 1400 m in the eastern tropical Pacific. In the southeastern Atlantic, models 478 tend to be biased low at the shallower depths, and show somewhat more variation at greater 479 480 depths (Figure 2). Overall, [O₂] biases tend to be positive over large areas of ocean with the 481 exception of some EBC regions, implying that models exaggerate the extent to which remineralization is concentrated in these regions. An alternate version of Figure 2 that shows the 482 modelled concentrations is given in Supplementary Figure S2. 483

484

The zonal mean oxygen concentration, saturation concentration, and apparent oxygen utilization
(AOU) are shown in Figure 3 for the same four cases. Again, the models generally show a

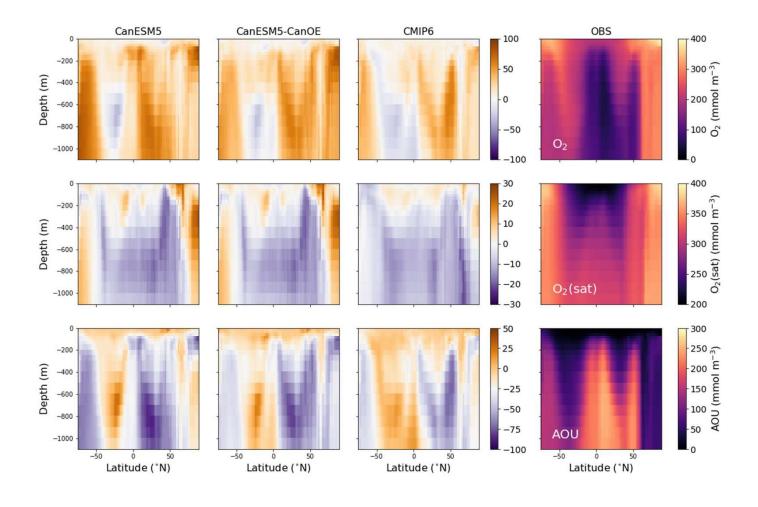


Figure 3 - Latitude-depth distribution (surface to 1750 m) of zonal mean oxygen concentration (O_2), oxygen concentration at saturation (O_2 (sat)), and apparent oxygen utilization (AOU) in mmol m⁻³. Observations (WOA2018) are in the right hand column; other columns show the difference from the observations of CanESM5-CanOE, CanESM5, and the mean for other CMIP6 models. Note different colour scales for different rows. Model concentrations are shown in Supplementary Figure S3.

488 positive bias in $[O_2]$, particularly in high-latitude deep waters. The major ocean circulation features are reproduced fairly well in all cases (e.g., weaker ventilation of low-latitude 489 subsurface waters, greater vertical extent of well-ventilated surface waters in the subtropics). The 490 saturation concentration (a function of temperature and salinity) generally shows relatively little 491 bias, implying that the bias in $[O_2]$ arises mainly from remineralization and/or ventilation. AOU 492 493 is lower than observed over much of the subsurface ocean. Regional biases are quite consistent across models, but are slightly greater in CanESM5 than in CanESM5-CanOE or the MEM, 494 except in the Arctic Ocean. Again, Supplementary Figure S2 includes a version of this plot that 495 496 shows the modelled concentration fields.

497

The skill of each model with respect to the distribution of O_2 at different depths is represented by 498 Taylor diagrams (Taylor, 2001) in Figure 4. These diagrams allow us to assess how well the 499 500 model reproduces the spatial distribution at a range of depths, because different physical and biogeochemical processes determine the distribution in different depth ranges. All of the CMIP6 501 models that were shown as an ensemble mean in Figures 2 and 3 are shown individually. The 502 large blue dots represent CanESM5, red CanESM5-CanOE, and grey the MEM; the smaller grey 503 dots represent the individual models. CanESM5-CanOE shows slightly higher pattern correlation 504 than CanESM5 at all depths. Both models compare favourably with the full suite of CMIP6 505 models, with r>0.85 for CanESM5 and r>0.9 for CanESM5-CanOE at all depths examined, and a 506 normalized standard deviation within $\pm 25\%$ of unity. 507

508

509 The total volume of ocean with $[O_2]$ less than 6 mmol m⁻³ (the threshold for denitrification

510 (Devol, 2008)) and 60 mmol m^{-3} (a commonly used index of hypoxia) is shown in Figure 5. The

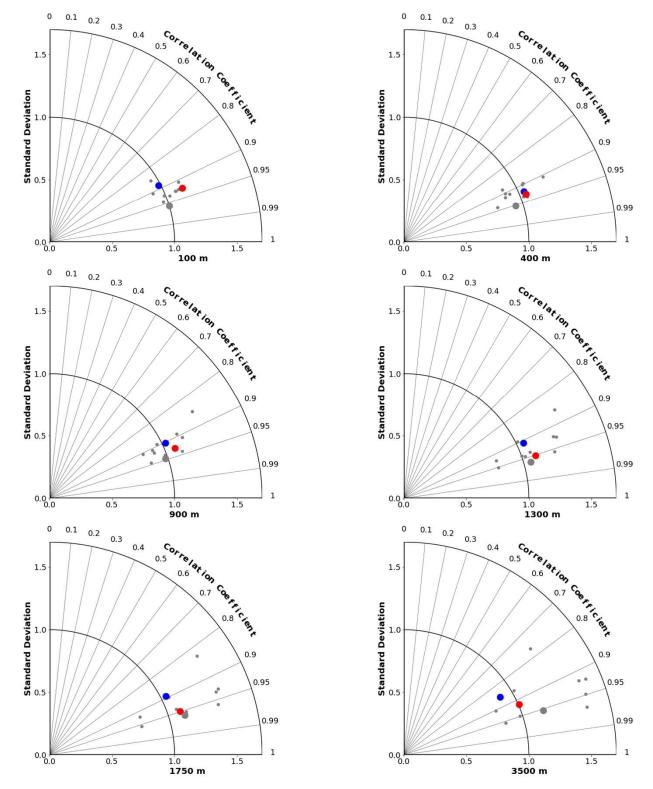


Figure 4 - Taylor diagrams (Taylor, 2001) comparing modelled and observed distributions of oxygen at specific depths from 100 to 3500 m. Angle from the vertical indicates spatial pattern correlation. Distance from the origin indicates ratio of standard deviation in modelled vs. observed (WOA2018) fields. Red dots represent CanESM5-CanOE, blue dots CanESM5, small grey dots other CMIP6 models, and large grey dots the model ensemble mean for all CMIP6 models except CanESM5 and CanESM5-CanOE. 29

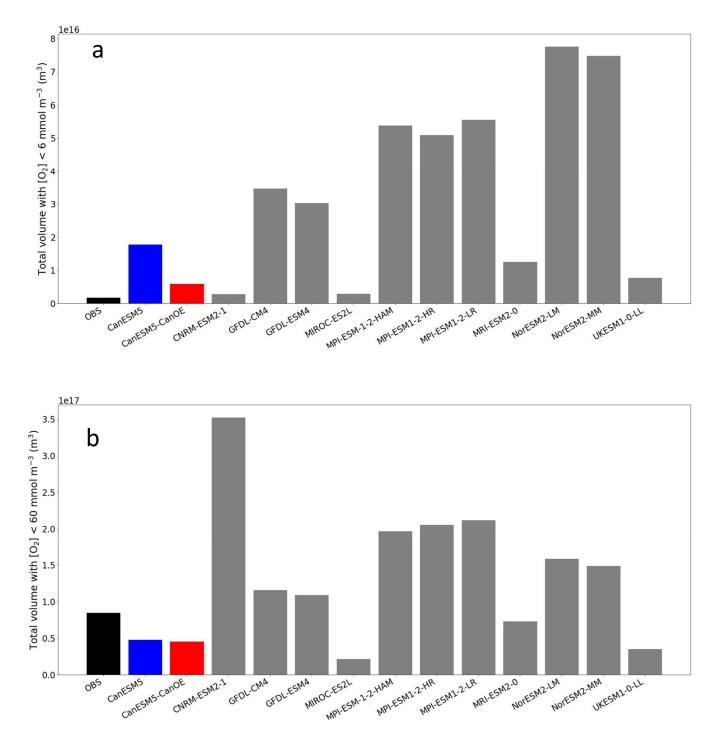


Figure 5 - Total volume of ocean with oxygen (O_2) concentration less than (a) 6 mmol m⁻³ (mean for last 30 years of the historical experiment) and (b) 60 mmol m⁻³. Observation are from WOA2018.

513	total volume is highly variable among models (note, however, that there are several clusters of
514	related models with quite similar totals). CanESM5 and CanESM5-CanOE have among the
515	lowest total volumes (i.e., the interior ocean is relatively well ventilated) and are among the
516	nearest to the observed total. For $[O_2] \leq 60 \text{ mmol m}^{-3}$ the bias is, nonetheless, quite large (i.e., the
517	observed volume is underestimated by almost 50% in both models). The volume of water with
518	[O ₂] below the denitrification threshold is overestimated in both CanESM5 and CanESM5-
519	CanOE; CanESM5-CanOE has a much smaller total that is closer to the observed value. The bias
520	in the spatial pattern of hypoxia (not shown) is generally similar to the bias in dissolved oxygen
521	distribution (Figure 2). The low-oxygen regions are generally more concentrated in the eastern
522	tropical Pacific in the models than in observations, and the low-oxygen region in the northwest
523	Pacific is not well reproduced in CanESM models.

524

525 **3.2 Distribution of DIC, alkalinity, and CaCO3 saturation**

526

527 The spatial distribution of aragonite saturation state (Ω_A) at selected depths is shown in Figure 6. The first two depths are the same as in Figure 2, but a much greater depth is also included, as the 528 529 length scale for CaCO₃ dissolution is greater than for organic matter remineralization. In this 530 case the observations are a combination of GLODAPv2 (Key et al., 2015; Lauvset et al., 2016) 531 for DIC and alkalinity, and WOA2018 for temperature and salinity. CanESM5 and CanESM5-532 CanOE generally compare well with other models and observations. The low saturation bias in the eastern tropical Pacific is substantially reduced in CanESM5-CanOE compared to CanESM5. 533 On the other hand CanESM5 generally does better than CanESM5-CanOE, or the MEM, at 534 535 reproducing the low saturation states in the northwestern Pacific and the Bering Sea. Both

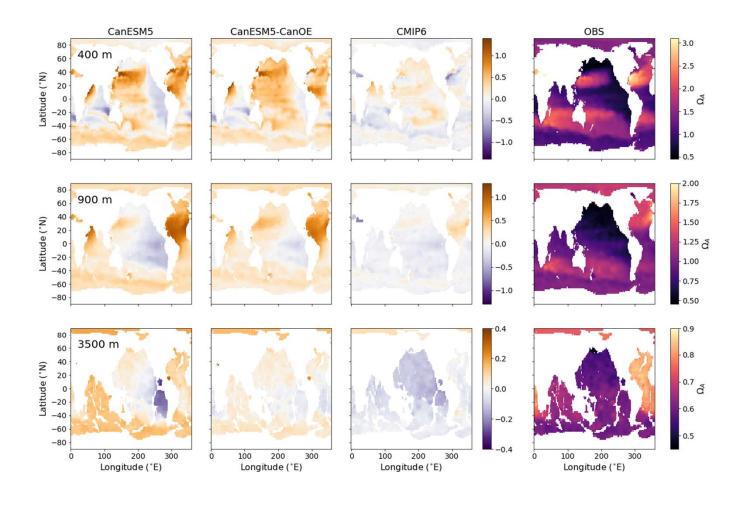


Figure 6 - Global distribution of aragonite saturation (Ω_A) at 400, 900, and 3500 m for CanESM5-CanOE, CanESM5, the mean for other CMIP6 models, and observations (GLODAPv2 + WOA2018). Note different colour scales for different depths. Difference from the observation-based fields are shown in Supplementary Figure S3.

537 CanESM models show a high saturation state bias in the North Atlantic and the well-ventilated

regions of the north Pacific subtropical gyre; these biases are slightly smaller in CanESM5-

539 CanOE. Maps of the calcite and aragonite saturation horizon (Ω =1) depth are shown in

540 Supplementary Figure S3; these generally confirm the same biases noted in Figure 6.

541

Zonal mean distributions of aragonite saturation state (Ω_A), calcite saturation state (Ω_C), and 542 carbonate ion concentration ([CO₃⁻⁻]) and the differences of the models from the observations are 543 shown in Figure 7 (Supplementary Figure S2 includes versions of Figures 6 and 7 that show the 544 modelled fields). The models generally compare well with the observations in the representation 545 546 of the latitude/depth distribution of high and low saturation waters. CanESM5 has a high saturation bias in low-latitude surface waters that is somewhat reduced in CanESM5-CanOE. 547 Both CanESM5 models show a high saturation bias in Northern Hemisphere intermediate (e.g., 548 549 200-1000 m) depth waters that is larger than in the MEM. This is primarily a result of low Ω in the North Atlantic Ocean (Figure 6). 550

551

552 Taylor diagrams for a range of depths are shown for DIC in Figure 8 and for Ω_A in Figure 9 (for 553 alkalinity, see Supplementary Figure S4). As expected, the MEM generally compares favourably 554 with the individual models (e.g., Lambert and Boer, 2001). CanESM5 and CanESM5-CanOE 555 compare favourably with the full suite of CMIP6 models. CanESM5-CanOE shows a gain in 556 skill relative to CanESM5, and both show improvement relative to CanESM2. At 400 m, CanESM2 stands out as having extremely high variance, which is mostly due to extremely high 557 DIC concentrations occurring over a limited area in the eastern equatorial Pacific (not shown). 558 559 This bias is present in CanESM5 and in CMIP6 models generally (Figure 6) but involves much

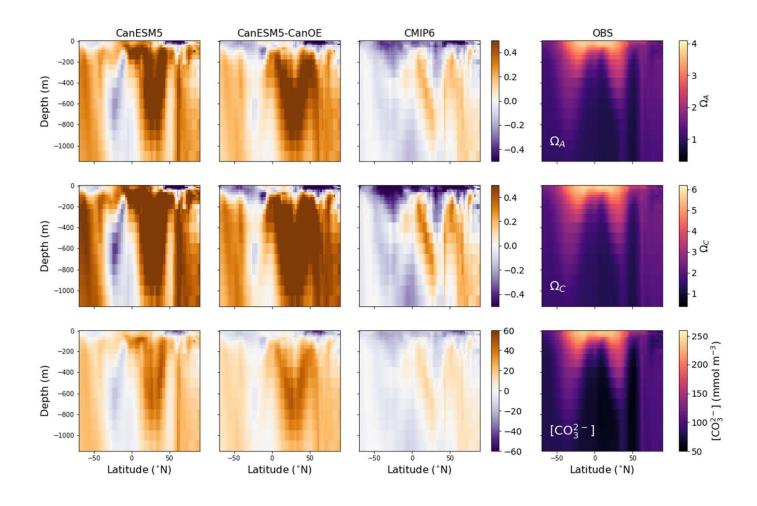


Figure 7 - Latitude-depth distribution of zonal mean (surface to 1150 m) aragonite saturation state (Ω_A), calcite saturation state (Ω_C), and carbonate ion concentration ([CO3⁻⁻]) in mmol m⁻³ for CanESM5-CanOE, CanESM5, the mean for other CMIP6 models, and observations (GLODAPv2 + WOA2018). Difference from the observation-based fields are shown in Supplementary Figure S3.

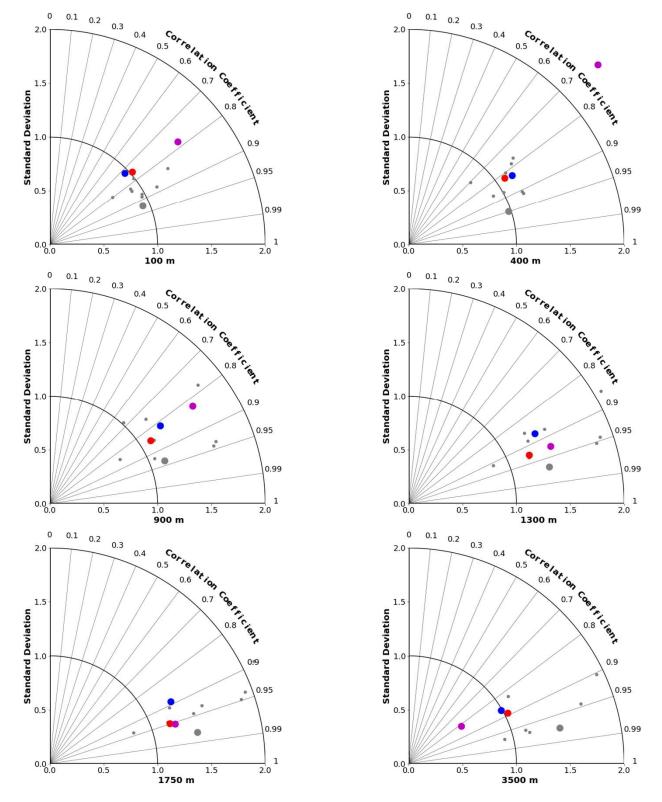


Figure 8 - Taylor diagrams comparing modelled and observed distributions of DIC at specific depths from 100 to 3500 m. Observations are from GLODAPv2 (Key et al., 2015; Lauvset et al., 2016). Red dots represent CanESM5-CanOE, blue dots CanESM5, magenta dots CanESM2, small grey dots other CMIP6 models, and large grey dots the model ensemble mean for all CMIP6 models except CanESM5 and CanESM5-CanOE.

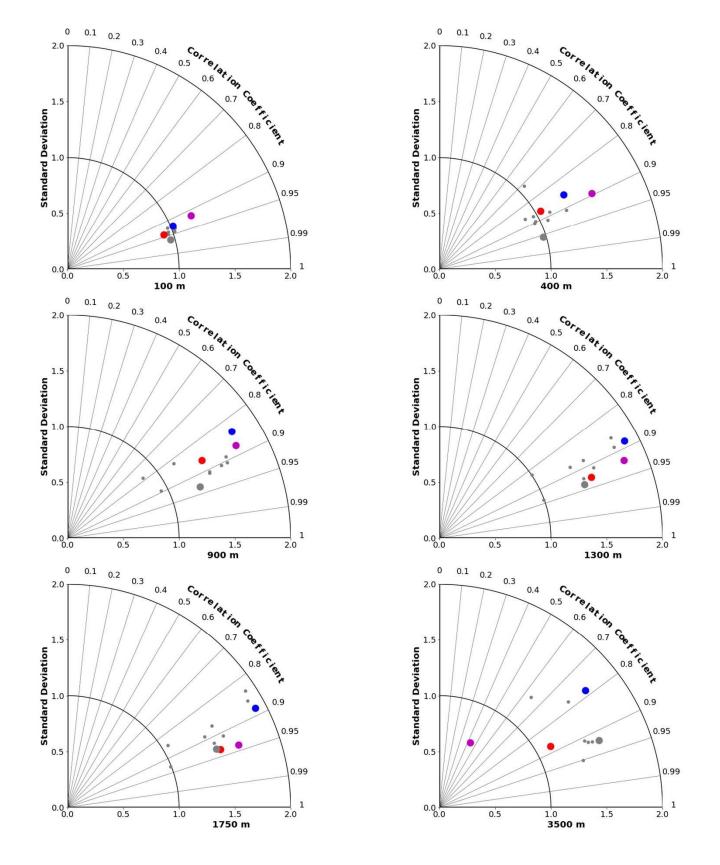


Figure 9 - Taylor diagrams comparing modelled and observed (GLODAPv2 + WOA2018) distributions of Ω_A at specific depths from 100 to 3500 m. Symbol colours as in Figure 8.

563 lower concentrations spread over a larger area.

3.3 N and Fe cycles

567	An important difference between CanESM5 and CanESM5-CanOE is the inclusion of a
568	prognostic Fe cycle. The CMOC iron mask (Zahariev et al., 2008) was a pragmatic solution in
569	the face of resource limitations but is inherently compromised as it can not evolve with a
570	changing climate. The first order test of a model with prognostic, interacting Fe and N cycles is
571	whether it can reproduce the distribution of HNLC regions and the approximate surface
572	macronutrient concentrations within these. CanESM5-CanOE succeeded by this standard,
573	although the surface nitrate concentrations are biased low in the subarctic Pacific and equatorial
574	Pacific and high in the Southern Ocean and in the global mean (Figure 10).
575	
576	The seasonal cycle of the zonal mean surface nitrate concentration for a selection of CMIP6
577	models is shown in Figure 11. CanESM5, CanESM5-CanOE, and CNRM-ESM2-1 reproduce the
578	equatorial enrichment and the low concentrations in the tropical-subtropical latitudes fairly well.
579	Some models either have very weak equatorial enrichment (MPI-ESM1-2-LR) or too high a
580	concentration in the off-equatorial regions (UKESM1-0-LL, NorESM2-LM). UKESM1-0-LL
581	has very high concentrations throughout the low-latitude Pacific, which biases the ensemble
582	mean (Figure 11). Supplementary Figure S6 shows the same data as Figure 11 but for a more
583	limited latitude range to better illustrate model behaviour in the tropics. CanESM5, CanESM5-
584	CanOE, and CNRM-ESM2-1 reproduce the seasonal cycle of tropical upwelling (e.g., Philander
585	and Chao, 1991), with highest concentrations in summer.

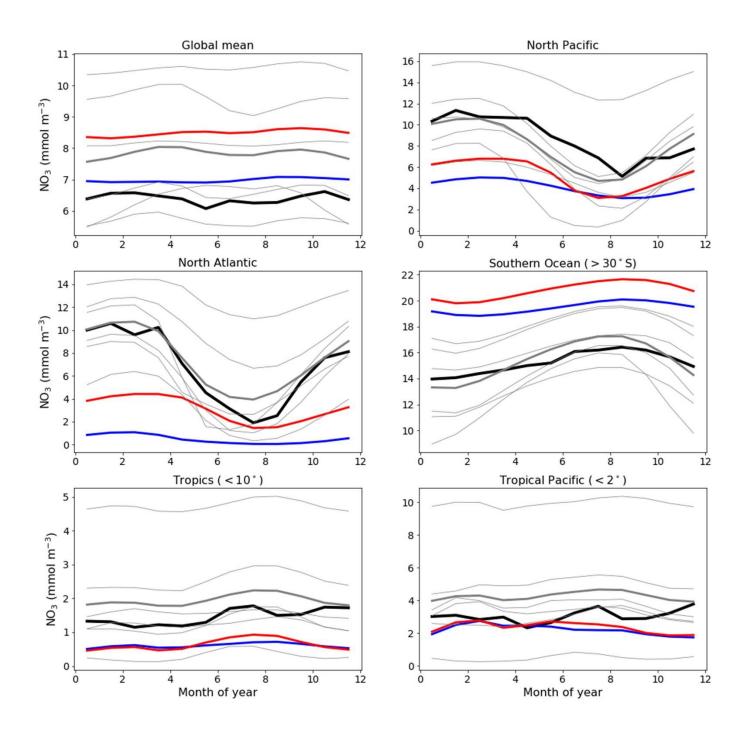


Figure 10 - Climatological seasonal cycle of surface nitrate concentration averaged for selected ocean regions. Thick red line represents CanESM5-CanOE, thick blue line CanESM5, thick black line observations (WOA2018), thin grey lines individual CMIP6 models, and thick grey line the model ensemble mean (excluding CanESM5 and CanESM5-CanOE). Regional boundaries are given in Supplementary Table S5 and Supplementary Figure S5.

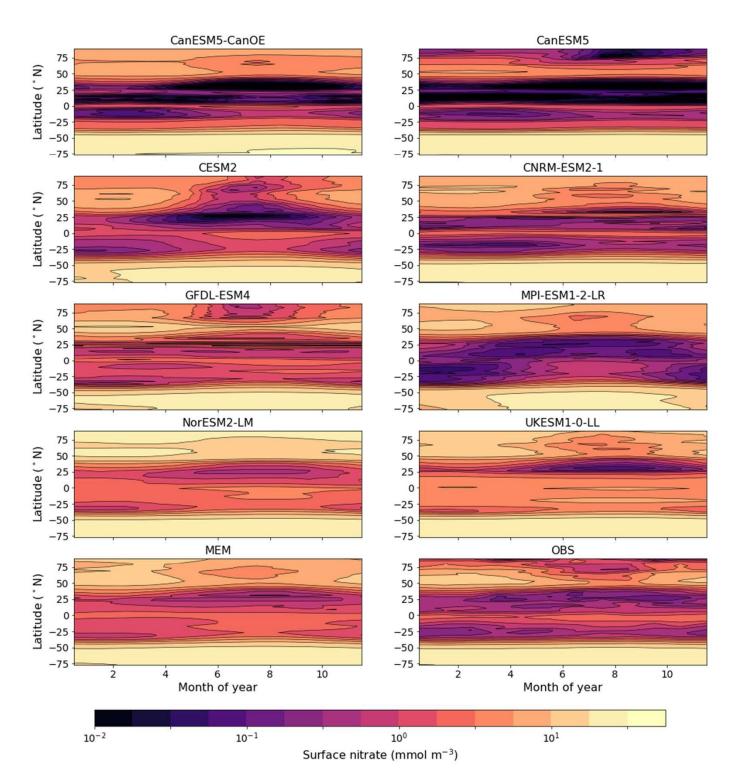


Figure 11 - Climatological seasonal cycle of zonal mean surface nitrate concentration for a selection of CMIP6 models, a model ensemble mean (MEM) excluding CanESM5 and CanESM5-CanOE, and an observation-based data product (WOA2018). An alternate version showing only latitudes <20° is given in Supplementary Figure S6.

588 The surface distribution of dissolved iron (dFe) in various CMIP6 models is shown in Figure 12. For Fe there is no observation-based global climatology with which to compare the model 589 solutions (some comparisons to available profile data are shown in Supplementary Figures S9b-590 h). CanESM5-CanOE shows a similar overall spatial pattern to other models, and generally falls 591 in the middle of the spread, particularly regarding concentrations in the Southern Ocean. Several 592 593 models show extremely high concentrations in the tropical-subtropical North Atlantic (Sahara outflow region). CanESM5-CanOE, along with CNRM-ESM2-1 and CESM2, has much less 594 elevated concentrations in this region, due to lower deposition or greater scavenging or both. 595 596 CanESM5-CanOE has its lowest concentration in the eastern subtropical South Pacific, which is common to many models (Figure 12). The area of strong surface depletion is generally more 597 spatially restricted in CanESM5-CanOE than in other models, and surface dFe concentrations are 598 599 greater over large areas of the Pacific. Both the north-south and east-west asymmetry of distribution in the Pacific is greater in CanESM5-CanOE than in most other models, some of 600 which show the South Pacific minimum extending westward across the entire basin, and others 601 into the Northern Hemisphere. Only in CESM2 is this minimum similarly limited to the 602 southeast Pacific. 603

604

The mean depth profiles of dFe are shown in Figure 13. Some models show more of a "nutrienttype" (increasing with depth due to strong near-surface biological uptake and subsequent remineralization) profile, some a more "scavenged-type" (maximal at the surface, declining with depth) profile (cf. Li, 1991; Nozaki, 2001), and others a hybrid profile (increasing downward but with a surface enrichment). CanESM5-CanOE is at the "nutrient-type" end of spectrum with a generally monotonic increase with depth to a near-constant deep-water concentration of 0.6 nM

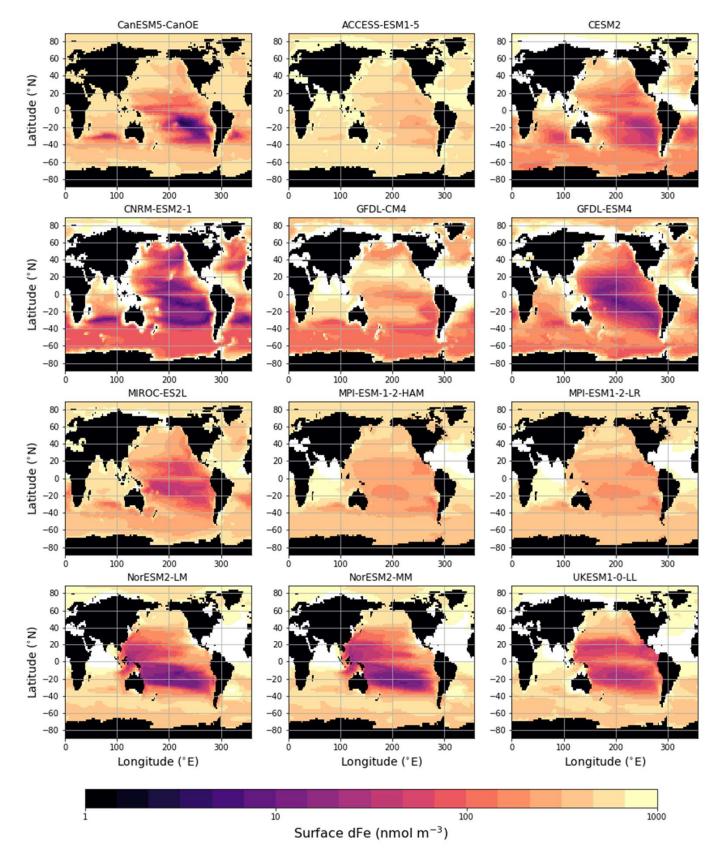


Figure 12 - Global distribution of dissolved iron (dFe) concentration (log10 of concentration in nmol m⁻³) at the ocean surface for CanESM5-CanOE and other CMIP6 models that published this field. Concentrations exceeding 1000 nmol m⁻³ are masked white. CanESM5 is not included because it does not have prognostic iron. 41

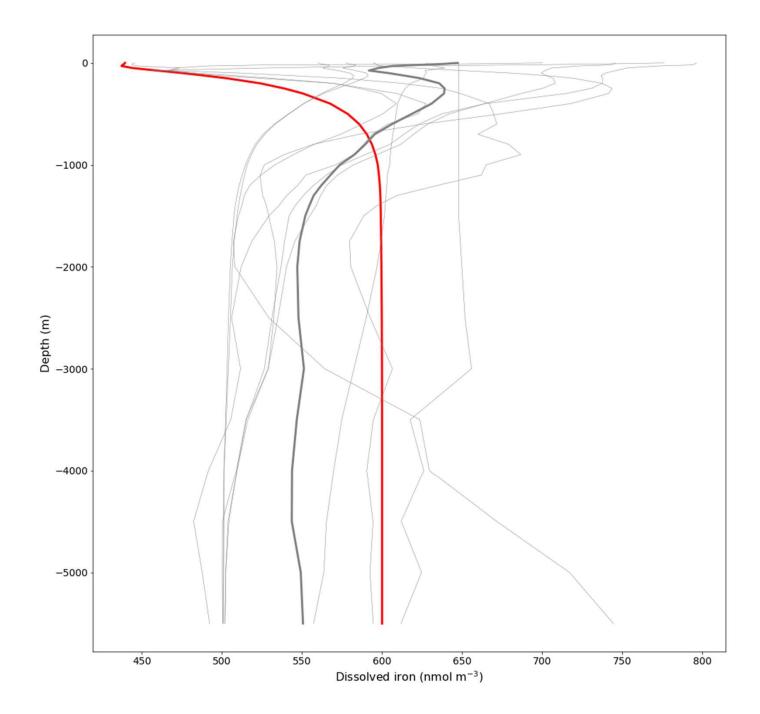


Figure 13 - Global mean depth profiles of dissolved iron concentration for CanESM5-CanOE and other CMIP6 models that published this field. GFDL-CM4 is excluded because it has very high concentrations (>2000 nmol m⁻³) near the surface. Thick red line represents CanESM5-CanOE, thin grey lines individual CMIP6 models, and the thick grey line the model ensemble mean (excluding CanESM5-CanOE and GFDL-CM4).

and a very slight near-surface enrichment (see also Supplementary Figures S9b,c).

614

Mean surface nitrate and dFe concentrations for selected ocean regions are shown in Figure 14. 615 CanESM5-CanOE shows concentrations that are within the range of CMIP6 models, although in 616 some cases at the higher or lower end. Surface nitrate concentrations generally compare 617 favourably with the observation-based climatology, but are biased low in HNLC regions other 618 619 than the Southern Ocean. These biases are not necessarily a consequence of having too much or too little iron. For example, in the Southern Ocean CanESM5-CanOE has among the highest 620 surface nitrate concentrations, but it also has some of the highest dFe concentrations, and the 621 622 high nitrate bias is present in CanESM5 as well. Comparisons with the limited GEOTRACES data available suggest that near surface dFe concentrations in the Southern Ocean are biased high 623 624 rather than low in CanESM5-CanOE (not shown). One region where there does seem to be a 625 strong correlation between surface nitrate and dFe concentrations is the western subarctic Pacific. All but two models (CNRM-ESM2-1, NorESM2-LM) fall along a spectrum from high 626 Fe / low nitrate to low Fe / high nitrate. CanESM5-CanOE falls near the high Fe / low nitrate end 627 of the range. 628

629

Surface nitrate concentrations along the Pacific equator during the upwelling season (JuneOctober) for CanESM5 and CanESM5-CanOE are shown in Figure 15. The range of other
CMIP6 models is not shown here because it is large and therefore adds little information (see
Figure 11 and Supplementary Figure S6). CanESM5-CanOE better represents the east-west
gradient, while CanESM5 has slightly higher concentrations in the core upwelling region. Both
models underestimate the highest concentrations around 100°W. Although some localized

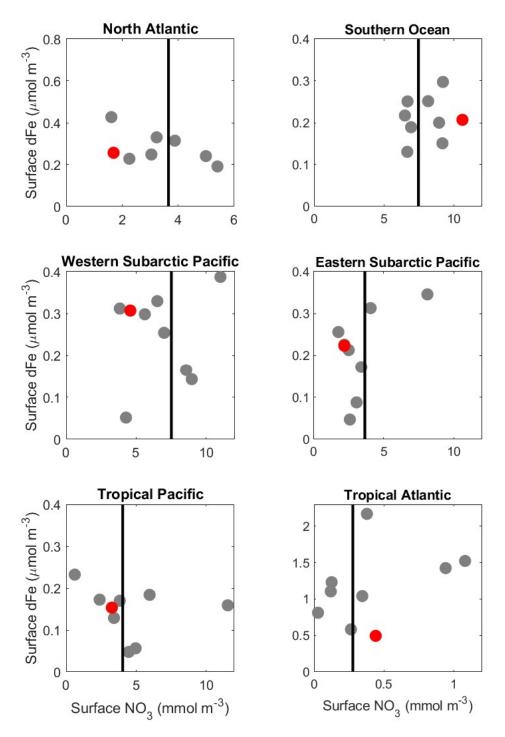


Figure 14 - Mean surface nitrate (NO3) vs. dissolved iron (dFe) concentrations in different oceans, including the major high nutrient / low chlorophyll (HNLC) regions. CanESM5-CanOE is shown as a red dot and other CMIP5 models as grey dots (CanESM5 is not included because it does not have iron). Observed NO3 is shown as a vertical black line as there are no observational estimates of dFe concentration. For GFDL-CM4, nitrate is estimated as phosphate x 16. Region definitions are given in Supplementary Table S5 and Supplementary Figure S5.

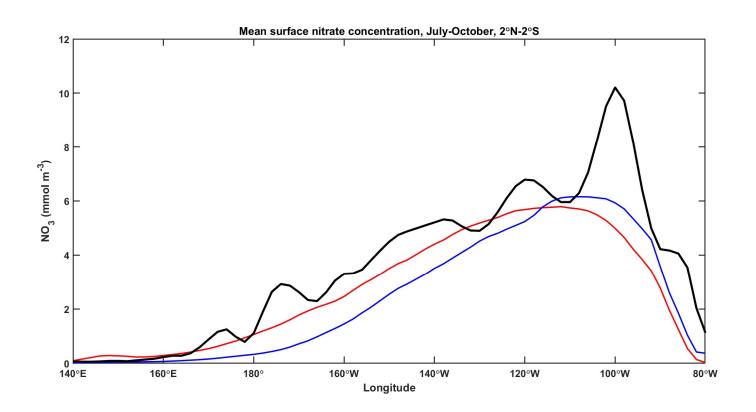


Figure 15 - Surface nitrate (NO3) concentrations along the Pacific equator (mean from 2°S-2°N) during the upwelling season (June-October) for CanESM5-CanOE (red), CanESM5 (blue), and WOA2018 observations (black).

maxima in this data product are due to undersampling, equatorial upwelling is strong at this
location (e.g., Lukas, 2001) and the spatial coherence of the data strongly suggests that this
maximum accurately reflects reality. It should be noted that CanESM5 iron limitation is
calculated from a version of the same data product; however, the Fe mask is based on the
minimum nitrate concentration over the annual cycle, whereas the data shown here are for the
upwelling season.

644

645 3.4 Plankton biomass, detritus, and particle flux

646

The relative abundance of the four living plankton groups are shown in Figure 16 for a range of ocean regions. Both CanESM models mostly compare favourably with observation-based estimates of phytoplankton biomass, except in the tropics where CanESM5-CanOE has very high biomass. Both CanESM models have low phytoplankton biomass in the North Atlantic. In the North Pacific and the Southern Ocean, CanESM5-CanOE reproduces the observation-based estimates well, and CanESM5 slightly less well. The general pattern is that large and small phytoplankton have similar abundance, and are substantially more abundant than zooplankton.

Part of the rationale for multiple food chains is that they better represent the way that actual
plankton communities adapt to different physical ocean regimes and therefore are better able to
simulate distinct ocean regions with a single parameter set (e.g., Chisholm, 1992; Armstrong,
1994; Landry et al., 1997; Friedrichs et al., 2007). The expectation is that small phytoplankton
will be more temporally stable and large phytoplankton will fluctuate more strongly between
high and low abundances. The mean annual cycles of surface chlorophyll largely conform to this

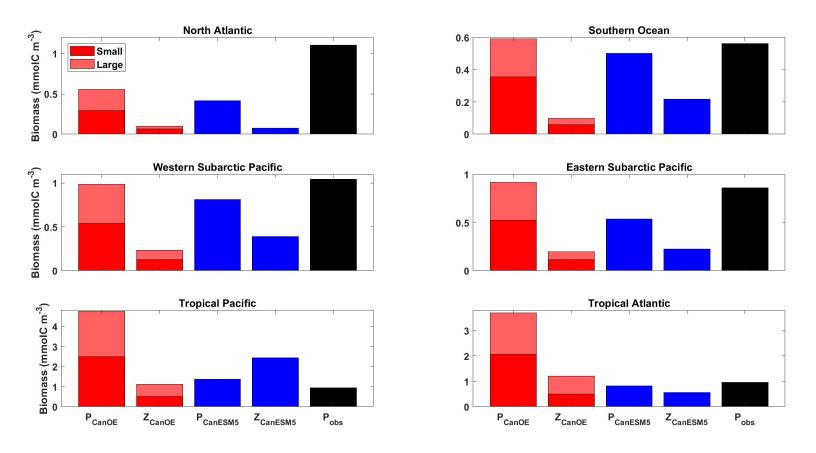


Figure 16 - Annual mean surface ocean concentration of large and small phytoplankton and zooplankton in CanESM5-CanOE (red) and of phytoplankton and zooplankton in CanESM5 (blue) for the representative ocean regions shown in Figure 14. Observational estimates (black) are for phytoplankton biomass calculated from satellite ocean colour estimates of surface chlorophyll (SeaWiFS/MODIS; Tesdal et al. 2016), assuming a carbon-to-chlorophyll ratio of 50 g/g. Region definitions are given in Supplementary Table S5 and Supplementary Figure S5. 662 pattern, e.g., in the North Atlantic and the western subarctic Pacific large phytoplankton are dominant in summer and much more variable over the seasons (Figure 17). Compared to 663 observations, CanESM5 models underestimate the amplitude of the seasonal cycle in the North 664 Atlantic and overestimate it in the North Pacific. CanESM5 shows a stronger and earlier North 665 Atlantic spring bloom compared to CanESM5-CanOE; the observations are in between the two 666 667 in terms of timing, and both models underestimate the amplitude (Figure 17). In the tropics, the seasonal cycle is weak. CanESM5-CanOE in the tropical Atlantic shows the expected seasonal 668 cycle but not the expected dominance of large phytoplankton in summer. CanESM5-CanOE 669 670 generally overestimates the total near surface chlorophyll in both the tropical Pacific and the tropical Atlantic. 671

672

Zooplankton biomass (especially microzooplankton) is also somewhat difficult to test against 673 674 observations, but our model concentrations appear to be biased low. Stock et al. (2014) estimated depth-integrated biomass of phytoplankton, mesozooplankton, and microzooplankton for a range 675 of oceanic locations in which intensive field campaigns have occurred (estimates of 676 microzooplankton biomass are relatively sparse). They found that in most locations 677 phytoplankton and (combined) zooplankton biomass are of comparable magnitude, whereas in 678 679 CanESM5-CanOE zooplankton biomass is consistently lower (Figure 16). The global integral biomass of mesozooplankton is about an order of magnitude less than the 0.19 PgC estimated by 680 681 Moriarty and O'Brien (2013). The CanESM5 total of 0.14 Pg is relatively close to the Moriarty 682 estimate but implicitly includes microzooplankton.

683

684 Surface chlorophyll and POC for CanESM5-CanOE and for ocean colour observational data are

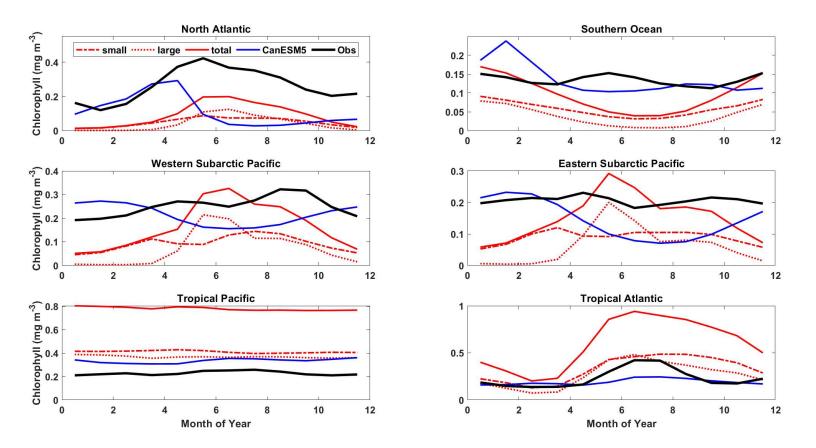


Figure 17 - Mean annual cycle of surface chlorophyll for the representative ocean regions shown in Figures 14 and 16. CanESM5-CanOE large and small phytoplankton concentrations are shown separately and combined (red) along with CanESM5 (blue) and observational estimates (black). Region definitions are shown in Supplementary Table S5 and Supplementary Figure S5.

686 shown in Figure 18 (POC in the model is the sum of phytoplankton, microzooplankton, and detrital carbon). The observations have a lower limit for POC that is not present in the model 687 (~17 mgC m⁻³), which is unsurprising given the processes neglected in the model, i.e., in regions 688 of very low chlorophyll there is still substantial dissolved organic carbon, bacteria that consume 689 it, and microzooplankton that consume the bacteria and produce particulate detritus. The 690 691 observational data show a fairly linear relationship at low concentrations, but with a curvature that implies a greater phytoplankton fraction in more eutrophic environments (cf. Chisholm, 692 1992). The model, by contrast, shows a fairly linear relationship over the whole range of 693 694 concentrations. In other words, the phytoplankton share of POC is higher and more constant in the model than in the observations. The living biomass (phytoplankton + microzooplankton) 695 fraction of total POC in CanOE is generally in excess of 50% (not shown), which is implausible 696 for a real-world oceanic microbial community (e.g., Christian and Karl, 1994) but consistent 697 with the relatively low rates of export from the euphotic zone. 698

699

Export production for a range of CMIP6 models is shown in Figure 19a. CanESM5-CanOE is at 700 the low end of the range. Observations are not shown because the range of observational 701 702 estimates covers the entire range of model estimates (e.g., Siegel et al., 2016). Note also that CanESM5 export is quite a bit lower than in CanESM2, which is relatively high for CMIP5 703 models (not shown). The difference between CanESM2 and CanESM5 is attributable primarily 704 705 to different circulation, although the different initialization fields for nitrate might also play a 706 small role. The lower rate in CanESM5-CanOE is consistent with the above results regarding plankton community structure (e.g., the concentration of detritus is generally low compared to 707 708 living biomass), as well as the lower sinking rate for small detritus. The latitudinal distribution of

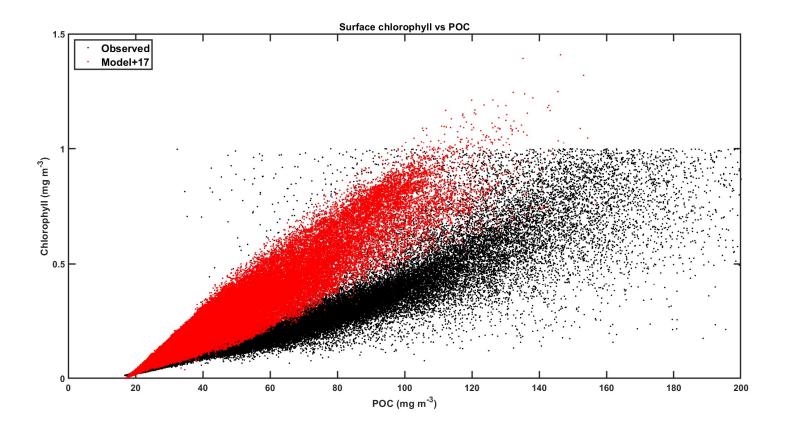


Figure 18 - Climatological surface particulate organic carbon (POC) vs. chlorophyll for CanESM5-CanOE (red) and observations (black). Data are for all ocean grid points ($2x2^{\circ}$ uniform global grid) for all months of the year where observational data are available. Model POC is offset 17 mg m⁻³ for illustrative purposes. Observed chlorophyll concentrations >1 mg m⁻³ are excluded as they largely represent coastal areas poorly resolved by coarse resolution global ocean models.

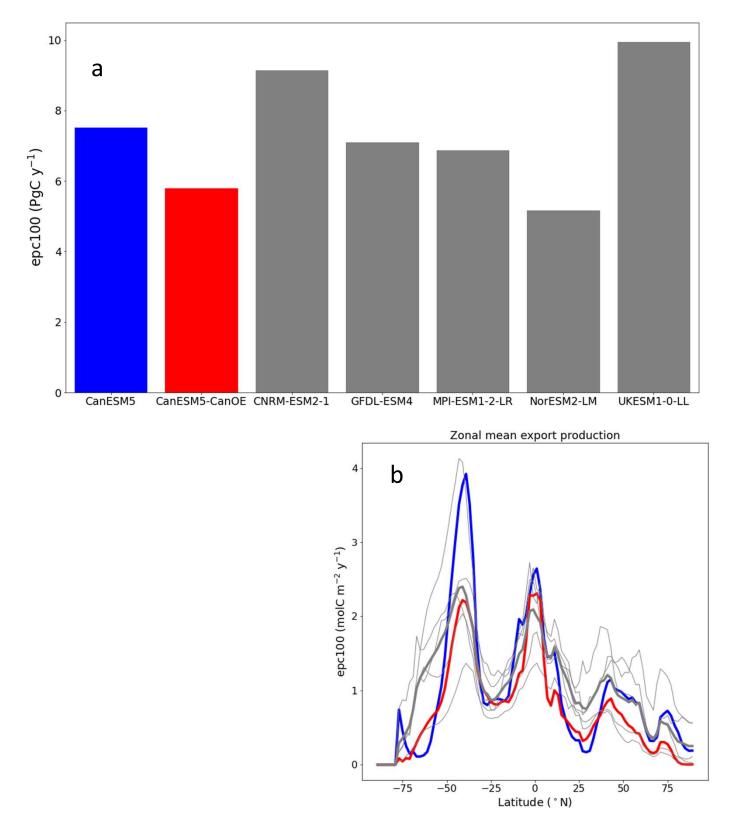


Figure 19 - (a) Global total export production (epc100) in PgC y⁻¹ (b) and zonal mean export production in molC m⁻² y⁻¹ according to selected CMIP6 models (mean for 1985-2014 of historical experiment). Thick red line represents CanESM5-CanOE, thick blue line CanESM5, thin grey lines individual CMIP6 models, and thick grey line the model ensemble mean (excluding CanESM5 and CanESM5-CanOE).

711	export is shown in Figure 19b. CanESM5 shows very high export in the mid-latitudes of the
712	Southern Ocean, similar to CanESM2 (not shown). Both CanESM5 and CanESM5-CanOE show
713	latitudinal patterns consistent with the range of other CMIP6 models. CanESM5 has slightly
714	greater export in the equatorial zone; in both CanESM5 and CanESM5-CanOE the equatorial
715	enrichment attenuates very rapidly with latitude and the rates are low in the subtropics.
716	
717	3.5 Historical trends
718	
719	Cumulative ocean uptake of CO ₂ is shown in Figure 20 for the historical experiment (1850-
720	2014). CanESM models are biased low relative to observation based estimates (~145 PgC, see

Friedlingstein et al., 2020) and the MEM (144 PgC, Figure 20), but fall well within the spread of

722 CMIP6 models. CanESM5-CanOE has lower cumulative uptake than CanESM5 by ~10 PgC. As

the models were not fully equilibrated when the historical run was launched, this difference does

not necessarily arise from the biogeochemical model structure; part of the difference can be

attributed to differences in the spinup protocol (cf. Séférian et al., 2016). The drift in the

piControl experiment over the 165 years from the branching off of the historical experiment is -

10.0 PgC in CanESM5-CanOE and -5.1 PgC in CanESM5 (see Supplementary Table S6), so

drift accounts for about half (48%) of the difference in net ocean CO_2 uptake. The spatial

distribution of anthropogenic DIC is very similar between CanESM5 and CanESM5-CanOE

730 (Supplementary Figure S7).

731

The long-term trend in global total export production is shown in Figure 21. The model valuesmust be normalized in order to compare trends, since the differences among means are large

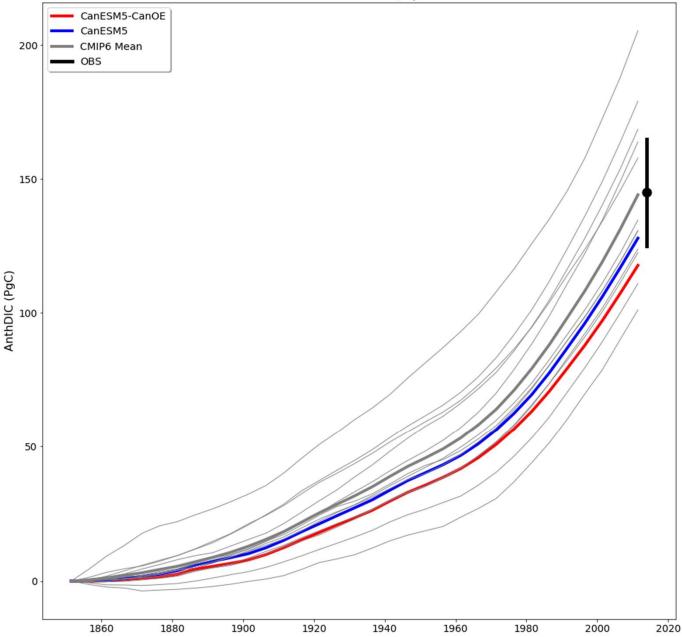


Figure 20 - Cumulative ocean uptake of carbon dioxide (CO_2) as anthropogenic dissolved inorganic carbon (AnthDIC) in PgC over the course of the historical experiment (1850-2014). Data are shown as successive fiveyear means. CMIP6 mean (thick grey line) indicates ensemble mean for CMIP6 models (thin grey lines) excluding CanESM5 (blue) and CanESM5-CanOE (red). An observation-based estimate of 145±20 PgC (Friedlingstein et al., 2020) is shown for nominal year 2014 (black).

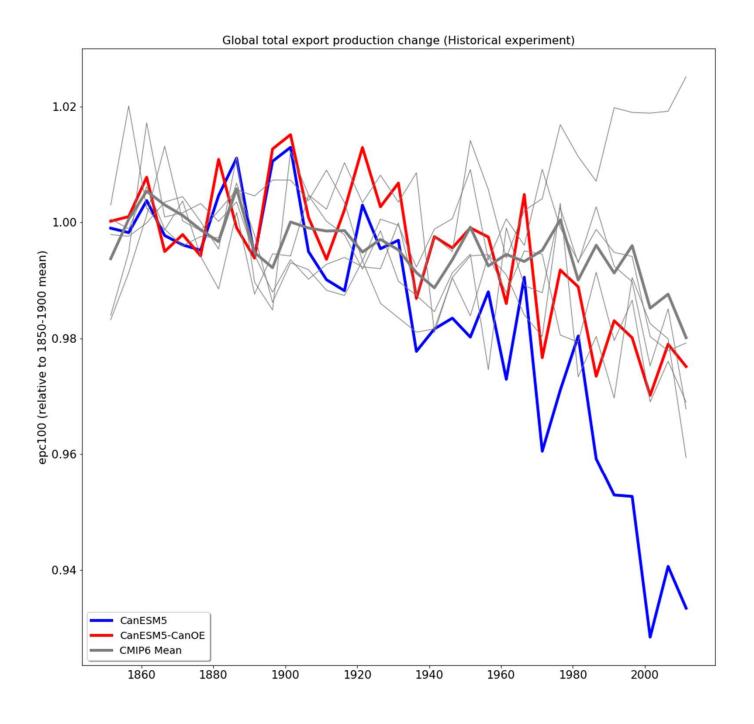


Figure 21 - Change in export production (epc100) over the course of the historical experiment (1850-2014), normalized to the 1850-1900 mean. Data are shown as successive five-year means. Thick red line represents CanESM5-CanOE, thick blue line CanESM5, thin grey lines other CMIP6 models, and thick grey line the ensemble mean of non-CanESM models.

736 compared to the changes over the historical period (Figure 19). Such trends are difficult or impossible to meaningfully constrain with observations, but the general expectation has been that 737 export will decline somewhat due to increasing stratification (e.g., Steinacher et al., 2010). 738 739 CanESM5 shows a greater decline than most other CMIP6 models, while CanESM5-CanOE is more similar to non-CanESM models. The change in CanESM5 is geographically widespread 740 and not concentrated in a specific region or regions: export is maximal in the tropics and the 741 northern and southern mid-latitudes (Figure 19b) and declines over the historical period in all of 742 these regions (Supplementary Figure S8). In CanESM5-CanOE, export declines in the same 743 744 regions, but the magnitude of the change is smaller, and in the Southern Ocean increases and decreases in different latitude bands largely offset each other. 745

746

The trend in the volume of ocean water with O₂ concentration less than 6 or 60 mmol m⁻³ is 747 748 shown in Figure 22. Again, the totals are normalized to a value close to the preindustrial, as the differences among models are large (Figure 5). For the volume with <60 mmol m⁻³, CanESM 749 models show relatively little change; in CanESM5 the volume actually declines slightly, while in 750 751 CanESM5-CanOE it increases, but the total change is <1% in each case. As with the baseline volumes, the range among models is large, with one model showing an increase approaching 752 10% of the total volume estimated for WOA2018 (Figures 5b and 22b). For the volume with <6 753 mmol m⁻³ (Figure 22a), CanESM models are among the most stable over time. In CanESM5, the 754 755 volume again declines, although this is within the range of internal variability. Again some 756 models show fairly large excursions, but in this case none shows a strong secular trend over the 757 last half-century.

758

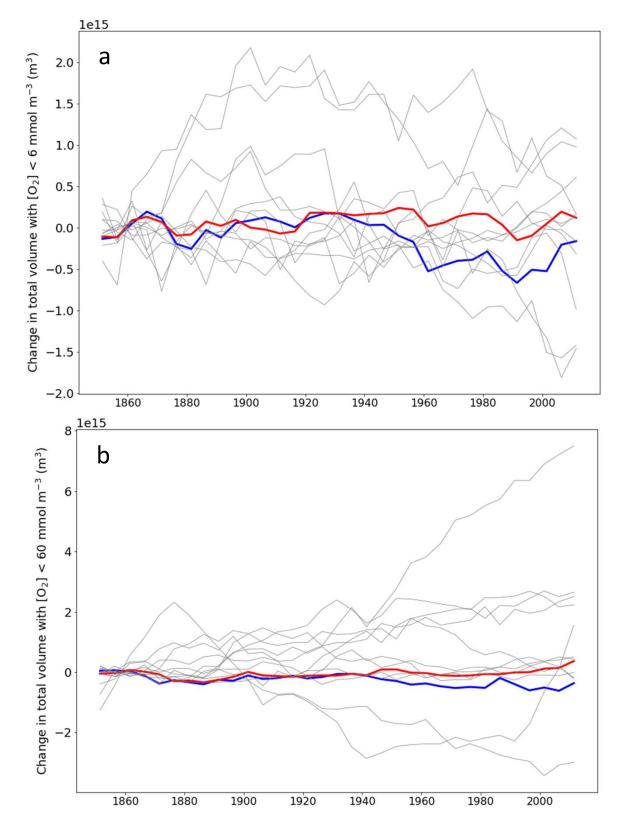


Figure 22 - (a) Change in total ocean volume with oxygen (O_2) concentration less than (a) 6 mmol m⁻³ and (b) 60 mmol m⁻³ over the course of the historical experiment (1850-2014), normalized to the 1850-1870 mean. Data are shown as successive five-year means. Thick red line represents CanESM5-CanOE, thick blue line CanESM5, and thin grey lines other CMIP6 models.

762	CanESM5 and CanESM5-CanOE are new coupled ocean-atmosphere climate models with
763	prognostic ocean biogeochemistry. The two have the same physical climate (in experiments with
764	specified atmospheric CO ₂) and differ only in their ocean biogeochemistry components.
765	CanESM5-CanOE has a much more complex biogeochemistry model including a prognostic iron
766	cycle. We have presented results that assess how these two models simulate the overall
767	distribution of major tracers like DIC, alkalinity, nitrate and oxygen, as well as analyses of the
768	interaction of the iron and nitrogen cycles, plankton community structure, export of organic
769	matter from the euphotic zone, and historical trends over 1850-2014.
770	
771	The overall distribution of major tracers indicates that both models do a reasonable job of
772	simulating both biogeochemical (e.g., export and remineralization of organic matter) and
773	physical (e.g., deep and intermediate ocean ventilation) processes. The volume of ocean with
774	oxygen concentration below 6 or 60 μ M compares favourably with other CMIP6 models (Figure
775	5), and is among the most stable over historical time (Figure 22). CanESM5-CanOE has a
776	substantially lower volume of water with $[O_2] \le 6 \mu M$ than CanESM5 and much closer to
777	observation-based estimates (Figure 22a). Both models are biased slightly low in terms of
778	historical uptake of anthropogenic CO2, which may indicate weak Southern Ocean upwelling or
779	too shallow remineralization of DIC or both (Figure 20). The spatial distribution of
780	anthropogenic DIC is very similar between the two models (Supplementary Figure S7), which is
781	expected as it is mainly a function of the physical ocean model circulation. However, CanESM5
782	has higher concentrations in the main areas of accumulation, particularly the North Atlantic and

the Southern Ocean. This probably indicates more efficient removal and export of 'natural' DIC
by the plankton, particularly in the Southern Ocean upwelling zone (Figure 19), and deeper
average remineralization, with the caveat that the preindustrial control simulations had different
degrees of equilibration when the historical experiment was launched (cf. Séférian et al., 2016,
Supplementary Table S6).

788

789 Analysis of phytoplankton and zooplankton biomass concentrations show that CanESM5 and CanESM5-CanOE compare somewhat favourably with available observational data but do have 790 distinct biases. In particular, both zooplankton biomass and detrital organic matter concentration 791 792 tend to be very low in CanESM5-CanOE; the total biomass of the plankton community and the standing crop of particulate organic matter are dominated by phytoplankton (e.g., Figure 17). 793 Regional biases differ between the two models, with CanESM5-CanOE showing excessively 794 795 large phytoplankton biomass in the tropics. We note, however, that the seasonal cycle of equatorial upwelling and the formation of the equatorial Pacific HNLC are reproduced rather 796 well by our models (e.g., Figures 11, 15 and S6), and that CanESM5-CanOE is the first CanESM 797 model to have genuinely simulated this as an emergent property (see section 3.3). In CanESM5-798 799 CanOE, decoupling of large and small phytoplankton populations associated with seasonal 800 upwelling or convection (see below) is observed in some regions but not others.

801

Global export production is biased low, particularly in CanESM5-CanOE. This is due in part to
the biogeochemical model and in part to ocean circulation. CanESM5 has the same ocean
biology as CanESM2 but a different physical ocean model, and global ocean export production is
substantially lower in CanESM5. It is lower still in CanESM5-CanOE (Figure 19). We note that

CanESM5 performs better than CanESM2 on most metrics of physical ocean model evaluation 806 (Swart et al., 2019a), and shows a more realistic distribution of major tracers like DIC (Figure 8). 807 While the range of observation-based estimates of global ocean export production is large, and 808 encompasses the full range of CMIP5 and CMIP6 models, the change between CanESM2 and 809 CanESM5 is large. Changes in the physical ocean are not entirely independent of the 810 811 biogeochemistry model even when the latter is ostensibly identical. In CanESM2 and CanESM5, iron limitation is specified as a spatially static 'mask' based on the observed distribution of 812 surface nitrate, and it is possible that in these two models ocean upwelling occurs in different 813 814 places relative to the specified boundary of the region of Southern Ocean iron imitation (Figure 3 of Zahariev et al., 2008). It is also possible that the lower export production in CanESM5-CanOE 815 is due to low iron supply to the surface waters of the Southern Ocean, but comparison with 816 available observations do not suggest that this is the case. Several biases are common to 817 CanESM5 and CanESM5-CanOE that relate to Southern Ocean upwelling (high Southern Ocean 818 819 surface nitrate concentration, low export production, weak anthropogenic CO₂ uptake) and so are probably more attributable to the physical ocean model than to the Fe submodel. The difference 820 between CanESM2 and CanESM5 bears this out. 821

822

The development of CanOE was undertaken in response to some of the most severe limitations of CanESM2, and in light of our collective experience. In addition to CMOC (Zahariev et al., 2008), previous models developed by members of our group include Denman and Peña (1999; 2002), Christian et al. (2002a; 2002b), Christian (2005), and Denman et al. (2006). Christian et al. (2002a) had a prognostic Fe cycle and multiple phytoplankton and zooplankton species, but had fixed elemental ratios. Christian (2005) incorporated a cellular-regulation model, but only for a single species and without Fe limitation. Christian (2005) had prognostic chlorophyll
whereas Denman and Peña (1999; 2002) and Christian et al. (2002a) used an irradiancedependent diagnostic formulation. Christian et al. (2002a) used multiplicative (Franks et al.,
1986) grazing, which creates stability in predator-prey interactions but severely limits
phytoplankton biomass accumulation under nutrient-replete conditions.

834

One of the most important lessons from Christian et al. (2002a; 2002b) was that when a fixed 835 Fe/N ratio is employed, sensitivity to this parameter is extreme. Because Fe cell quotas are far 836 more variable than N, P, or Si quotas, treating this parameter as constant results in the specified 837 838 value influencing the overall solution far more than any other parameter. CanESM5-CanOE largely succeeded in creating a prognostic Fe-N limitation model that produces HNLC conditions 839 in the expected regions (Figures 10, 11, 14, 15, S6), although surface nitrate concentration is low 840 841 relative to observation-based estimates in some cases. External Fe sources and scavenging parameterizations will be revisited and refined in future versions. In CanESM5-CanOE the 842 scavenging model is very simple, with distinct regimes for concentrations greater or less than 0.6 843 nM; scavenging rates are very high above this threshold which causes deep-water concentrations 844 845 to converge on this value. The generally nutrient-like profile suggest that in CanOE the scavenging rate is quite low for concentrations below 0.6 nM (Figure 13; see also Supplementary 846 Figure S9h). We note that the aeolian mineral dust deposition field employed here is derived 847 from the CanESM atmosphere model; these processes are not presently interactive but could be 848 849 made so in the future.

850

851	A particular issue with CanESM2 was that extremely high concentrations of nitrate occurred
852	under the EBC upwelling regions. This error resulted from spreading denitrification out over the
853	ocean basin so that introduction of new fixed N from N2 fixation would balance denitrification
854	losses within each vertical column, whereas in the real world denitrification is highly localized in
855	the low oxygen environments under the EBCs. CanESM2 did not include oxygen, but CanESM5
856	incorporates oxygen as a 'downstream' tracer that does not feed back on other biogeochemical
857	processes. The incorporation of a more process-based denitrification parameterization in
858	CanESM5-CanOE is independent of the many other processes that are present in CanESM5-
859	CanOE but not in CanESM5: a CMOC-like model with prognostic denitrification is clearly an
860	option. We chose not to include explicit, oxygen-dependent denitrification in CanESM5 because
861	we wanted to maintain a CMOC-based model as close to the CanESM2 version as possible, and
862	because oxygen would not then be a downstream tracer that does not affect other processes.
863	
863	
863 864	Plankton community structure in CanESM5-CanOE is somewhat biased toward high
863 864 865 866	Plankton community structure in CanESM5-CanOE is somewhat biased toward high
863 864 865 866 867	Plankton community structure in CanESM5-CanOE is somewhat biased toward high concentrations of phytoplankton, low concentrations of zooplankton and detritus, and low export
863 864 865 866 867 868	Plankton community structure in CanESM5-CanOE is somewhat biased toward high concentrations of phytoplankton, low concentrations of zooplankton and detritus, and low export (Sect. 3.4). In the development phase, a fair number of experiments were conducted with various
863 864 865 866 867 868 869	Plankton community structure in CanESM5-CanOE is somewhat biased toward high concentrations of phytoplankton, low concentrations of zooplankton and detritus, and low export (Sect. 3.4). In the development phase, a fair number of experiments were conducted with various values of the grazing rates and detritus sinking speeds. A wide range of values of these
863 864 865 866 867 868 869 870	Plankton community structure in CanESM5-CanOE is somewhat biased toward high concentrations of phytoplankton, low concentrations of zooplankton and detritus, and low export (Sect. 3.4). In the development phase, a fair number of experiments were conducted with various values of the grazing rates and detritus sinking speeds. A wide range of values of these parameters was tested, with no resulting improvement in the overall results. Possibly the detrital
863 864 865 866 867 868 869 870 871	Plankton community structure in CanESM5-CanOE is somewhat biased toward high concentrations of phytoplankton, low concentrations of zooplankton and detritus, and low export (Sect. 3.4). In the development phase, a fair number of experiments were conducted with various values of the grazing rates and detritus sinking speeds. A wide range of values of these parameters was tested, with no resulting improvement in the overall results. Possibly the detrital remineralization rates are too high, although primary production is also on the low end of the
863 864 865 866 867 868 869 870	Plankton community structure in CanESM5-CanOE is somewhat biased toward high concentrations of phytoplankton, low concentrations of zooplankton and detritus, and low export (Sect. 3.4). In the development phase, a fair number of experiments were conducted with various values of the grazing rates and detritus sinking speeds. A wide range of values of these parameters was tested, with no resulting improvement in the overall results. Possibly the detrital

⁶ 'replacement', i.e., small phytoplankton and their grazers do not become much more abundant in
more nutrient-rich environments but rather stay at about the same level and are joined by larger
species that are absent in more oligotrophic conditions (see also Chisholm, 1992; Landry et al.,
1997; Friedrichs et al., 2007). The results presented here suggest that this was partially achieved
but further improvement is possible (Figure 17).

879

As to whether the gains in skill with CanESM5-CanOE justify the extra computational cost, 880 Taylor diagrams (Figures 4, 8, 9, and Supplementary Figure S4) show a modest but consistent 881 gain in skill at simulating the major biogeochemical species $(O_2, DIC, alkalinity)$ across 882 883 variables and depths, especially for alkalinity at mid-depths (Supplementary Figure S4), for which CanESM5 displays the least skill relative to other fields or depths. Other processes that are 884 highly parameterized in CanESM5, such as calcification and CaCO₃ dissolution, were not 885 886 addressed in detail in this paper, but are an important factor in determining the subsurface distribution of alkalinity. Again, we emphasize that we are simulating as an emergent property 887 something that is parameterized in CanESM5 (as previously noted for surface nitrate 888 concentration in HNLC regions), and doing at least as well in terms of model skill. As a general 889 890 rule, the potential for improving skill and achieving better results in novel environments (e.g., 891 topographically complex regional domains like the Arctic Ocean and the boreal marginal seas), is expected to be greater in less parameterized models (e.g., Friedrichs et al., 2007; Tesdal et al., 892 2016). 893

894

An updated version of CanESM5 with prognostic denitrification is clearly possible. However, for
the reasons discussed above, a prognostic Fe cycle with a fixed phytoplankton Fe/N remains

problematic, and the model would still have a single detritus sinking speed and remineralization 897 length scale. We are also developing CanOE for regional downscaling applications (Hayashida, 898 2018; Holdsworth et al., 2021), and it is likely that the simplification of having a single particle 899 sinking speed is not well suited to a domain with complex topography and prominent continental 900 shelf and slope. The number of tracers in CanOE is not particularly large compared with other 901 CMIP6 models. We expect to further refine CanOE and its parameterizations, evaluate it against 902 new and emerging ocean data sets (e.g., GEOTRACES, biogeochemical Argo), and 903 incrementally improve CMOC (which we will maintain for a wide suite of physical-climate 904 905 experiments for which ocean biogeochemistry is not central to the purpose). For CMIP6, we chose to keep CMOC as close to the CanESM2 version as possible. This strategy allows us to 906 quantify how much of the improvement in model skill is due to the physical circulation, as is 907 illustrated by greater skill with respect to DIC (Figure 8) and alkalinity (Supplementary Figure 908 S4), particularly at intermediate depths (400-900 m). The CanESM terrestrial carbon model is 909 also undergoing important new developments (e.g., Asaadi and Arora, 2021) and we expect 910 CanESM to continue to offer a credible contribution to global carbon cycle studies, as well as 911 advancing regional downscaling and impacts science. 912

913

914 *Code availability*. The full CanESM5 source code is publicly available at

gitlab.com/cccma/canesm; within this tree the ocean biogeochemistry code can be found at

916 gitlab.com/cccma/cannemo/-/tree/v5.0.3/nemo/CONFIG/CCC_CANCPL_ORCA1_LIM_CMOC

917 or CCC_CANCPL_ORCA1_LIM_CANOE (last access: 21 September 2021). The version of the

code which can be used to produce all the simulations submitted to CMIP6, and described in this

919	paper, is tagged as v5.0.3 and has the associated DOI: https://doi.org/10.5281/zenodo.3251113
920	(Swart et al., 2019b).
921	
922	Data availability. All simulations conducted for CMIP6, including those described in this paper,
923	are publicly available via the Earth System Grid Federation (source_id = CanESM5 or
924	CanESM5-CanOE). All observational data and other CMIP6 model data used are publicly
925	available.
926	
927	Author contributions. Formulation of the overall research goals and aims: JRC, KLD, NS, NCS;
928	Implementation and testing of the model code: JRC, HH, AMH, WGL, OGJR, AES, NCS;
929	Carrying out the experiments: JRC, WGL, OGJR, AES, NCS; Creation of the published work:
930	JRC, HH, AMH, AES, NS, NCS.
931	
932	Competing interests. The authors declare that they have no conflict of interest.
933	
934	Disclaimer. CanESM has been customized to run on the ECCC high-performance computer, and
935	a significant fraction of the software infrastructure used to run the model is specific to the
936	individual machines and architecture. While we publicly provide the code, we cannot provide
937	any support for migrating the model to different machines or architectures.
938	
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